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PRINCIPAL INVESTIGATOR: John A. Katzenellenbogen, Ph.D.

CONTRACTING ORGANIZATION: University of Illinois Champaign, Illinois 61820-6242

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13. Abstract (Maximum 200 Words	s) (abstract should contain no proprie	etary or confidential inform	mation)	1	
We have conceived of an approach to prepare by combinatorial methods, libraries of novel ligands for the estrogen receptor, by the creation of simple amide or five-membered ring heterocyclic core structures that display peripheral substituents (phenols, aliphatic groups, etc.) commonly found in non-steroidal estrogens. These novel estrogens might be useful in the treatment or prevention of breast cancer.					
We have made excellent progress on the preparation of novel estrogens of the diphenyl carboxamide class, the diphenylsulfonamide class, the phenyl benzylcarboxamide and sulfonamide classes, and the pyrazole, oxazole, thiazole, and imidazole classes. Members of some classes have high affinity for the estrogen receptor, and some of them show high binding and potency selectivity for the estrogen receptor subtype alpha and can be adapted as selective estrogen receptor alpha antagonists. We have also developed a convenient solid phase synthesis of the pyrazole class, so that we can prepare conveniently and rapidly larger libraries of the members of what appear presently to be the most promising of these classes of novel estrogens.					
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FINAL REPORT: September 1, 1997 – August 31, 2001

PRINCIPAL INVESTIGATOR: John A. Katzenellenbogen

TITLE: Combinatorial Synthesis For The Expedited Discovery Of Novel Selective Antiestrogens For Breast Cancer Prevention And Therapy

ORGANIZATION: University Of Illinois

INTRODUCTION, GOALS OF THE PROJECT AND APPROACH

[NOTE 1: This project was funded for three years, and a final report was submitted September, 2000, covering the progress for the initial three year period. However, I received a one-year no-cost extension on this project, to complete some of the studies that were still ongoing on August 31, 2000. <u>Thus, this "second" final report is largely the same as the</u> <u>previous final report submitted in September, 2000. It has simply been updated to indicate the</u> <u>additional progress that was made during the year of the no-cost extension. To facilitate</u> <u>review of this second final report, those sections that have been added or modified are so</u> <u>indicated with underlining.]</u>

[NOTE 2: For the purpose of continuity, the three Schemes in the attached Appendix have been taken directly from the original proposal. The details of this final report are contained in seven reprints and three manuscripts that are appended. Specific reference is made to these as "Publications No. 1-10".]

Antiestrogens such as tamoxifen are widely used in the treatment of hormone responsive breast cancer and were recently shown to be effective in breast cancer prevention¹⁻⁵. Tamoxifen, however, as well as the pure ICI antiestrogens, are not ideal agents, because they cause vaginal atrophy and menopausal host flashes, induce osteoporosis (pure antiestrogens), and may cause endometrial and liver cancer (tamoxifen) $^{6-14}$. Thus, there is a need for the development of selective antiestrogens with an improved endocrine profile for use in the treatment and prevention of breast cancer $^{15-16}$. Recent advances in our understanding of the molecular pharmacology of estrogens and the development of new selective antiestrogens for menopausal bone maintenance, suggest that new selective antiestrogens of this type can be discovered ¹⁶⁻²⁵. Up to now, however, this search has not been approached in a systematic fashion ²⁶. Furthermore, the structures of antiestrogens that have been studied to date are quite complex and their synthesis sufficiently challenging, so as not to be amenable to synthesis by solid-phase combinatorial means. Combinatorial synthesis is the fastest growing new technology in pharmaceutical chemistry, and is proving to be a highly expeditious and promising approach to new drug discovery ²⁷⁻³⁵.

In preparing for this project, we analyzed the structures of many selective antiestrogens and found that they possess three common peripheral groups (a phenol, a second aromatic group, and a basic side chain) attached to various core structures. Because the core structure appears to function as a scaffold simply to hold these other appendages together, we then designed six novel core structures that will perform the same scaffold function for the peripheral groups, yet are sufficiently simple that they can be readily prepared by solid-phase combinatorial synthesis.

In this project, we proposed to prepare six novel classes of ligands for the estrogen receptor, based on four functional group so far unexplored in the antiestrogen literature: a carboxamide, a sulfonamide, a pyrazole, and an oxazole (and related thiazole and imidazole). Solution-phase syntheses were first to be developed, and then adapted to solid-phase synthesis using an acid-labile linker attached to the common phenol function. Libraries containing larger numbers of the best of these classes were then to be prepared, and all of these compounds then to be assayed for their binding affinity for the estrogen receptor. The estrogen agonist and antagonist activity of those members with high affinity was then later to be determined in cell transfection and proliferation assays, and those with the most appropriate endocrine activity tested in a uterotrophic assay.

The combination of this novel structural insight leading to the design of new core structures for estrogen receptor ligands that can be readily prepared by combinatorial synthesis, together with a set of simple, but effective assays to establish their hormonal activity, should assist in the discovery of novel selective antiestrogens for the treatment and prevention of breast cancer.

BODY

Experimental Approach

A General Structural Description of Selective Estrogen Receptor Ligands Suggests Alternate Core Structures that Can be Prepared by Solid Phase Combinatorial Chemistry Methods

As a class, selective antiestrogens can be envisioned as having four structural components (Schemes 1 and 2 in attached Appendix [taken from the original proposal]): a core structure (A) onto which are appended three other structural elements, a phenol (B), a second aromatic group (C) and a basic side chain (D). The achievement of an appropriate balance of estrogenic vs antiestrogenic activities in each of these series of selective antiestrogens appears to involve a delicate interaction between these component parts of their structure. Curiously, whereas components B, C, and D are rather similar in almost all of the selective antiestrogens (cf. Scheme 1, selective antiestrogens), the central core structure A, which links together the other three components like a scaffold, is quite variable. This suggests that other core structures could replace this central component in selective antiestrogens.

In the original proposal we proposed to explore new antiestrogens specifically designed to have novel core structures that are readily amenable to solid phase combinatorial synthesis and which may prove to be more tissue selective and efficacious for breast cancer. As explained in Scheme 3 in attached Appendix [taken from the original proposal], we identified three simple structure motifs that are found in ER ligands, around which we designed six new classes of potential ligands for the ER that are based on four new core structures (Scheme 3, right column). We anticipated that these four core structures—a carboxamide, a sulfonamide, a pyrazole, and an oxazole (and a related thiazole and imidazole)—would provide a suitable molecular scaffold for the other three components typically found in selective antiestrogens—the phenol, the second aromatic unit, and the basic side chain—so that these new structural classes would also have selective antiestrogenic activity. The unique feature of these new core structures is that, unlike most antiestrogens develop to date, they can be prepared by combinatorial chemistry means.

The first of the motifs, *motif A is an anti-bibenzyl system*, a structural subunit that is found in the potent estrogen hexestrol, as well as in the antiestrogen hydroxytamoxifen (Scheme 3). The structural analogs of this motif that we planned to explore, the diphenylcarboxamide and diphenylsulfonamide (Classes I and II), are well suited to three-component combinatorial solid phase synthesis methods (see below and Scheme 6). *Motif B is the homolog of the bibenzyl motif*, a substructure that is exemplified in the potent estrogen benzestrol and the selective estrogen raloxifene, but is otherwise largely undeveloped. We planned to explore several structural analogs of the homobibenzyl motif that are better suited for solid phase three-component combinatorial synthesis, benzylic homologs of the carboxamides and sulfonamides (Classes III and IV) and pyrazoles (Class V) (see below and Schemes 6 and 7). Finally, *Motif C, a syn-bibenzyl system* found in tamoxifen and centchroman, is the motivation for the three-

component combinatorial synthesis of a series of heterocyclic analogs, oxazoles, thiazoles, and imidazoles (Class VI; see below and Scheme 8).

As we discuss below, we have made good progress evaluating the synthetic feasibility and ER binding affinity of members of all of these classes, as well as some others.

Results and Discussion for total project, years 1-3, plus one year no-cost extension.

The original statement of work for years 1-3 of this project are listed below.

ORIGINAL STATEMENT OF WORK

Project Period: July 1, 1997–June 30, 2000 (3 years)

Year 1

- Synthesis of representative members of Class I-VI ligands by solution phase methods.
- Isolate, purify, and fully characterize these members.
- Measure estrogen receptor binding affinity of representative members of Class I-VI ligands
- Begin to adapt solution phase syntheses to solid phase.

Year 2

- Complete adaptation of solution phase synthesis to solid phase.
- Isolate and fully characterize representative members produced by solid phase synthesis. Determine yield and characterize impurities.
- Compare estrogen receptor binding of representative members of Class I-VI ligands prepared by solution vs solid phase.
- Begin synthesis of full Class I-VI libraries.
- Begin cell proliferation and cell-based transfection assays.

Year 3

- Complete synthesis of full Class I-VI libraries.
- Complete estrogen receptor assay of full libraries at two concentrations.
- Reassay the members with detectable estrogen receptor binding affinity by quantitative titration assay.
- Assay the members with high estrogen receptor binding affinity in the cell proliferation and cell-based transfection assays.

• Assay uterotrophic activity of the most promising members in rats.

Summary — Through the work we have engaged in on this project over the past three years, we have completed all of the goals of the original proposal, as summarized in the original Statement of Work. This has involved an exploration of the promise of all of the proposed acyclic amide and 5-membered ring heterocyclic systems, and the development of solid phase synthesis methods and large library synthesis and analysis. In addition, we have discovered in the process that some of the most promising heterocyclic ligands, the pyrazoles, have high affinity and potency selectivity for one of the estrogen receptor subtypes, ER α . Because the development of estrogen receptor subtype selective ligands is a very important issue and one of great current interest, we have devoted considerable efforts in years 2, 3, and 4 to pursue this further. We have completed a careful structure-activity relationship study on the pyrazoles, obtaining ultimately a compound that is almost a completely specific activator of ERa, having essentially no activity on ER β at doses where it maximally stimulated ER α , and, based on these findings, we have developed a series of pyrazole basic side chain derivatives that hold considerable promise as potency selective antagonists of ERa. Such compounds could be used to block the action of estradiol through this receptor subtype without interfering with its effect on ERβ.

The work that we have done on this project has also spawned several related lines of inquiry that are currently being pursued in my laboratory with independent support. These include investigations of other 5-membered ring heterocyclic systems as estrogen receptor ligands, namely furans, thiophenes, and pyrroles, as well as carbocyclic analogs, cyclopentadienes and related cyclopentadienones. In addition, based on the favorable results of the work under this project, we have begun to synthesize 6-membered ring heterocycles such as pyrazines, pyrimidines, and pyridazines as estrogen receptor ligands. This work is <u>now</u> beginning to mature, and we already know that we have compounds that show high affinity for the receptor, <u>showing high ER subtype selectivity</u>, and are therefore potentially quite interesting, because their structures are unique in the estrogen receptor field.

We certainly appreciate the support that has been provided to our research efforts in this area through this US Army grant. It has served as a stimulus to us to thrust our laboratory research into new directions that have proved to be very fruitful and potentially useful for the prevention and treatment of breast cancer.

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The details of this final report are contained in <u>seven</u> reprints and <u>three</u> manuscript that are appended. Specific reference is made to these as "Publications No. 1-<u>10</u>".

Publication No. 1:	B. E. Fink, D. S. Mortensen, S. R. Stauffer, Z. D. Aron, J. A.		
	Katzenellenbogen. Novel Structural Templates for Estrogen-Receptor		
	Ligands and Prospects for Combinatorial Synthesis of Estrogens. Chem.		
	& Biol., 1999, 6, 205-219.		
Publication No. 2:	S. R. Stauffer, J. Sun, B. S. Katzenellenbogen, J. A. Katzenellenbogen.		
	Acyclic Amides as Estrogen Receptor Ligands: Synthesis, Binding,		
	Activity, and Receptor Interaction. Bio. Med. Chem., 2000, 8, 1293-1316.		
Publication No. 3:	S. R. Stauffer, J. A. Katzenellenbogen. Solid-Phase Synthesis of Tetra-		
	Substituted Pyrazoles, Novel Ligands for the Estrogen Receptor. J. Comb.		
	Chem., 2000, 2, 318-329.		
Publication No. 4:	S. R. Stauffer, C. J. Coletta, J. Sun, B. S. Katzenellenbogen, J. A.		
	Katzenellenbogen. Pyrazole Ligands: Structure-Affinity/Activity		
	Relationships of Estrogen Receptor-a Selective Agonists. J. Med. Chem.,		
	<u>2000, <i>43</i>, 4934-4947.</u>		
Publication No. 5:	S. R. Stauffer, Y. Huang, C. J. Coletta, R. Tedesco, J. A.		
	Katzenellenbogen. Estrogen Pyrazoles: Defining the Pyrazole Core		
	Structure and the Orientation of Substituents in the Ligand Binding Pocket		
	of the Estrogen Receptor. Bio. Med. Chem., 2001, 9, 141-150.		
Publication No. 6:	S. R. Stauffer, Y. R. Huang, Z. D. Aron, C. J. Coletta, J. Sun, B. S.		
	Katzenellenbogen, J. A. Katzenellenbogen. Triarlpyrazoles with Basic		
	Side Chains: Development of Pyrazole-Based Estrogen Receptor		
	Antagonists. Bio. Med. Chem., 2001, 9, 151-161.		
Publication No. 7:	Y. R. Huang, J. A. Katzenellenbogen. Regioselective Synthesis of 1,3,5-		
	Triaryl-4-alkylpyrazoles: Novel Ligands for the Estrogen Receptor. Org.		
	Lett, 2000, <u>2, 2833-2836.</u>		
Publication No. 8:	D. S. Mortensen, A. L. Rodriquez, K. E. Carlson, J. Sun, B. S.		
	Katzenellenbogen, J. A. Katzenellenbogen. Synthesis and Biological		
	Evaluation of a Novel Series of Furans: Ligands Selective for Estrogen		
	Receptor Alpha. J. Med Chem., 2001, In Press.		
Publication No. 9:	D. S. Mortensen, A. L. Rodriquez, J. Sun, B. S. Katzenellenbogen, J. A.		
	Katzenellenbogen. Furans with Basic Side Chains: Synthesis and		
	Biological Evaluation of a Novel Series of Antagonists with Selectivity for		
	the Estrogen Receptor Alpha. Bioorg. Med. Chem. Letters, 2001, In Press		

Publication No. 10:J. Sun, Y. R. Huang, W. R. Harrington, S. Sheng, B. S. Katzenellenbogen, aJ. A. Katzenellenbogen. Antagonists Selective for Estrogen ReceptorAlpha. Endocrinology. 2001, Submitted

Investigation of Novel Core Structures of Estrogen Receptor Ligands [Publications No. 1 and 2]

In the first of these publications [**Publication No. 1**], we surveyed a series of heterocyclic systems to identify which core structure from among those originally proposed (pyrazole, imidazole, oxazole, thiazole, isoxazole and isothiazole classes) were capable of giving ligands with high affinity for the estrogen receptor. This work definitively identified the pyrazole class as the most promising one, and one tetrasubstituted pyrazole, in particular, had a binding affinity that reached 19% that of estradiol, a truly remarkable finding. This compound is numbered **38d** in this publication.

The second of these publications [**Publication No. 2**] we surveyed a series of amides (carboxamides, thiocarboxamides, and sulfonamides) as potential ligands for the estrogen receptor. Again, we found that high affinity for the receptor was restricted to the diphenylcarboxamide class and then only to those analogs having appropriate substituents. The most promising agend, compound **16g** in this publication, has an affinity 15% that of estradiol. This work included extensive analysis of the conformation of these flexible ligands with the aim of understanding how they fit into the ligand binding pocket.

Development of Solid Phase Synthesis Methods to Prepare the Most Promising Ligands [Publication No. 3]

In this publication, we describe the systematic approach that we took to develop an efficient method to synthesize the pyrazole system on a solid phase. In this included a careful determination of the conditions required to attach the precursor to the support and to effect all of the ensuing reactions, including cleavage of the final product. We used NMR and IR analysis directly on the derivatized beads to assess the progress of the reactions. Finally, we used the optimized methods to prepare a small library (12 members) and then a large library (96 members) of the tetrasubstituted pyrazole class. We also measured the receptor binding affinities of these 96 compounds and thereby developed an interesting structure affinity relationship [See Figure 6, **Publication No. 3**]. The highest affinity ligands are those in which at least two of the aromatic rings bear a 4-hydroxy group.

Structure-Activity Relationships of Tetrasubstituted Pyrazoles and Development of an Estrogen Receptor Alpha Specific Ligand [Publication No. 4]

In this publication, we have taken the most favorable results from Publication Nos. 1 and 3 and extended further the structure-activity analysis of the tetrasubstituted pyrazole ligands. In addition, we have assayed their affinity for the estrogen receptor subtypes ER α and ER β . Remarkably, we find that some of the pyrazoles have especially high affinity for ER α , reaching as high as 75% that of estradiol, yet very low affinity for ERβ. The best of these compounds [numbered 4g (PPT) in Publication No. 4] has a 410-fold affinity preference for ER α . This compound also shows extremely high potency selectivity for ER α in cell-based transcription assays [Figure 2 in **Publication No. 4**]. This compound has proved to be of great interest to the molecular endocrinology community as a probe for assessing the biological roles of ER α and ER β . We have supplied samples of this ER α selective ligand to more than 20 laboratories who are engaged with use in collaborative studies. In this study we also endeavored to use molecular modeling to supplement the results of our structure-activity study so as to be able to predict the orientational mode with which these rather symmetrical pyrazole core ligands are bound by the estrogen receptor. We also have developed a productive collaboration with Wyeth-Ayerst to evaluate PPT further in a series of biological assays, such as uterotropic, bone mineralization, neuroprotection model, hot flush model, ER-subtype specific gene induction in the brain, etc. PPT functions as a full agonist in most of these assays. This work is still underway and is, so far unpublished. A joint publication is anticipated once the studies have been completed.

Synthesis of Pyrazole Core Isomers in a Further Structure Activity Relationship Study of Tetrasubstituted Pyrazole Estrogen Receptor Ligands [Publication No. 5]

Originally, we explored only one isomeric series of tetrasubstituted pyrazoles as ligands for the estrogen receptor, ones in which the three aryl groups were at N-1, C-3 and C-5, and the alkyl group at C-4. In this publication [**Publication No. 5**] we explore pyrazoles of the isomeric series (i.e., 1,3,4-triaryl-5-alkyl). This required the development of a completely new synthetic method, because the original method, when applied to the corresponding monoaryl 1,3-diones underwent a Fischer indole reaction rather than a pyrazole cyclization. We were able to over come this side reaction by performing the synthesis on a 1,3-dione of lesser substitution and then to introduce the final aryl group by a Suzuki coupling method. The overall process proved to be both regioselective for this series and efficient. The most interesting products from this study were those listed a compounds **10a** and **10b** in **Publication No. 5**.

Investigation of Pyrazoles with Basic Side Chains as Potential Potency-Selective Estrogen Receptor Alpha Antagonists [Publications No. 6 and 10]

Because of the high affinity and potency selectivity of certain of the pyrazoles as agonists for ER α , we were interested in seeing whether we could convert them to ER α potency selective

antagonists by adding a basic side chain to the pyrazole core structure. In this publication **[Publication No. 6]**, we attached a basic side chain to the four substituent positions on the tetrasubstituted pyrazoles, and we found that only one of these showed high affinity for the estrogen receptor. This compound showed about a 10-fold potency preference as an antagonist for ER α vs ER β [see compound **5** and Figure **3** in **Publication No. 6**]. It is the basis for continued studies on pyrazole ligands with basic side chains, which stimulated in part the development of the regioselective method for pyrazole synthesis discussed below. In addition, we have investigated further the biological activity of this series of pyrazoles with basic side chains [Publication No. 10], and we have found that one of these, termed methyl-piperidinyl-pyrazole (MPP), is an ER α -selective antagonist. This compound will be a useful pharmacological tool for studying the biological roles of the two ER subtypes.

Development of a Regioselective Synthesis of Pyrazoles [Publication No. 7]

The method we originally devised for the synthesis of the pyrazoles involved the reaction of a hydrazine with a 1,3-diketone and was, therefore, not regioselective. This increasingly became a problem as we wanted to prepare more and more non-symmetrical pyrazoles. In this publication, we have developed an efficient regioselective synthesis of these pyrazoles by the addition of the hydrazine to an enone to give an pyrazoline intermediate. The corresponding anion of this pyrazoline can be alkylated and then oxidized to produce unsymmetrical pyrazoles in high yield and with complete regioselectivity. We have used this method to prepare more than 20 different unsymmetrical pyrazoles, including a variety of ones having different basic side chain substituents on the position where it is most favorable (see discussion above).

Novel Furan Ligands for the Estrogen Receptor [Publication No. 8 and 9]

We have also developed novel ligands for ER based on a different heterocyclic system, furans. In this system we also discovered high affinity ER ligands, some of which are very selective for ERa, as well as basic side chain substituted analogs that are ERa-selective antagonists. This work complements our studies on the pyrazoles.

KEY RESEARCH ACCOMPLISHMENTS

Progress in Relation to the Statement of Work

The complete three year Statement of Work, presented in the original proposal of July 1996, is shown below. Our accomplishments are noted in *italics*:

ORIGINAL STATEMENT OF WORK

Project Period: <u>September 1, 1997– August 31, 2001 (3 years, with a one-year no-cost</u> extension)

Year 1

- Synthesis of representative members of Class I-VI ligands by solution phase methods. *completed*
- Isolate, purify, and fully characterize these members. completed
- Measure estrogen receptor binding affinity of representative members of Class I-VI ligands. *completed*
- Begin to adapt solution phase syntheses to solid phase. completed for the most promising series (pyrazoles)

Year 2

- Complete adaptation of solution phase synthesis to solid phase. completed for the most promising series (pyrazoles)
- Isolate and fully characterize representative members produced by solid phase synthesis. *completed for the most promising series (pyrazoles)*Determine yield and characterize impurities. *completed for the most promising series (pyrazoles)*
- Compare estrogen receptor binding of representative members of Class I-VI ligands prepared by solution vs solid phase. *completed for the most promising series (pyrazoles)*
- Begin synthesis of full Class I-VI libraries. completed for the most promising series (pyrazoles)
- Begin cell proliferation and cell-based transfection assays. completed for the most promising series (pyrazoles)

Year 3

- Complete synthesis of full Class I-VI libraries. completed for the most promising series (pyrazoles)
- Complete estrogen receptor assay of full libraries at two concentrations. *completed for the most promising series (pyrazoles)*

- Reassay the members with detectable estrogen receptor binding affinity by quantitative titration assay. *completed for the most promising series (pyrazoles)*
- Assay the members with high estrogen receptor binding affinity in the cell proliferation and cell-based transfection assays. *completed for the most promising series (pyrazoles)*
- Assay uterotrophic activity of the most promising members in rats. <u>In vivo assays have been</u> <u>completed through a collaboration with Wyeth-Ayerst.</u>

Please see **Summary** at the conclusion of the preceeding section that outlines additional accomplishments beyond those outlined in the original Statement of Work.

REPORTABLE OUTCOMES

- <u>Seven papers have been published, two</u> manuscripts have been accepted, and <u>one</u> has recently been submitted. These publications are attached as the appendix and are listed as **Publications No. 1-10**.
- Presentations have been made at <u>four</u> American Chemical Society meeting by the PI and principal co-workers, Shaun Stauffer <u>and Deborah Mortensen</u>, and by Stauffer at a meeting on solid phase synthesis. Numerous seminars given by the PI have included work developed under this project.
- A PCT patent has been submitted to the US Patent Office covering the 5-membered ring heterocyclic estrogens.
- Shaun Stauffer, my principal co-worker on this project, has completed his Ph.D. degree in the fall of 1999.
- <u>Deborah Mortensen, another major co-worker on this project, has completed her Ph.D.</u> <u>degree in the fall of 2000.</u>
- <u>After completing a very successful 2-year postdoctoral fellowship with Professor John</u> <u>Hartwig at Yale University, supported by a postdoctoral fellowship from the National</u> <u>Institutes of Health, Shaun Stauffer joined Merck. He had numerous offers from the best</u> <u>pharmaceutical industries</u>. The topic of Shaun's NIH postdoctoral fellowship derived from the experience he developed working on this project.
- On the basis of a presentation that she gave at the ACS meeting in San Francisco, Deborah Mortensen received two job offers. She is now working for Signal Pharmaceuticals in San Diego on a nuclear receptor project.

- Ying Huang, a postdoctoral fellow in my laboratory who worked on this project, has completed her studies and has moved to a job in new drug discovery at Schering-Plough in New Jersey.
- Rosanna Tedesco, a graduate student who has worked on this project, will complete her Ph.D. studies in October, 2000, and she has <u>moved to</u> a position in new drug discovery at SmithKline Beecham.
- Christopher Coletta and Zachary Aron, two undergraduate students who worked on this
 project for their senior research, have graduated with high honors, based in part on their
 excellent undergraduate theses. Chris is working as a B. S. Chemist in new drug discovery at
 Pfizer, in Connecticut, and Zachary is in a Ph.D. graduate program at the University of
 California, Irvine, working with Professor Larry Overman.

CONCLUSIONS [The Summary, presented earlier, is repeated for convenience]

Summary — Through the work we have engaged in on this project over the past three years, we have completed all of the goals of the original proposal, as summarized in the original Statement of Work. This has involved an exploration of the promise of all of the proposed acyclic amide and 5-membered ring heterocyclic systems, and the development of solid phase synthesis methods and large library synthesis and analysis. In addition, we have discovered in the process that some of the most promising heterocyclic ligands, the pyrazoles, have high affinity and potency selectivity for one of the estrogen receptor subtypes, ER α . Because the development of estrogen receptor subtype selective ligands is a very important issue and one of great current interest, we have devoted considerable efforts in years 2 and 3 to pursue this further. We have completed a careful structure-activity relationship study on the pyrazoles, obtaining ultimately a compound that is almost a completely specific activator of ER α , having essentially no activity on ER β at doses where it maximally stimulated ER α , and, based on these findings, we have developed a series of pyrazole basic side chain derivatives that hold considerable promise as potency selective antagonists of ERa. Such compounds could be used to block the action of estradiol through this receptor subtype without interfering with its effect on ERβ.

The work that we have done on this project has also spawned several related lines of inquiry that are currently being pursued in my laboratory with independent support. These include investigations of other 5-membered ring heterocyclic systems as estrogen receptor ligands, namely furans, thiophenes, and pyrroles, as well as carbocyclic analogs,

cyclopentadienes and related cyclopentadienones. In addition, based on the favorable results of the work under this project, we have begun to synthesize 6-membered ring heterocycles such as pyrazines, pyrimidines, and pyridazines as estrogen receptor ligands. This work is still in an early state of development, but already we know that we have compounds that show high affinity for the receptor and are therefore potentially quite interesting, because their structures are unique in the estrogen receptor field.

We certainly appreciate the support that has been provided to our research efforts in this area through this US Army grant. It has served as a stimulus to us to thrust our laboratory research into new directions that have proved to be very fruitful and potentially useful for the prevention and treatment of breast cancer.

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Publications, Patents, Meeting Abtracts, and Personnel Supported

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Publication No. 1:	B. E. Fink, D. S. Mortensen, S. R. Stauffer, Z. D. Aron, J. A.
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John A. Katzenellenbogen, August, 2001 Army Breast Cancer Annual Report

Personnel Supported: Shaun R. Stauffer Deborah Mortensen Ying R. Huang •

Scheme 1. Estrogens, Pure Antiestrogens, and Selective Antiestrogens - From the Original Proposal



Scheme 2. Structural Components of Selective Antiestrogens – From the Original Proposal



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Estrogen Receptor Ligand	Structural Motif	Combinatorial Analog (Class)
HO HEXESTROL	HO Notif Aanti-Bibenzyl	I. Diphenyl Carboxamides R1 R2 R2 R2 R2
Horizanoviten		II. Diphenyl Sulfonamides
HO CH'S HO CH'S Benzestrol	HO HO Motif BHomobibenzyi	III. Phenyl Benzylcarboxamides & Sulfonamides R_1 R_2 R_2 R_3 R_3 R_2 R_3 R_3 (or SO ₂ in place of CO)
и С С С С С С С С С С С С С С С С С С С		IV. Benzyl Phenylcarboxamides & Sulfonamides $R_1 \cup R_2$ $R_2 \cup R_3$ (or SO ₂ in place of CO)
Haloxifene		V. Pyrazoles
HO Hydroxy-Tarnoxifen	HO Good Motif Csyn-Bibenzyl	VI. Oxazoles, Thiazoles, and Imidazoles R1 HO HO R2
		(X = O, S, NR)

Scheme 3. Structural Motifs for Estrogen Receptor Ligands and Their Combinatorial Analogs – From the Original Proposal

Novel structural templates for estrogen-receptor ligands and prospects for combinatorial synthesis of estrogens

Brian E Fink, Deborah S Mortensen, Shaun R Stauffer, Zachary D Aron and John A Katzenellenbogen

Introduction: The development of estrogen pharmaceutical agents with appropriate tissue-selectivity profiles has not yet benefited substantially from the application of combinatorial synthetic approaches to the preparation of structural classes that are known to be ligands for the estrogen receptor (ER). We have developed an estrogen pharmacophore that consists of a simple heterocyclic core scaffold, amenable to construction by combinatorial methods, onto which are appended 3–4 peripheral substituents that embody substructural motifs commonly found in nonsteroidal estrogens. The issue addressed here is whether these heterocyclic core structures can be used to prepare ligands with good affinity for the ER.

Results: We prepared representative members of various azole core structures. Although members of the imidazole, thiazole or isoxazole classes generally have weak binding for the ER, several members of the pyrazole class show good binding affinity. The high-affinity pyrazoles bear close conformational relationship to the nonsteroidal ligand raloxifene, and they can be fitted into the ligand-binding pocket of the ER–raloxifene X-ray structure.

Conclusions: Compounds such as these pyrazoles, which are novel ER ligands, are well suited for combinatorial synthesis using solid-phase methods.

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Introduction

Estrogens are endocrine regulators of the vertebrate reproductive system that have important effects in many nonreproductive tissues as well (bone, liver, cardiovascular system, CNS and so on). Many estrogen pharmaceuticals, based on both natural and synthetic substances, have been developed as agents for regulating fertility, preventing and controlling hormone-responsive breast cancer, and menopausal hormone replacement. These substances display a spectrum of agonist to antagonist activity that can show remarkable tissue and cell selectivity [1].

The molecular target of estrogens is the estrogen receptor (ER), of which there are now known to be two subtypes, ER- α and ER- β , that have different patterns of tissue expression and somewhat different ligand-binding specificities [2,3]. ER is a transcription factor that binds to specific estrogen-response elements in the promoter region of estrogen-regulated genes and whose activity is modulated by the estrogen ligands [4]. The capacity of ERligand complexes to activate gene transcription is mediated by a series of co-regulator proteins [5]. These co-regulators have interaction functions that tether ER to the RNA polymerase II preinitiation complex and enzymatic activities to modify chromatin structure [6]. It is the fact that each cell and each gene presents to an ER(subtype)ligand complex a unique combination of these effector components-various estrogen-response elements and

co-regulators—that appears to underlie, in part, the cell and gene selectivity of various estrogens [7].

Among known ligands for ER, the natural estrogens are the simplest of the steroidal hormones, distinguished by their phenolic A-ring (Figure 1). Synthetic estrogens, especially those of nonsteroidal nature, generally retain a phenolic function (at least for those of high potency), but otherwise span a remarkable range of structural motifs that encompass simple acyclic core structures of various lengths and sizes, as well as a variety of ring-size fused and nonfused carbocyclic and heterocyclic systems [8-10]. It is clear from many decades of medicinal-chemistry investigations that minor changes in the structure and stereochemistry of these ligands can have profound effects on both their affinity and their biocharacter (i.e. the agonist versus antagonist balance in various tissues). Major efforts have been directed at optimizing ER ligand structure to obtain desired profiles of tissue selectivity, but, even so, the ideal profile for various uses has not yet been achieved [1,11,12].

As currently explored, ER ligands are, by and large, not well suited for synthesis using combinatorial approaches, because their preparation generally involves a series of carbon–carbon bond forming reactions that do not give uniformly high yields, nor are they well adapted to solidphase-synthesis methods. There are two examples of the preparation of estrogen combinatorial libraries on solid





Examples of ligands with high affinity for the estrogen receptor. In each case, the binding affinity relative to that of estradiol (100%) is given in parentheses. The compounds are grouped according to their activity in a standard rat uterine weight gain assay as (a) agonists, (b) mixed or selective agonist/antagonists or (c) pure antagonists.

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phase, both involving stilbene-like structures [13,14], but the application of combinatorial approaches for the preparation of ER ligands has, so far, been limited.

To expand possible combinatorial approaches to the synthesis of ER ligands, we have begun investigating simple structural motifs that might be used for the construction of molecules with high affinity for ER. The goal was to identify core structures that could be readily prepared by the types of simple condensation reactions that typify those used in solid-phase combinatorial approaches for the preparation of drug-like molecules, and from these to select ones that would support the development of high-affinity ligands for ER.

Here, we describe the investigation of prototype 1,2- and 1,3-azole systems as potential ligands for the ER. We examine the issue of whether ER ligands can be considered

simply as an assembly of a phenol unit together with 2–3 auxiliary peripheral groups linked together by a functionally inert core scaffold, or whether the core scaffold itself plays an integral role in ligand binding. In the process, we have discovered a new class of high-affinity ligands for ER, 4-alkyl-1,3,5-triarylpyrazoles, that bear an unexpected topological resemblance to the nonsteroidal estrogen raloxifene and show an interesting structure-binding affinity pattern.

Results and discussion

Structural motifs found in estrogen-receptor ligands and proposed heterocyclic surrogates

Selected examples of nonsteroidal ligands for the estrogen receptor are shown in Figure 1, together with an indication of their ER-binding affinity and their agonist (Ag) versus antagonist (Antag) character in a standard rat uterine weight gain assay. Collectively, these molecules exemplify a recognizable structural gestalt (Figure 2): a core structure (A) onto



Estrogen ligand pharmacophore model.

which are attached other, peripheral structural elements, a phenolic unit (B) that is always preserved, a second aromatic group (C) that is usually present, and another substituent (D) or two (D'), one of which might be aromatic. In the case of ER antagonists or mixed agonist/antagonists, one of the substituents generally contains a basic or polar function.

Comparisons of the various specific manifestations of this basic structure (Figure 1) suggest that, to achieve high ER-binding affinity, the peripheral substituents (B-D') need to be displayed in a certain geometric arrangement, so that they will be 'in register' with their corresponding subsites in the ligand-binding pocket in ER. It seems that this peripheral group 'display function' can be accomplished by using core elements that encompass a considerable structural variety. This raises the interesting question of whether the core element itself plays any direct role in ER binding or whether it serves merely as an inert molecular scaffold whose function is simply to display these peripheral elements with appropriate topology. If the latter is true, it should be possible to replace the core scaffold with a variety of other units, providing they also are able to display the peripheral elements with the appropriate geometry. Some of these core scaffolds, namely small-ring heterocycles, could be assembled by facile condensation reactions from simpler components, a situation that is favorable for the development of large chemical libraries of related compounds by combinatorial synthesis approaches.

In Table 1 we have outlined two (out of many possible) manifestations of this conceptual approach, based on the incorporation of certain substructural motifs into 1,2- and

Structural motifs found in ER ligands and proposed surrogates. Structural motif Combinatorial analog (class) Estrogen-receptor ligand 3,5-Diaryl pyrazoles Benzestro Motif A. Homobibenzyl² 3,5-Diaryl isoxazoles Raloxifene 2,4-Diaryl imidazoles, thiazoles, oxazoles NMe₂ Hydroxytamoxifen HO 4,5-Diaryl imidazoles, oxazoles, thiazoles Motif B. Bibenzyl*

*These structural motifs are meant to highlight alternative atom connections between the phenol and a second aromatic substituent, without specific consideration of conformational factors. Structural motifs found in ER ligands and proposed surrogates.

Table 1



Figure 3

1,3-azole systems. Here, the homobibenzyl motif A, exemplified in the nonsteroidal ligands benzestrol and raloxifene, is represented in various 3,5-diaryl-1,2-azoles (pyrazoles and isoxazoles) and 2,4-diaryl-1,3-azoles (imidazoles, thiazoles and oxazoles). Similarly, the bibenzyl motif B is represented in various 4,5-diaryl-1,3-azoles. In each case, the diazole N,N-systems (namely pyrazoles and imidazoles) can accommodate up to four peripheral substituents, whereas the N,O- and N,S-heterocycles (oxazoles, isoxazoles and thiazoles) are limited to three substituents.

Synthesis of representative diaryl and triaryl 1,2- and 1,3-azoles as potential ligands for the estrogen receptor *Imidazoles*

The synthesis of representative symmetrical members of the imidazole class and their N-alkyl analogs was accomplished by a well-precedented approach [15] shown in Figure 3. Refluxing 4,4'-dimethoxybenzil (1) in formamide in the presence of *para*-formaldehyde afforded the 4,5-disubstituted imidazole 2 [16], which upon deprotection with BBr₃ in CH₂Cl₂ afforded imidazole 3 in good yield. A similar reaction using 4-methoxybenzaldehyde afforded the 2,4,5- tri-substituted imidazole 4 [17–19] in good yield. To prepare tetra-substituted systems, the sodium salt of imidazole 4 was alkylated with ethyl, propyl and butyl iodide, and then deprotected to afford free phenols **6a–d**.

Two additional, unsymmetrical, imidazoles were synthesized as outlined in Figure 4. The top reaction sequence illustrates the synthetic approach to N-ethyl imidazole **12** Reaction of 4-methoxy-deoxybenzoin (**7**) [20] with bromine

Figure 4



Synthesis of imidazoles 12 and 17.



and a trace of AlCl₃ in Et₂O gave α -bromoketone 8 [21], which, upon reaction with sodium azide in acetone, afforded the corresponding azide 9. The azido-ketone 9 was treated with one equivalent of Et₃N and imine 10 in tetrahydrofuran (THF). Removal of solvent and excess Et₃N followed by treatment of the crude intermediate 2,5-dihydro-2-hydroxyimidazole with TFA in CH₂Cl₂, according to the procedure of Patonay and Hoffman [22], resulted in the formation of *N*-ethyl imidazole 11. Deprotection with BF₃•SMc₂ in CH₂Cl₂ produced imidazole 12 in good yield.

The synthesis of N-aryl substituted imidazole 17 is also shown in Figure 4. Refluxing 4'-methoxy- α -bromobutyrophenone (13) with *p*-anisidine in acctone gave the α -amino-ketone 14, which was converted into the benzamide 15 upon reaction with benzoyl chloride and base. Cyclization with ammonium acetate in refluxing acetic acid afforded the 1,2,4,5 tetra-substituted imidazole 16, which upon deprotection with BF₃•SMe₂ in CH₂Cl₂ produced the free phenol 17.

Thiazoles

The synthesis of representative thiazoles is shown in Figure 5. Thioamide 19, derived from 4-methoxybenzonitrile (18) [23], was condensed with 4'-methoxy- α -bromoacetophenone (20) or 4'-methoxy- α -bromobutyrophenone (13) in refluxing DMF to give high yields of the 2,4-disubstituted thiazole 21a [24] or 2,4,5- tri-substituted thiazole 21b, respectively. Deprotection with BBr₃ afforded moderate yields of the free phenols 22a and 22b.

Oxazoles

Two representative oxazoles were synthesized as shown in Figure 6. Reaction of the lithium anion of dithiane 23with *p*-methoxybenzyl bromide gave the alkylated product 24, which upon hydrolysis afforded 4'-methoxy-deoxybenzoin (25) [25] in excellent yield. Conversion to the

Figure 6



Synthesis of oxazoles 29 and 31.

bromide 26 [26] and azide 27 was accomplished as described for analogous compounds 8 and 9 above. The azido-ketone 27 was then treated with one equivalent of Et_3N and *p*-anisaldehyde, and then with TFA to afford oxazole 28 [27]. Oxazole 30 resulted from the condensation of bromo-ketone 26 with *p*-methoxybenzamide in refluxing toluene, analogous to the thiazole synthesis discussed above. Deprotection of 28 and 30 with $BF_3 \cdot SMe_2$ gave oxazoles 29 and 31, respectively.

Pyrazoles

The synthesis of the pyrazoles involves the condensation of a hydrazine with a 1,3-diketone [28]. Using the method of Beak and coworkers [29], we obtained 1,3-diketone **33** from the reaction of the methyl thioester **32** and lithium tetramethylpiperidide in good yield (Figure 7). Condensation of the diketone with hydrazine hydrochloride or N-substituted hydrazine hydrochlorides in refluxing DMF/ THF (3:1) afforded the 3,5-disubstituted pyrazole **34a** or 1,3,5- tri-substituted pyrazoles **34b–d**; yields were higher with aryl-substituted hydrazines than with hydrazine itself. Deprotection of **34a–d** with BBr₃ afforded the free phenols **35a–d** in moderate yield.

The introduction of a 4-alkyl substituent was accomplished through the alkylation of diketone **33** with TBAF and ethyl iodide to afford **36** in moderate yield [30,31]. Attempts to increase the yield of this alkylation were





unsuccessful. Conversion of diketone **36** to the corresponding pyrazoles was accomplished as with the unsubstituted case, to afford pyrazoles **38a–d**.

Isoxazoles

The preparation of a single isoxazole is shown in Figure 8 [32]. Double deprotonation of the ketoxime **39** derived from 4-methoxyacetophenone with nBuLi, followed by addition of methyl 4-methoxybenzoate afforded the 3,5-disubstituted isoxazole **40** in low yield [33]. Deprotection with BBr₃ afforded the free phenol **41** in moderate yield [34].

Estrogen-receptor binding

The binding affinities of the heterocycles prepared above for the estrogen receptor are shown in Tables 2–4, organized according to heterocyclic core structure. The binding values were obtained from a competitive radiometric binding assay, using [³H]estradiol as the tracer, dextran-coated charcoal

Figure 8



to adsorb free tracer and lamb uterine cytosol as a source of ER. The values are expressed as relative binding affinities (RBA), with estradiol having an affinity of 100% [35]. In replicate assays, these values are reproducible with a coefficient of variation of 30% (K.E. Carlson and J.A.K., unpublished observations).

Imidazoles, oxazoles and thiazoles

The receptor-binding data for the imidazole series are shown in Table 2. Although the members of this series have rather low affinity, there is an increase in RBA with the addition of alkyl substituents at the 1-position (R_3 ; **6a-d**); this trend reaches a maximum for propyl **6c**, reversing for the butyl substituent **6d**. Such trends, where affinity

Table 2

Estrogen receptor binding data for imidazoles 3, 6a-d, 12 and 17.

			R ³ ∕R ⁴		
Compo	ound R ¹	R ²	R ³	R⁴	RBA*
3	4′-HO-C ₆ H ₄	4′-HO-C ₆ H₄	н	н	< 0.001
6a	4'-HO-C ₆ H ₄	4'-HO-C ₆ H ₄	Н	4'-HO-C ₆ H ₄	0.007
6b	4'-HO-C ₆ H ₄	4'-HO-C ₆ H ₄	C_2H_5	4'-HO-C ₆ H ₄	0.38
6c	4'-HO-C ₆ H ₄	4'-HO-C ₆ H ₄	C_3H_7	4'-HO-C ₆ H ₄	0.62
6d	4'-HO-C ₆ H ₄	4'-HO-C ₆ H ₄	C₄H ₉	4'-HO-C ₆ H ₄	0.17
12	C ₆ H ₅	4'-HO-C ₆ H ₄	C_2H_5	4'-HO-C ₆ H ₄	0.25
17	4'-HO-C ₆ H ₄	C ₂ H ₅ 4'	-HO-C ₆ H	H ₄ C ₆ H ₅	0.37

*RBA, relative binding affinity (estradiol = 100%).

Table 3

29

31

Estrogen receptor binding data for thiazoles 22a, 22b and oxazoles 29 and 31.						
HO N R ¹						
Compound	Х	R ¹	R ²	RBA		
22a	S	4′-HO-C ₆ H₄	Н	0.018		
22b	S	4'-HO-C ₆ H ₄	C_2H_5	0.041		

4'-HO-C₆H₄

 C_6H_5

 C_6H_5

4'-HO-C₆H₄

< 0.001

0.027

0

0

increases with substituent size up to a point, are well known both in steroidal systems (11 β - and 16 α -substituents) [36] and in other non-steroidal ligand series (such as 2-phenylindoles [37], tetrahydrochrysenes [38] and so on), and probably represent the filling of preformed pockets of limited volume in the receptor by these substituents [37]. The principal difference in binding, however, is between the tetra-substituted imidazoles (6b-d, 12 and 17) and the di- or tri-substituted imidazole (3 and 6a), the tetra-substituted ones having much higher affinity. There is little difference in binding between imidazoles 12 and 17, which have a different arrangement of nitrogen atoms in the heterocyclic core but display their four substituents in an identical fashion (Figure 9). The overall low binding affinity of the imidazoles as a class might be the result of the high inherent polarity of this heterocyclic system as compared with the pyrazoles, reflected by their higher chromatographic polarity in both normal and reversed phases systems. It is also of note that the dipole moment for imidazole is very

Table 4

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Compound	Х	R	R ¹	RBA	
35a	Ν	н	н	0.009	
35b	Ν	C_6H_5-	н	0.028	
35c	Ν	C ₆ H ₅ CH ₂ -	н	< 0.007	
35d	Ν	4'-HO-C ₆ H ₄	н	0.059	
38a	Ν	Н	C_2H_5	0.015	
38b	Ν	C ₆ H ₅ -	C_2H_5	14.0	
38c	Ν	C ₆ H ₅ CH ₂ -	C_2H_5	0.150	
38d	Ν	4'-HO-C ₆ H ₄	C_2H_5	19.0	
41	0	-	C_2H_5	0.006	

Estrogen receptor binding affinity data for pyrazoles and an





Comparisons between ring pyrazoles (**38b**, **38d**) and imidazoles (**6b**, **12** and **17**).

large, 5.56 D [39], and this might be unfavorable for binding to the estrogen receptor.

Table 3 shows the binding data for the two thiazoles and oxazoles. Although affinities are again very low, the more highly substituted thiazole again has the higher affinity (22a compared with 22b). The oxazole 29 has undetectable affinity for ER. The isomer 31, however, does have measurable, albeit low, binding. In contrast to imidazoles, thiazoles and oxazoles do not have very high dipole moments [39]; so overall polarity is not likely to be the cause of their low ER binding affinity, although heteroatom orientation appears to play a role (29 compared with 31). In the imidazole series, the compounds with the highest affinities were all tetra-substituted, however. As it is only possible to trisubstitute a thiazole or oxazole, this core structure might be unable to present sufficient peripheral substituents to afford ligands with good ER binding affinities, at least as far as we have investigated.

The low binding affinities of the imidazoles, thiazoles and oxazoles are disappointing, although not surprising, considering the relatively poor affinity of the most closely related benzothiazole reported by von Angerer [40] (Figure 1). The sparsely substituted monocyclic or polycyclic aromatic systems are also expected to be rather



Comparison of the estrogen receptor binding affinity of four related pyrazoles with increasing substitution.

planar. It is generally agreed that good ligands for the estrogen receptor need to have some degree of 'thickness' in the central portion of the ligand [38]. When alkyl substituents are added to either the imidazoles or thiazoles, their RBA increases (Tables 2,3). This increased binding could be due to an increase in steric bulk around the central portion of the molecule, the result, in part, of a twisting of some of the aromatic substituents (see below) or to an increase in lipophilicity. Regardless, the effects are not great, and in general the 1,3-azoles seem to present some special challenges that might best be investigated further by combinatorial approaches.

Pyrazoles and isoxazoles

The RBA data for the 1,2-azoles are presented in Table 4. Immediately apparent is the relatively high binding affinity of pyrazoles **38b** and **38d**. An interesting comparison can be made among compounds **35a**, **35b**, **38a** and **38b** (Figure 10). The disubstituted progenitor **35a** has very low affinity; addition of a third substituent, 1-phenyl in **35b** or 4-ethyl in **38a**, causes only a twofold or threefold increase in binding affinity, respectively. By contrast, addition of the fourth substituent (to give **38b**) causes either an 900- or 500-fold increase in binding affinity, respectively. Clearly, this is not additive behavior—two groups that each alone raise binding affinity twofold and threefold, together raise binding not sixfold but 1600-fold. This suggests that to achieve high binding affinity there needs to be a detailed





Ab initio calculated conformations for 35a, 35b, 38a and 38b.

and proper match between the peripheral substituents and several subsites on the receptor, and in the azole systems we have explored, it appears that this requires a tetra-substituted ring (see below). Consistent with this is the low affinity of the isoxazole **41**, whose affinity is similar to the most closely related tri-substituted pyrazole **38a**.

There are other interesting trends in the pyrazole series: replacement of the N-phenyl substituent (38b) with an N-benzyl group (38c) causes a dramatic 100-fold reduction in binding. Both of these compounds are tetra-substituted pyrazoles, and they contain the homobibenzyl motif A that was considered to be an important factor for receptor binding (as do all of the other compounds in Table 1). The decrease in binding affinity in 38b compared with 38c again suggests the need for a detailed match between ligand substituents and receptor subsites: the extra 'kink' in the benzyl substituent in 38c might be repositioning the peripheral substituents in a less favorable geometry. The addition of a hydroxyl group at the *para* position of the N-phenvl substituent (compound 38d compared with 38b) causes a minor increase in binding, indicating that polarity is well tolerated in this region of the receptor.

Structural comparisons between high-affinity pyrazole ligands and other nonsteroidal ligands

The conformation of pyrazoles **35a,b** and **38a,b** was determined by *ab initio* calculations at the 3-G21* level (Figure 11). The action of A-strain is evident in these structures: even in the disubstituted system **35a**, the 5-phenyl group is twisted ~60° out of the plane; this twisting increases as the third (**35b**, **38a**) and fourth (**38b**) substituents are added to the pyrazole ring, so that in **38b** the 5-phenyl substituent is nearly perpendicular to the pyrazole core. In this conformation **38b** resembles the general propeller-type conformation of the triarylethylene nonsteroidal estrogens such as tamoxifen [41]. Although the comparison of the conformations of tamoxifen with pyrazole **38b** might give some idea of the reason **38b** binds to ER with high affinity, it does not adequately explain why **35a**, **35b** and **38a** have so much lower binding affinities. When the latter three pyrazoles are compared with **38b**, it is apparent that there is little difference in conformation that could account for the large discrepancy in binding within the series. The reasons for the marked changes in RBA in response to very small changes in structure, noted especially for the pyrazole series, are therefore not intuitively obvious from an examination of the structure of the ligands alone, although they are not entirely unexpected on the basis of the behavior of other nonsteroidal ligands and the detailed fit of these congeners in the ligandbinding pocket of the estrogen receptor (see below).

Analysis of the X-ray structure of the estrogen-receptor ligand-binding domain complexed with estradiol and the nonsteroidal ligand raloxifene

The explanation that was previously proposed — that highaffinity binding to ER derives from a proper match between the peripheral substituents on the ligand and their complementary binding regions on the receptor—can now be considered in some detail, because, recently, the X-ray structures of the estrogen receptor- α bound to estradiol and the nonsteroidal ligand raloxifene have been reported [42].

The ligand-binding pocket in the ER-estradiol structure has a volume of approximately 450 Å³, which is ~200 Å³ larger than the volume of estradiol [43] (Figure 12a). As a result, there is a large hydrophobic space around the central portion of the binding pocket, especially in the regions corresponding to the 11 β position and, to a lesser degree, 7α position (Figure 12a), which is consistent with the tolerance that ER shows for binding steroids with large substituents at these positions [36]. A view of the ERraloxifene structure is shown in Figure 12b. The core of this ligand is oriented in the same manner as estradiol and occupies much of the same space in the binding pocket, but the benzoyl substituent projects outward, askew of the ligand core, with the piperidinyl sidechain extending into an upper hydrophobic region, which is much more open due to the displacement of helix 12 [42].

Comparison between raloxifene and the tetra-substituted pyrazole (38b)

The structure of the high-affinity pyrazole (38b) can be overlayed onto the structure of raloxifene (Figure 13a). In such a superposition, it is evident that the centroids of all three aromatic rings lie quite close to one another. This structural alignment can be used to place the pyrazole 38b into the ligand-binding pocket of the ER-raloxifene structure (Figure 13b). With a minimal, energetically reasonable rearrangement of nearby residues (see Figure 13 legend), this ligand can fit quite comfortably in the raloxifene pocket (compare Figure 13b with Figure 12b).





Ligand binding pockets for (a) estradiol and (b) raloxifene. Structures were prepared from the crystallographic coordinates [43] by generating a solvent-accessible surface for the protein (green-blue dot surface) and for the ligand (purple).

In the arrangement shown in Figure 13b, the two hydroxyl groups are positioned in such a manner that they could engage the same protein hydrogen-bonding partners as do the corresponding hydroxyl groups in raloxifene; the 3-(p-hydroxyphenyl) substituent of **38b** is mimicking the estradiol A-ring surrogate of raloxifene (i.e. the fused phenol of the benzothiophene unit) and the 5-(p-hydroxyphenyl) group of **38b** is mimicking the pendant p-hydroxyphenyl group at position 2 of the benzothiophene (compare with Figure 12b). The 1-phenyl group of **38b** overlies the benzoyl arene of raloxifene and projects into channel that exists in roughly the 11 β direction in the ER-raloxifene





(a) Overlay of pyrazole 38b with raloxifene and (b) pyrazole 38b in the binding pocket of the ER-raloxifene structure. (a) The overlay of the two ligands was obtained by a least-squares multifitting seven atoms in each molecule (rms=0.924 Å): in each of the three benzene rings, the two atoms selected were the ones at the site of attachment and the ones *para* to this site; in addition, the *p*-hydroxy group on the 5-phenyl substituent in pyrazole 38b was overlaid with the benzothiophene hydroxyl group.
(b) Pyrazole 38b was pre-positioned into the ER-raloxifene crystal structure [42] from the overlay shown in (a). From this structure additional docking studies using FlexiDock (Tripos, St Louis, MO) followed by a three-step minimization using the TRIPOS Forcefield were conducted to afford the final model (see the Materials and methods section). This minimization caused only small changes in the protein and ligand, but reduced the ligand-protein interaction energy to a reasonable level.

structure. The 4-ethyl substituent on pyrazole **38b** projects outwards from the heterocycle from a position that corresponds roughly to the sulfur atom of the benzothiophene ring system, and extends into an open pocket in the ER.

When analyzed in this manner, one can appreciate that the pyrazole core can display these four substituents in a manner congruent with the same regions in the binding pocket of the estrogen receptor that accommodate corresponding portions of raloxifene. In addition, the high cost in binding affinity that results from the absence of either the 1-phenyl or the 4-ethyl substituent (compare with Figure 10) suggests that proper registration of each of the four peripheral substituents into its appropriate binding subsite is supported by the interaction of the other three. By this analysis, it is therefore not surprising that all of the other heterocyclic systems that were only di-substituted or tri-substituted were low affinity ligands, at least with the substituents we have thus far investigated. The low-affinity of the tetra-substituted imidazoles, however, most likely derives from their high polarity, as noted (see below).

The importance of core structural element in ligand binding: passive or active?

The question raised initially-does the core scaffold in these novel ER ligands play only a passive role in their binding, merely displaying the peripheral substituents in an appropriate topology, or is its role more active or functional?-can be answered reasonably definitively from the results we have obtained so far. Clearly, with the substituents we have examined, high-affinity binding was obtained only with those azoles that afforded the possibility of tetra-substitution. But of these, the 1,2-diazoles (pyrazoles) and the 1,3-diazoles (imidazoles), only the pyrazoles gave good binding. Although there are not many direct comparisons that can be made between these two diazole systems, pyrazole 38b and imidazoles 12 and 17 are, in fact, just ring nitrogen isomers of one another, having otherwise identical peripheral groups and functionality (Figure 9). The same is true for pyrazole **38d** and imidazole **6b**. In both cases, however, the pyrazole partner binds to ER with ~30-36-fold higher affinity than the isomeric imidazole(s). This would suggest that the core structure does play more than a passive role in ER binding, although, as noted before, the high polarity and significant dipole moment of the imidazole might be the principal reason for the difference in this case. The issue of the functional role of the core scaffold in ER binding needs to be investigated further in these and other heterocyclic systems.

Significance

Compounds with a remarkable variety of structures bind with high affinity to the estrogen receptor (ER), and many nonsteroidal ligands have been prepared in the search for agents that have improved tissue-selectivity profiles. The application of combinatorial approaches to the development of selective ER ligands, however, is still in a state of infancy.

Based on a simple pharmacophore model consisting of a core scaffold and various peripheral groups we have designed systems that display the peripheral groups on simple heterocyclic core systems, which are readily prepared by simple condensation reactions. Here, we describe the synthesis and ER-binding affinity of various substituted 1,2- and 1,3-azoles.

No significant binding was found for members of the imidazole, thiazole or isoxazole classes, and this is rationalized either by the inherent polarity of these compounds (imidazoles) or by their inability to carry a sufficient number of the types of peripheral substituents we have explored so far (thiazoles, oxazoles and isoxazoles). Several members of the pyrazole class did show good binding affinity, however, the best being a tetra-substituted pyrazole **38d**. Both **38b** and **38d** bear an unexpectedly close conformational relationship to the nonsteroidal ligand raloxifene.

Compounds such as **38b** and **38d** are well suited to combinatorial synthesis using solid-phase methods. The large differences in binding affinity that result from small structural changes suggest that a thorough investigation of many possible combinations of core structures and peripheral substituents will be needed to identify novel high-affinity ligands for the ER that can be evaluated for their selective biological activity. The solid-phase combinatorial synthesis of pyrazole libraries is currently underway and has yielded other high-affinity ligands for the estrogen receptor (S.R.S. and J.A.K., unpublished observations). Some of these heterocycles have also shown intriguing biological activity [43].

Materials and methods

General methods

All reactions using water- or air-sensitive reagents were conducted under an Ar atmosphere with dry solvents. Solvents were distilled under N₂ as follows: CH_2Cl_2 from CaH_2 , tetrahydrofuran (THF) from sodium benzophenone ketyl, dimethylformamide (DMF) from MgSO₄, and hexanes from CaSO₄. Triethylamine was distilled over CaH₂. All other reagents were purchased from commercial suppliers and used without further purification. Reactions were all monitored using thin-layer chromatography (TLC), performed on 0.25 mm silica gel glass plates containing F-254 indicator. Visualization on TLC was achieved by UV light (254 nm), iodine vapors, or phosphomolybdic acid indicator. Flash chromatography was performed using Woelm 32–63 μ m silica gel packing unless otherwise noted.

¹H NMR and ¹³C NMR spectra were recorded on a Varian U400, Varian U500 or Varian INOVA 750. NMR spectra chemical shifts (δ) are reported in parts per million downfield from TMS and referenced with either TMS internal standard for CDCl₃, acetone-*d*₆, MeOD-*d*₄, or DMSO-*d*₆ solvent peak. NMR coupling constants are reported in Hertz. Electron ionization (EI) spectra were obtained using a Finnigan---MATCH5 spectrometer at 70 eV. Fast atom bombardment (FAB) were recorded on a VG ZAB-SE spectrometer. High-pressure liquid chromatography (HPLC) was performed on a SpectraPhysics P100 solvent delivery system with ultraviolet detection at 254 nm. Elemental analysis was performed by the Microanalytical Service Laboratory at the University of Illinois. All characterized compounds are chromatographically homogeneous.

Relative binding affinities

Assays were performed as reported previously [44] using lamb uterine cytosol diluted to approximately 1.5 nM of receptor, which was incubated with buffer of several concentrations of unlabeled competitor together with 10 nM [³H]estradiol for 18–24 h. Free ligand was removed by adsorption onto dextran-coated charcoal. Unlabeled competitors were prepared in 1:1 DMF:TEA to ensure solubility.

Molecular modeling and docking studies

Solvent-accessible surfaces were generated (Figures 12 and 13) using the QCPE Connolly Program module (Indiana University) in Sybyl 6.5 (Tripos, St. Louis, MO). Figure 13b: the pre-positioned pyrazole **38b** was used for additional docking studies using the Tripos FlexiDock module. Both hydrogen-bond donors and acceptors within the pocket surrounding the ligand and the ligand itself in addition to select rotable torsional bonds were defined in order to afford an optimal docked-structure prior to molecular mechanics minimization. With the protein backbone held rigid, the ligand and the protein residues within 8 Å of the ligand were then minimized using a step-wise approach: first torsional bonds about the ligand were minimized holding the receptor fixed, followed by minimization of the receptor holding the ligand fixed, and then minimization of both the ligand and receptor. Minimizations were done using the TRIPOS Forcefield (as implemented in the program Sybyl) with the Powell gradient method and default settings (final RMS < 0.05 kcal/mol-Å).

Representative chemical synthesis

4,5-*Di*(4-*methoxyphenyl*)-1*H*-*imidazole* (2). To 4,4'-dimethoxybenzil (1) (2.0 g, 7.4 mmol) and *p*-formaldehyde (1.0 g, 11.1 mmol) was added formamide (50 ml). The bright yellow suspension was heated to reflux (220°C) for 2 h. The reaction mixture was then cooled to room temperature then to 0°C. The crystals that formed were filtered and recrystallized from EtOAc to afford 2 (2.4 g, 86%). mp 183–184°C (lit [16] mp 183–184°C); ¹H NMR (400 MHz, MeOH-*d*₄) δ 7.64 (s, 1H), 7.44 (d, 4H, *J* = 7.50), 6.89 (d, 4H, *J* = 7.50), 3.79 (s, 6H); ¹³C NMR (100 MHz, MeOH-*d*₄) δ 158.4, 135.2, 129.1, 128.9, 122.6, 114.2, 55.3.

General demethylation procedure using BBr₃. To a stirring solution of the methyl-protected heterocycle (1 equiv) in CH_2Cl_2 at $-78^{\circ}C$ was added a solution of BBr₃ (4–5 equiv) as a 1N solution in CH_2Cl_2 . The reaction were allowed to warm to room temperature and stirred for 18 h. After quenching with H₂O, the layers were separated and the aqueous layer extracted with EtOAc (3 × 5 ml). The combined organic layers were dried over Na₂SO₄, fitered and concentrated to afforded the crude phenols. Flash chromatography afforded the demethylated products.

4,5-*Di*(4-*hydroxyphenyl*)-1*H*-*imidazole* (3). Imidazole 2 (100 mg, 0.35 mmol) afforded 3 (52 mg, 59%) by the general BBr₃ demethylation procedure. ¹H NMR (400 MHz, CDCl₃) δ 14.56 (br s, 1H), 9.93 (br s, 1H), 9.24 (s, 1H), 7.24 (d, 4H, *J* = 8.47), 6.82 (d, 4H, *J* = 8.40); MS (FAB) *m/z* (relative intensity, %) 253.1 (MH⁺ 24), 169.2 (100).

2,4,5-Tri(4-methoxyphenyl)-1H-imidazole (4). A suspension of 4,4'-dimethoxybenzil (1) (4.0 g, 15 mmol) and *p*-anisaldehyde (20 ml, 164 mmol) and formamide (100 ml) was heated to reflux (220°C) for 2 h, during which time the reaction mixture became homogeneous. The reaction was then cooled to 0°C and the precipitated product 4 was filtered. The light yellow powder was recrystallized from MeOH/H₂O to afford 3.80 g of 4 [19] (66%). mp 89–91°C (lit [19] mp 88–94°C). ¹H NMR (400 MHz, Acetone-*d*₆) δ 7.98 (d, 2H, *J*=8.88), 7.42 (d, 4H, *J*=8.52), 7.01 (d, 2H, *J*=8.83), 6.92 (br s, 4H), 3.79 (s, 3H), 3.75 (s, 6H); ¹³C NMR (100 MHz Acetone-*d*₆) δ 162.4, 159.9, 158.9, 145.4, 131.0, 129.0, 128.9, 126.6, 132.8, 114.0, 113.7, 113.6; MS (EI, 70 eV) *m/z* (relative intensity, %) 386.2 (M⁺, 100), 371 (30), 280 (100), 265 (30); HRMS calc'd for C₂₁H₁₆N₂O₃: 345.123800, found: 345.123918.

General *N*-alkylation procedure for imidazoles. A solution of imidazole 4 (200 mg, 0.52 mmol) in THF (10 ml) and DMF (1.5 ml) was cooled to 5°C. NaH (31 mg, 0.78 mmol) was added as 60% dispersion in mineral oil. The reaction mixture was warmed to room temperature for 1 h and respective alkyl halide (0.04 ml, 0.62 mmol) was added. The resulting suspension was heated to reflux for 12 h, then cooled to room temperature. The light precipitate was filtered, and the filtrate was concentrated under vacuum to a yellow solid which was flashed on silica (30% EtOAc/Hexanes) to afford alkylated products **5b-d** in 80–90% yields.

1-Ethyl-2,4,5-tri(4-methoxyphenyl)-imidazole (**5b**). ¹H NMR (400 MHz, CDCl₃) δ 7.60 (AA'XX', 2H, J_{AX} = 8.88, J_{AA} = 2.51), 7.46 (AA'XX', 2H,
$\begin{array}{l} J_{\rm AX} \approx 8.97, \, J_{\rm AA} = 2.56), \, 7.32 \, ({\rm AA'XX'}, \, 2{\rm H}, \, J_{\rm AX} = 8.88, \, J_{\rm AA} = 2.56), 7.00 \\ ({\rm AA'XX'}, \, 2{\rm H}, \, J_{\rm AX} = 8.88, \, J_{\rm AA} = 2.51), \, 6.99 \, ({\rm AA'XX'}, \, 2{\rm H}, \, J_{\rm AX} = 8.88, \, J_{\rm AA} = 2.56), \, 6.74 \, ({\rm AA'XX'}, \, 2{\rm H}, \, J_{\rm AX} = 8.97, \, J_{\rm AA} = 2.56), \, 3.87 \, ({\rm q}, \, 2{\rm H}, \, J = 7.32), \, 3.87 \, ({\rm s}, \, 3{\rm H}), \, 3.85 \, ({\rm s}, \, 3{\rm H}), \, 1.00 \, ({\rm t}, \, 3{\rm H}, \, J = 7.14); \, ^{13}{\rm C} \, {\rm NMR} \\ (100 \, {\rm MHz}, \, {\rm CDCl}_3) \, \delta \, 160.0, \, 159.7, \, 158.1, \, 146.8, \, 137.2, \, 132.4, \, 130.5, \, 127.9, \, 127.5, \, 123.8, \, 123.7, \, 114.5, \, 114.0, \, 113.5, \, 55.3, \, 55.2, \, 55.1, \, 39.5, \, 16.2. \end{array}$

2,4,5-*Tri*(4-*hydroxyphenyl*)-1*H*-*imidazole* (**6***a*). According to the general BBr₃ demethylation procedure above, imidazole **4** (3.0 g, 7.8 mmol) afforded **6***a* as a green-orange solid that darkened upon exposure to air (1.8 g, 68%). mp 203–205°C; ¹H NMR (400 MHz, Acetone-*d*₆) δ 7.93 (AA'XX', 2H, *J*_{AX} = 8.97, *J*_{AA} = 2.47), 7.40 (AA'XX', 4H, *J*_{AX} = 8.60, *J*_{AA} = 2.47), 6.89 (AA'XX', 2H, *J*_{AX} = 8.97, *J*_{AA} = 2.47), 6.80 (AA'XX', 2H, *J*_{AX} = 8.60, *J*_{AA} = 2.47); ¹³C NMR (100 MHz, Acetone-*d*₆) δ 157.9, 156.4, 145.9, 129.0, 126.8, 124.0, 123.4, 121.3, 115.0, 114.7; MS (FAB) *m/z* (relative intensity, %) 345.1 (M+H⁺, 10), 353 (10), 169 (100); HRMS calc'd for C₂₁H₁₆N₂O₃: 345.123800, found: 345.123918.

1-*Ethyl-2,4,5-tri(4-hydroxyphenyl)-imidazole* (6b). According to the general BBr₃ demethylation procedure above, imidazole 5b (185 mg, 0.46 mmol) afforded **5b** (107 mg, 62%). mp 150–153°C; ¹H NMR (400 MHz, Acetone-*d*₆) & 7.51 (AA'XX', 2H, *J*_{AX} = 8.52, *J*_{AA} = 2.42), 7.34 (AA'XX', 2H, *J*_{AX} = 8.71, *J*_{AA} = 2.39), 7.24 (AA'XX', 2H, *J*_{AX} = 8.63, *J*_{AA} = 2.41), 6.97 (AA'XX', 2H, *J*_{AX} = 8.58, *J*_{AA} = 2.31), 6.90 (AA'XX', 2H, *J*_{AX} = 8.63, *J*_{AA} = 2.41), 6.97 (AA'XX', 2H, *J*_{AX} = 8.58, *J*_{AA} = 2.31), 6.90 (AA'XX', 2H, *J*_{AX} = 8.75, *J*_{AA} = 2.31), 6.64 (AA'XX', 2H, *J*_{AX} = 8.96, *J*_{AA} = 2.43), 3.93 (q, 2H, *J* = 7.19), 0.98 (t, 3H, *J* = 7.12); ¹³C NMR (100 MHz, Acetone-*d*₆) & 158.2, 157.9, 155.9, 146.5, 136.8, 132.5, 130.3, 127.9, 127.8, 126.6, 122.5, 122.4, 116.0, 115.4, 114.8, 17.9, 15.4; MS (FAB) *m/z* (relative intensity, %) 372.1 (MH⁺, 100), 343 (15), 275 (10), 214 (25), 162 (30), 148 (30); HRMS calc'd for C₂₃H₂₀N₂O₃: 372.147251, found: 372.147393.

1-Ethyl-2,5-(4-methoxyphenyl)-4-phenyl imidazole (11). Azido-ketone 9 (50.0 mg, 0.187 mmol) and imine 10 (92.0 mg, 0.564 mmol) were dissolved in THF (15 ml). Et_3N (29.0 $\mu\lambda,$ 0.208 mmol) was added via syringe and reaction stirred at room temperature for 48 h. The reaction mixture was then poured into H₂O and extracted with CH₂Cl₂, organic fractions were pooled, dried over Na2SO4, filtered and solvent removed under reduced pressure. The intermediate, 2,5-dihydro-2-hydroxyimidazole, used in next step without further purification or characterization, was taken up CH₂Cl₂ (10 ml). Solution was cooled to 0°C and TFA (14.4 µl, 0.187 mmol) was added via syringe. Reaction stirred at 0°C for 36 h. The mixture was diluted with CH2Cl2 (10 ml) and washed with H₂O, sat. NaHCO₃, and sat. NaCl successively. The organic fraction was dried over Na2SO4, filtered and solvent removed under reduced pressure. Purification by flash column chromatography (1:2 EtOAc:Hexanes) and recrystallization from CH2CI2/ Hexanes afforded imidazole 11 as a white solid (24.6 mg, 34% yield from azide 9). ¹H NMR (500 MHz, CDCl₃) δ 7.63 (AA'XX', 2H, J_{AX} = 8.81, $J_{XX'}$ = 2.53), 7.54 (m, 2H), 7.34 (AA'XX', 2H, J_{AX} = 8.80, $J_{XX'}$ = 2.45), 7.20 (m, 2H), 7.12 (m, 1H), 7.02 (AA'XX', 2H, J_{AX} = 8.78, J_{AA'} = 2.54), 7.01 (AA'XX', 2H, J_{AX} = 8.43, $J_{AA'}$ = 2.57), 3.90 (q, 2H, J = 7.08), 3.89 (s, 3H), 3.87 (s, 3H), 1.02 (t, 3H, J = 7.17); ¹³C NMR $(125 \text{ MHz}, \text{CDCl}_3) \delta 200..8, 160.0, 159.8, 147.0, 134.8, 132.3 (2),$ 130.5 (2), 129.3, 128.9, 128.0 (2), 126.6 (2), 126.0, 123.6, 114.5 (2), 114.0 (2), 55.33, 55.28, 36.4, 16.2; MS (EI, 70 eV) m/z 384.2 (M⁺); Anal. calc'd for C₂₅H₂₄N₂O₂, C: 78.10%, H: 6.29%, N: 7.29%, found, C: 77.91%, H:6.28%, N: 7.28%.

General demethylation procedure using BF₃•SMe₂. To a stirring solution of the methyl protected heterocycle (1 equiv) in CH₂Cl₂ (8 ml) at room temperature was added BF₃•SMe₂ complex (75 equiv). After stirring for 24 h, solvent and excess reagent were evaporated under nitrogen stream in hood. Residue was taken up in EtOAc and washed with H₂O and sat. NaCl. Organic extract was dried over Na₂SO₄, filtered and solvent removed under reduced pressure. The resulting residue was purified though a silica plug, eluting with EtOAc. Solvent evaporation afforded the deprotected products.

1-*Ethyl*-2,5-(4-*hydroxyphenyl*)-4-*phenyl imidazole* (12). Imidazole 11 (12.0 mg, 0.031 mmol) was demethylated according to the general BF₃•SMe₂ procedure to afford imidazole 12 as an off-white powder (10.6 mg, 95%). ¹H NMR (500 MHz, Acetone-*d*₆) δ 7.80 (AA'XX', 2H, *J*_{AX}=8.81, *J*_{XX}:=2.44), 7.47-7.49 (m, 2H), 7.44 (AA'XX', 2H, *J*_{AX}=8.65, *J*_{XX}:=2.44), 7.37-7.40 (m, 3H), 7.14 (AA'XX', 2H, *J*_{AX}=8.80, *J*_{AA'}=2.43), 7.06 (AA'XX', 2H, *J*_{AX}=8.68, *J*_{AA'}=2.46), 4.25 (q, 2H, *J*=7.28), 1.17 (t, 3H, *J*=7.29); MS (FAB) *m/z* 357.2 (M+H⁺); HRMS calc'd for C₂₃H₂₁N₂O₂: 357.160303, found: 357.160000.

5-Ethyl-1,4-(4-methoxyphenyl)-2-phenyl imidazole (**16**). Keto-amide **15** (110.0 mg, 0.273 mmol) and ammonium acetate (105.0 mg, 1.362 mmol) were heated to reflux in acetic acid (10 ml) for 48 h. Acetic acid was removed under reduced pressure, resulting residue was taken up in EtOAc, washed with sat. NaHCO₃, H₂O, and sat. NaCl. Organic extracts were dried over Na₂SO₄, filtered and solvent removed. Product was purified by flash column chromatography (1:4 EtOAc:Hexanes) and recrystallization from CH₂Cl₂/Hexanes to give imidazole **16** as a white solid (25.7 mg, 25%). ¹H NMR (500 MHz, CDCl₃) δ 7.72 (AA'XX', 2H, J_{AX} = 8.29, J_{XX'} = 2.55), 7.14 (m, 2H), 7.21 (m, 3H), 7.19 (AA'XX', 2H, J_{AX} = 9.33, J_{XX'} = 2.71), 6.98 (AA'XX', 2H, J_{AX} = 8.48, J_{AA'} = 2.52), 6.96 (AA'XX', 2H, J_{AX} = 8.62, J_{AA'} = 2.74), 3.87 (s, 3H), 3.85 (s, 3H), 2.67 (q, 2H, J = 7.48), 1.01 (t, 3H, J = 7.45); MS (EI, 70 eV) *m/z* 384.2 (M⁺).

5-*Ethyl-1*,4-(4-hydroxyphenyl)-2-phenyl imidazole (17). Imidazole 16 (25.0 mg, 0.065 mmol) was demethylated as outlined in general BF₃•SMe₂ procedure above to give deprotected imidazole 17 as an off-white powder (20.2 mg, 87%). ¹H NMR (400 MHz, Acetone- d_6) δ 9.04 (br s, 1H), 8.51 (br s, 1H), 7.64 (AA'XX', 2H, J_{AX} =8.73, J_{XX} =2.51), 7.50-7.47 (m, 2H), 7.31-7.27 (m, 5H), 7.01 (AA'XX', 2H, J_{AX} =8.94, $J_{AA'}$ =2.76), 6.94 (AA'XX', 2H, J_{AX} =8.73, $J_{AA'}$ =2.52), 2.69 (q, 2H, J=7.48), 1.02 (t, 3H, J=7.49); MS (FAB) m/z 357.1 (M+H⁺); HRMS calc'd. for C₂₃H₂₁N₂O₂: 357.1603, found 357.1602.

2,4-Di(4-methoxyphenyl)-thiazole (**21a**). A suspension of thioamide **19** (1.3 g, 7.9 mmol) and α -bromo-4'-methoxy-acetophenone (**20**) (1.8 g, 7.9 mmol) in DMF (10 ml) was heated to reflux for 1h, until it became homogeneous. The heat was removed and the reaction was stirred for 15 h at room temperature. The reaction mixture was poured into H₂O (50 ml) and the solid precipitate was filtered to afford crude **21a**. Recrystallization from CH₃NO₂ afforded pure **21a** as light yellow crystals (1.8 g, 81%). ¹H NMR (400 MHz, CDCl₃) δ 7.98 (AA'XX', 2H, J_{AX} = 8.87, J_{AA} = 2.53), 7.63 (AA'XX', 2H, J_{AX} = 8.94, J_{AA} = 2.48), 6.85 (AA'XX', 2H, J_{AX} = 8.97, J_{AA} = 2.53), 6.85 (AA'XX', 2H, J_{AX} = 8.94, J_{AA} = 2.48), 7.26 (s, 1H), 3.86 (s, 3H), 3.85 (s, 3H); ¹³C NMR (400 MHz, CDCl₃) δ 167.5, 160.9, 159.4, 155.6, 127.9, 127.6, 127.4, 126.6, 114.1, 113.9, 109.9, 55.3, 55.2; MS (EI, 70 eV) *m/z* (relative intensity, %) 297.1 (M⁺, 100), 282.1 (10), 164.1 (30), 149.1 (55), 133.1 (10), 121.1 (25), 77.1 (15); HRMS calc'd for C₁₇H₁₅NSO₂: 297.082469, found: 297.082351.

2,4-Di(4-hydroxyphenyl)-thiazole (22a). Thiazole 21a (1.0 g, 3.6 mmol) was demethylated using BBr₃ as outlined in the general procedure above to afford 22a (430 mg, 45%). mp 218–221°C; ¹H NMR (400 MHz, Acetone- d_6) δ 8.84 (br s, 2H), 7.92 (AA'XX', 4H, J_{AX} = 8.57, J_{AA} = 2.17), 7.58 (s, 1H), 6.96 (AA'XX', 2H, J_{AX} = 8.79, J_{AA} = 2.51), 6.92 (AA'XX', 2H, J_{AX} = 8.78, J_{AA} = 2.44); ¹³C NMR (100 MHz, Acetone- d_6) δ 168.2, 159.3, 157.2, 155.8, 127.7, 127.3, 126.2, 125.1, 115.2, 114.9, 109.3; MS (EI, 70 eV) *m/z* (relative intensity, %) 296.1 (M^{+.} 100), 150.1 (27), 121.1 (11), 78.1 (8); HRMS calc'd for C₁₅H₁₁NSO₂: 269.051163, found: 269.051051.

2,4-(4-Methoxyphenyl)-5-phenyl oxazole (28). Azido-ketone 27 (0.18 g, 0.673 mmol) and *p*-anisaldehyde (0.25 ml, 2.05 mmol) were dissolved in THF (15 ml). Et₃N (94.0 μ l, 0.674 mmol) was added via syringe and reaction stirred at room temperature for 48 h. The reaction mixture was then poured into H₂O and extracted with CH₂Cl₂, organic fraction was dried over Na₂SO₄, filtered and solvent removed under reduced pressure. Resulting intermediate 2,5-dihydro-5-hydroxyoxazole, used in next

step without further purification or characterization, was taken up CH₂Cl₂ (10 ml). Solution was cooled to 0°C and TFA (54.0 µl, 0.701 mmol) was added via syringe. Reaction stirred at 0°C for 36 h. The mixture was diluted with CH₂Cl₂ (10 ml) and washed with H₂O, sat. NaHCO₃, and sat. NaCl successively. Organic extracts were combined, dried over Na₂SO₄, filtered and solvent removed under reduced pressure. Purification by flash column chromatography (1:2 EtoAc: hexanes) and recrystallization from CH₂Cl₂/Hexanes afforded oxacole **28** as a white solid (72.4 mg, 30% yield from azide 27). mp 125–128°C (lit. [27] mp 126–127°C); ¹H NMR (500 MHz, CDCl₃) & 7.84 (AA'XX', 2H, J_{AX}= 8.89, J_{XX}:= 2.47), 7.43 (AA'XX', 2H, J_{AX}= 8.83, J_{XX}:= 2.48), 7.53 (m, 2H), 7.32 (m, 2H), 7.26 (tt, 1H, J= 7.03, 1.42), 6.97 (AA'XX', 2H, J_{AX}= 8.85, J_{AA'}= 2.53), 6.88 (AA'XX', 2H, J_{AX}= 8.70, J_{AA'}= 2.53), 3.84 (s, 3H), 3.81 (s, 3H).

2,4-(4-Hydroxyphenyl)-5-phenyl oxazole (29). Oxazole 28 (22.0 mg, 0.062 mmol) was demethylated according to the general BF₃•SMe₂ procedure above to give deprotected oxazole 29 as an off-white powder (18.8 mg, 93%). ¹H NMR (500 MHz, Acetone- d_6) δ 9.43 (br s, 1H), 8.90 (br s, 1H), 8.08 (AA'XX', 2H, J_{AX} =9.06, $J_{XX'}$ =2.62), 7.61 (m, 2H), 7.49 (m, 3H), 7.44 (AA'XX', 2H, J_{AX} =8.97, $J_{XX'}$ =2.44), 7.12 (AA'XX', 2H, J_{AX} =8.87, $J_{AA'}$ =2.50), 6.95 (AA'XX', 2H, J_{AX} =8.86, $J_{AA'}$ =2.42); MS *m/z* 329.1 (M⁺); HRMS calc'd. for C₂₁H₁₅NO₃: 329.1052, found 329.1285.

2,5-(4-Methoxyphenyl)-4-phenyl oxazole (**30**). A solution of bromoketone **26** (87.0 mg, 0.285 mmol) and *p*-methoxybenzamide (43.0 mg, 0.285 mmol) in toluene was heated to reflux for 36 h. Toluene was removed under reduced pressure and resulting residue purified by flash column chromatography (1:4 EtOAc:Hexanes). Recrystallization of desired product from CH₂Cl₂/hexanes afforded oxazole **30** as a colorless solid (52.9 mg, 52%). mp 147–149°C; ¹H NMR (500 MHz, CDCl₃) δ 8.08 (AA'XX', 2H, J_{AX} = 8.58, J_{XX}:= 2.24), 7.72 (m, 2H), 7.59 (AA'XX', 2H, J_{AX} = 8.63, J_{AA}:= 2.61), 7.39 (m, 2H), 7.33 (m, 1H), 6.99 (AA'XX', 2H, J_{AX} = 8.63, J_{AA}:= 2.62), 6.92 (AA'XX', 2H, J_{AX} = 8.68, J_{AA}:= 2.67), 3.88 (s, 3H), 3.85 (s, 3H); ¹³C NMR (100 MHz, CDCl₃) δ 161.2, 159.9, 159.7, 145.1, 135.3, 132.9, 128.6 (2), 128.1 (2), 128.0 (2), 127.96 (2), 127.9, 121.7, 120.3, 114.2 (2), 114.1 (2), 55.4, 55.3; MS *mlz* 357.2 (M⁺).

2,5-(4-Hydroxyphenyl)-4-phenyl oxazole (31). Oxazole 30 (22.0 mg, 0.062 mmol) was demethylated according to the general BF₃•SMe₂ procedure above to give deprotected oxazole 31 as an off-white powder (18.1 mg, 89%). ¹H NMR (500 MHz, Acetone-*d*₆) δ 8.93 (br s, 1H), 8.77 (br s, 1H), 7.99 (AA'XX', 2H, *J*_{AX}=8.83, *J*_{XX'}=2.41), 7.72 (m, 2H), 7.53 (AA'XX', 2H, *J*_{AX}=8.77, *J*_{XX'}=2.46), 7.40 (m, 2H), 7.34 (tt, 1H, *J*=7.36, 1.33), 6.99 (AA'XX', 2H, *J*_{AX}=8.70, *J*_{AA}=2.40), 6.92 (AA'XX', 2H, *J*_{AX}=8.83, *J*_{AA'}=2.46); MS *m*/z 329.1 (M⁺); HRMS calc'd for C₂₁H₁₅NO₃: 329.1052, found 329.1055.

General procedure for pyrazole synthesis. A suspension of diketone (1 equiv) and appropriate hydrazine hydrochloride (3–5 equiv) in a 3:1 mixture DMF:THF was heated to reflux for 16–24 h with reaction progress being monitored by TLC for disappearance of starting material. The reaction mixtures was cooled to room temperature and poured into iced sat. LiCl solution (10 ml) and EtOAc (10 ml). The layers were separated and the organic layer was washed with brine (10 ml), dried over MgSO₄, filtered and concentrated. Purification using flash column chromatography (EtOAc/hexanes systems) afforded the pyrazoles.

3,5-*di*(4-*methoxyphenyl*)-1*H*-*pyrazole* (34a). Diketone **33** (91 mg, 0.32 mmol) and hydrazine (0.1 ml, 3.2 mmol) were reacted as outlined in general pyrazole procedure to afford **34a** [45] as an off-white solid (32.6 mg, 38%). mp 172–175°C (lit [45] mp 174°C); ¹H NMR (400 MHz, CDCl₃) δ 7.73 (AA'XX', 4H, J_{AX} = 8.73, J_{AA} = 2.42), 6.97 (AA'XX', 4H, J_{AX} = 8.73, J_{AA} = 2.42), 6.80 (s, 1H), 3.72 (s, 6H); ¹³C NMR (100 MHz, CDCl₃) δ 159.9, 148.3, 126.8, 123.0, 112.9, 98.6, 54.5; MS (FAB) *m/z* (relative intensity, %) 281 (MH⁺, 100).

1-Phenyl-3,5-di(4-methoxyphenyl)-pyrazole (**34b**). Diketone **33** (100 mg, 0.35 mmol) and phenyl hydrazine hydrochloride (500 mg, 3.5 mmol) were reacted as outlined in general pyrazole procedure above to afford **34b** [46] (30 mg, 25%). mp 159–161°C (lit [46] mp 163°C); ¹H NMR (400 MHz, CDCl₃) δ 7.82 (AA'XX', 2H, J_{AX} = 8.96, J_{AA} = 2.44), 7.24–7.20 (m, 5H), 7.20 (AA'XX', 2H, J_{AX} = 8.79, J_{AA} = 2.46), 6.99 (AA'XX', 2H, J_{AX} = 8.79, J_{AA} = 2.46), 6.99 (AA'XX', 2H, J_{AX} = 8.79, J_{AA} = 2.46), 6.99 (AA'XX', 2H, J_{AX} = 8.96, J_{AA} = 2.44), 6.70 (s, 1H), 3.84 (s, 3H), 3.80 (s, 3H); ¹³C NMR (100 MHz, CDCl₃) δ 159.4, 151.5, 144.1, 140.0, 129.9, 128.8, 127.2, 126.9, 125.5, 125.2, 122.9, 113.9, 113.8, 104.1, 55.2, 55.1; MS (EI, 70 eV) *m/z* (relative intensity, %) 356 (M⁺, 100), 341 (19), 135 (89); HRMS calc'd for C₂₃H₂₀N₂O₂: 356.15241, found: 356.152478.

3,5-Di(4-hydroxyphenyl)-1H-pyrazole (35a). Pyrazole 34a (20 mg, 0.07 mmol) was demethylated with BBr₃ according to the general procedure to afford 35a [47] as an off-white solid (11 mg, 63%). ¹H NMR (400 MHz, Acetone-*d*₆) δ 8.58 (br s, 2H), 7.75 (AA'XX', 4H, *J*_{AX} = 8.95, *J*_{AA} = 2.46), 6.93 (AA'XX', 4H, *J*_{AX} = 8.95, *J*_{AA} = 2.46), 6.83 (s, 1H); ¹³C NMR (100 MHz, Acetone-*d*₆) δ 157.1, 148.3, 126.5, 123.0, 115.3, 97.5; MS (CI, CH₄) *m/z* (relative intensity, %) 253.1 (MH⁺, 100), 237(10), 161 (5), 123 (15).

1-Phenyl-3,5-di(4-hydroxyphenyl}-pyrazole (**35b**). Pyrazole **34b** (20 mg, 0.06 mmol) was demethylated with BBr₃ according to the general procedure to afford **34b** [47] as an off-white solid (11.5 mg, 58%). ¹H NMR (400 MHz, CDCl₃) δ 8.64 (s, 1H), 8.45 (s, 1H) 7.79 (AA'XX', 4H, J_{AX} = 8.78, J_{AA} = 2.38), 7.36 (m, 5H), 7.14 (AA'XX', 4H, J_{AX} = 8.60, J_{AA} = 2.47), 6.90 (AA'XX', 2H, J_{AX} = 8.78, J_{AA} = 2.38), 6.81 (AA'XX', 2H, J_{AX} = 8.78, J_{AA} = 2.47), 6.90 (AA'XX', 2H, J_{AX} = 8.78, J_{AA} = 2.38), 6.81 (AA'XX', 2H, J_{AX} = 8.78, J_{AA} = 2.47), 6.80 (s, 1H); ¹³C NMR (100 MHz, CDCl₃) δ 157.5, 157.3, 151.3, 144.1, 140.6, 129.9, 128.6, 126.8, 126.7, 124.9, 122.2, 122.1, 115.2, 115.1, 103.7; MS (EI, 70 eV) *m/z* (relative intensity, %) 362.1 (M⁺, 85), 328.1 (100).

4-*Ethyl*-3,5-*di*(4-*methoxyphenyl*)-1*H*-*pyrazole* (**37***a*). Diketone **36** (100 mg, 0.32 mmol) and hydrazine (0.12 ml, 3.2 mmol) were reacted as outlined in the general pyrazole procedure above to afford **37***a* as a while solid (69 mg, 70%). ¹H NMR (400 MHz, CDCl₃) δ 7.50 (AA'XX', 4H, J_{AX} = 8.84, J_{AA} = 2.48), 6.94 (AA'XX', 4H, J_{AX} = 8.90, J_{AA} = 2.50), 3.85 (s, 3H), 2.71 (q, 2H, J = 7.38), 1.07 (t, 3H, J = 7.44); ¹³C NMR (100 MHz, CDCl₃) δ 159.4, 129.0, 127.4, 124.1, 116.8, 113.9, 55.1, 16.6, 15.3; MS (EI, 70 eV) *m/z* (relative intensity, %) 308.1 (M⁺, 100), 293.1 (73), 160.1 (7), 134 (8).

1-Phenyl-4-ethyl-3,5-di(4-methoxyphenyl)-pyrazole (**37b**). Diketone **36** (100 mg, 0.35 mmol) and phenyl hydrazine hydrochloride (140 mg, 0.96 mmol) were reacted as outlined in the general pyrazole procedure above to afford 37b as an orange solid (109 mg, 87%). ¹H NMR (400 MHz, CDCl₃) δ 7.72 (AA'XX', 2H, J_{AX} = 9.03, J_{AA} = 2.44), 7.24 (m, 3H), 7.20 (m, 2H), 7.17 (AA'XX', 2H, J_{AX} = 8.79, J_{AA} = 2.44), 6.99 (AA'XX', 2H, J_{AX} = 8.79, J_{AA} = 2.44), 3.86 (s, 3H), 3.83 (s, 3H), 2.63 (q, 2H, J = 7.57), 1.04 (t, 3H, J = 7.57); ¹³C NMR (100 MHz, CDCl₃) δ 159.6, 159.4, 150.8, 141.2, 140.5, 131.5, 129.3, 128.8, 127.0, 126.7, 124.8, 123.5, 120.7, 114.2, 114.1, 55.5, 55.4, 17.3, 15.8; MS (EI, 70 eV) *m/z* (relative intensity, %) 356 (M⁺, 100), 341 (100), 328 (15), 196 (25), 77 (40); HRMS calc'd for C₂₅H₂₄N₂O₂: 384.183582, found: 384.183778.

*4-Ethyl-*3,5-*di*(*4-hydroxyphenyl*)-1*H-pyrazole* (*38a*). Pyrazole **37a** (69 mg, 0.22 mmol) was demethylated according to the general BBr₃ procedure to afford **38a** as a white solid (35 mg, 57%). ¹H NMR (400 MHz, Acetone-*d*₆) & 7.49 (AA'XX', 4H, *J*_{AX} = 8.85, *J*_{AA} = 2.46), 6.93 (AA'XX', 4H, *J*_{AX} = 8.864, *J*_{AA} = 2.40), 2.73 (q, 2H, *J* = 7.39), 1.07 (t, 3H, *J* = 7.47); ¹³C NMR (100 MHz, Acetone-*d*₆) & 156.9, 128.8, 124.3, 122.2, 115.4, 115.2, 16.5, 14.9; MS (CI, CH₄) *m/z* (relative intensity, %) 282.1 (M+H⁺, 100), 263.1 (10), 187.1 (20).

1-Phenyl-4-ethyl-3,5-di(4-hydroxyphenyl)-pyrazole (**38b**). Pyrazole **37b** (100 mg, 0.26 mmol) was demethylated according to the general BBr_3 procedure to afford **38b** as a white solid (50 mg, 54%). ¹H NMR

 $\begin{array}{l} (400 \text{ MHz}, \text{MeOH-}d_{4}) \ \& 7.51 \ (\text{AA'XX'}, 4\text{H}, J_{\text{AX}} = 8.67, J_{\text{AA}} = 2.47), \ 7.24-7.42 \ (\text{m}, 5\text{H}), \ 7.05 \ (\text{AA'XX'}, 4\text{H}, J_{\text{AX}} = 8.81, J_{\text{AA}} = 2.40), \ 6.88 \ (\text{AA'XX'}, 2\text{H}, J_{\text{AX}} = 8.67, J_{\text{AA}} = 2.46), \ 6.78 \ (\text{AA'XX'}, 2\text{H}, J_{\text{AX}} = 8.81, J_{\text{AA}} = 2.40), \ 2.60 \ (\text{q}, 2\text{H}, J = 7.53), \ 0.98 \ (\text{t}, 3\text{H}, J = 7.39); \ \text{MS} \ (\text{EI}, 70 \ \text{eV}) \ \textit{m/z} \ (\text{relative intensity}, \ \%) \ 356.1 \ (\text{M}^{+}, 100), \ 341.1 \ (100), \ 328.1 \ (15), \ 196.1 \ (25), \ 77 \ (40); \ \text{HRMS} \ \text{calc'd for } C_{23}\text{H}_{21}\text{N}_{2}\text{O}_{2}; \ 357.161155, \ \text{found:} \ 357.160303. \end{array}$

3,5-Di(4-methoxyphenyl)isoxazole (40). To a solution of oxime 39 (1.0 g, 6 mmol) in THF (20 mL) at 0°C was added nBuLi (9.11 ml, 13.3 mmol) as a solution in hexanes. The clear solution was stirred for 30 min at 0°C then methyl 4-methoxybenzoate (498 mg, 3 mmol) was added as a solution in THF (5 ml) over 5 min. The reaction mixture was stirred at 0°C for 30 min, then warmed to room temperature. 5 N HC (10 ml) was added and the biphasic reaction mixture was brought to reflux overnight (12 h). Upon cooling to 0°C, isoxazole 40 precipitated [33] and was collected via filtration (450 mg, 27%). mp 174-177°C (lit [33] mp 176-177°C); ¹H NMR (400 MHz, Acetone d_β) δ 7.66 (AA'XX', 2H, $J_{AX} = 8.88$, $J_{AA} = 2.44$), 7.63 (AA'XX', 2H, $J_{AX} = 9.1$, $J_{AA} = 2.15$), 6.86 (AA'XX', 2H, $J_{AX} = 8.88$, $J_{AA} = 2.44$), 6.85 (AA'XX', 2H, $J_{AX} = 9.1$, $J_{AA} = 2.44$), 6.57 (s, 1H), 3.73 (s, 3H), 3.72 (s, 3H); ¹³C NMR (100 MHz, Acetone-d₆) δ 169.8, 162.3, 160.8, 160.7, 127.9, 127.1, 121.5, 120.0, 114.1, 114.0, 95.7, 55.1, 55.0; MS (EI, 70 eV) m/z (relative intensity, %) 281.1 (M+, 5), 150.1 (20), 135.1 (100).

3,5-*Di*(4-*hydroxyphenyl*)*isoxazole* (41). Isoxazole 40 (300 g, 1.1 mmol) was demethylated according to the general BBr₃ procedure to afford 41 [34] as a white solid (152 mg, 56%). mp 267–269°C (lit [34] mp 255°C); ¹H NMR (400 MHz, Acetone-*d*₆) δ 10.07 (s, 1H), 9.91 (s, 1H), 7.90 (d, 4H, *J*=8.79), 7.21 (s, 1H), 6.88 (4H, t, *J*=9.38); ¹³C NMR (100 MHz, MeOH-*d*₄) δ 170.3, 162.8, 159.1, 127.7, 126.9, 119.9, 118.7, 116.3, 115.1, 95.3, 95.0; MS (EI, 70 eV) *m/z* (relative intensity, %) 253.1 (M⁺, 60), 133.1 (25), 121.1 (100), 93.0 (20), 77.0 (10), 65.0 (30).

Supplementary material available

Experimental detail regarding the preparation of all intermediates discussed in the synthesis of the above heterocycles is available with the online version of this paper.

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Acyclic Amides as Estrogen Receptor Ligands: Synthesis, Binding, Activity and Receptor Interaction

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Abstract—We have prepared a series of bisphenolic amides that mimic bibenzyl and homobibenzyl motifs commonly found as substructures in ligands for the estrogen receptor (ER). Representative members were prepared from three classes: N-phenyl benzamides, N-phenyl acetamides, and N-benzyl benzamides; in some cases the corresponding thiocarboxamides and sulfonamides were also prepared. Of these three classes, the N-phenyl benzamides had the highest affinity for ER, the N-phenyl acetamides had lower, and the N-benzyl benzamides were prone to fragmentation via a quinone methide intermediate. In the N-phenyl benzamide series, the highest affinity analogues had bulky N-substituents: a CF₃ group, in particular, conferred high affinity. The thiocarboxamides bound better than the corresponding carboxamides and these bound better than the corresponding sulfonamides. Binding affinity comparisons suggest that the p-hydroxy group on the benzoate ring, which contributes most to the binding, is playing the role of the phenolic hydroxyl of estradiol. Computational studies and NMR and X-ray crystallographic analysis indicate that the two anilide systems studied have a strong preference for the s-cis or exo amide conformation, which places the two aromatic rings in a syn orientation. We used this structural template, together with the X-ray structure of the ER ligand binding domain, to elaborate an additional hydrogen bonding site on a benzamide system that elevated receptor binding further. When assayed on the individual ER subtypes, ER α and ER β , these compounds show modest binding affinity preference for ER α . In a reporter gene transfection assay of transcriptional activity, the amides generally have full to nearly full agonist character on ER α , but have moderate to full antagonist character on ER β . One high affinity carboxamide is 500-fold more potent as an agonist on ER α than on ER β . This work illustrates that ER ligands having simple amide core structures can be readily prepared, but that high affinity binding requires an appropriate distribution of bulk, polarity, and functionality. The strong conformational preference of the core anilide function in all of these ligands defines a rather rigid geometry for further structural and functional expansion of these series. (C) 2000 Elsevier Science Ltd. All rights reserved.

Introduction

Recent advances in nuclear hormone receptor pharmacology and mechanism of action have redefined the classification of estrogen receptor (ER) ligands.¹ The selective estrogen receptor modulator class, or SERMs, are considered to be very important because of the potential of SERMs for maintaining bone mineral density and cardiovascular health, and for treating and preventing breast cancer and other hormone-dependent disorders, without adverse stimulation of the uterus and breast.² SERMs comprise several structurally diverse classes, including the triarylethylenes, triarylnaphthalenes, benzo[b]furans, benzopyrans, and various other tetracyclic manifestations of these core structures.³ Collectively, SERMs display a wide range of tissueselective agonist and antagonist activities, but major efforts continue to be directed toward optimizing ER ligand structure in order to obtain desired tissue-specific properties with minimal side-effects, which is an important concern in maintaining efficacy and patient compliance.⁴

In connection with our interests in novel templates for ER ligands, especially ones that might be suitable for

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combinatorial development,⁵ we wondered whether acyclic tertiary amides might be developed as mimics of common structural motifs found at the core of many ER ligands and thus support a combinatorial approach to SERM discovery and optimization. In this report, we describe the design and synthesis of three types of amide-core ER ligands, and we investigate their molecular structure, their structure binding-affinity relationships, and their activity in ER α and ER β transactivation assays. A number of interesting trends and high affinity amides were discovered during this investigation. One high affinity amide in particular (**16g**) was found to be an ER α -potency selective agonist.

Results and Discussion

Rationale and design

To the best of our knowledge, only two examples of ER ligands containing, either partially or exclusively, an amide core structure are known in the public literature. Hartmann and co-workers studied a small series of bisphenolic carboxamides^{6,7} containing an invariant N-(1,1,1-trifluoro-2-propyl) substituent (e.g., 1) and found them to be anti-estrogens with weak to moderate potency. An early example of an N-ethyl benzene sulfonamide $(2)^8$ was also reported and shown to be weakly estrogenic in vivo. Neglecting amide bond stereoisomers for the moment, these ligands represent potential mimics (1) for either the anti- or syn-bibenzyl motifs A and **B**, respectively (Fig. 1, right), that are found in many common ER ligands (Fig. 1, left). In addition to the bibenzyl structural template of these leads, we envisioned two homologues (II and III, Fig. 2, right) which mimic the homo-bibenzyl motif C (Fig. 2, II-III), a structural element that is also well represented in high affinity ligands for the ER (Fig. 2, left).



From this analysis, model *para*-substituted bis-phenolic carboxamides, sulfonamides and thioamides containing N-alkyl and trifluoroalkyl substituents (R and R') were considered as potential templates for combinatorial development (Fig. 3). Preparation for many of these compounds is straightforward and thus not unreasonable for adaptation to combinatorial library synthesis.

Chemical syntheses

N-Phenyl benzamides and benzene sulfonamides. To synthesize simple N-alkyl amide analogues in an expeditious manner, we used catalytic phase transfer conditions (Scheme 1). Thus, N-alkylation of the known secondary benzene sulfonamide 3^8 proceeded under mild conditions using *n*-Bu₄NSO₄H, NaOH, and excess alkyl halide in $CH_2Cl_2/H_2O.^9$ The target N-alkylated sulfonamides 5a-c were then obtained after deprotection. The less acidic carboxamide 6 was alkylated using a solid-liquid two-phase system, consisting of powdered NaOH and cat. n-Bu₄NBr in refluxing benzene, to afford the tertiary amides 9a-e in good yield.^{9,10} Carboxamides 9f and 9g were conveniently prepared from the secondary anilines 7 and 8. Aniline 7 was prepared via reductive alkylation of the corresponding aldimine using a modified literature procedure.¹¹ Aniline 8 was obtained by reductive amination, using NaBH(OAc)₃, *p*-anisidine, and 3-methyl-2-butanone. Thioamides were prepared in relatively good yield by treating the carboxamide (9a-g) with Lawesson's reagent.¹² Deprotection with BBr₃ or



Figure 1. Syn and anti-bibenzyl motifs and proposed acyclic amide analogues.



Figure 2. Homo-bibenzyl motif and proposed acylic amide analogues.



Figure 3. Potential amide ER ligand scaffolds for combinatorial chemistry (X = O, S).



Scheme 1. Synthesis of *N*-phenyl benzamides.

ethanethiolate afforded the target benzamides **10a–f** and thiobenzamides **11a–g**.

CF₃-containing N-phenyl benzamides. To further explore the effect that a trifluoromethyl group would have on receptor binding affinity, we prepared several carboxamide and thioamide analogues related to the weak antiestrogen (1) reported by Hartmann and co-workers.⁶ The synthetic route to CF₃-amide analogues, depicted in Scheme 2, was adapted from that reported by Hartmann and required introduction of the CF3-alkyl substituent at the aniline stage in two steps starting from the corresponding iminophosphoranes. The starting trifluoromethylketones were commercially available, with the exception of 2,2,2-trifluoro-1-(3-methoxyphenyl)ethanone, which was prepared from 3-bromoanisole according to the procedures described by Hatanaka et al. and references therein.¹³ Ylides **12a** and **12b** were readily obtained by treatment of the corresponding azides with Ph₃P.^{14,15} Subsequent Staudinger reaction to afford the CF₃-substituted imines 13a d, followed by LAH reduction, furnished the desired secondary anilines **14a**-d in good overall yield.

Amidation in warm toluene proceeded satisfactorily to provide carboxamides **15a–e**; however, unactivated anilines **14c** and **14d** required 1,2-dichlorobenzene as solvent and higher temperatures to afford benzamides **15f** and **15g** in satisfactory yields. Subsequent deprotection using BBr₃ afforded the desired phenols **16a–g** as before. Thionation of the protected, CF₃-substituted carboxamides proceeded in significantly lower yield (20–51%) than did those with simple *N*-alkyl substitution (>79%). Alternative thionation conditions, such as P₂S₅ and POCl₃/(TMS)₂S, did not give improved yields. Nevertheless, the two thioamides **17a** and **17b** could be obtained after deprotection with BBr₃.

N-Phenyl acetamides and *N*-benzyl benzamides. Acetamides were attractive because they offer an additional site for structural diversity that is not available in benzamides (Scheme 3). *N*-Ethylation of amide 18 using PTC conditions as before afforded 19 in excellent yield. Subsequent α -alkylation using LDA at -78 °C then afforded the disubstituted carboxamides 20a-b in high yield. Thionation and/or deprotection as before furnished







Scheme 3. Synthesis of N-phenyl-acetamides.

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the desired bis-phenolic amides **21a-b** and **22a-c** without complication.

The *N*-benzyl-*N*-alkylbenzamides **24** (Scheme 4), which are essentially amide isomers of the acetamide class, and related benzenesulfonamides (not shown), were also investigated. These *N*-benzyl amides, protected as methyl ethers, could be prepared by routes that are analogous to those used in Schemes 1–3. However, these systems proved to be unstable to various deprotection conditions: BBr₃, ethanethiolate, HBr/AcOH and TMSI all led to decomposition. The only isolatable products were the corresponding secondary amides, presumed to result from an elimination of the 4-hydroxy-*N*-benzyl substituent via a quinone-methide intermediate (Scheme 4). Due to their propensity for elimination, additional efforts were not made to prepare members of the *N*benzyl benzamide class.

CF₃-containing *N*-**phenyl acetamides.** Incorporation of the CF₃-group was also explored in the acetamide series, and the synthesis of α -substituted acetamides containing an *N*-(1,1,1-trifluoro-2-propyl) substituent is shown in Scheme 5. Thus, the *N*-(1,1,1-trifluoro-2-propyl) substituted aniline **12a** was treated with 4-methoxy-phenylacetyl chloride to afford carboxamide **25**. α -Alkylation as before using LDA at -78 °C and MeI proceeded in excellent yield to give **26** as a \sim 1:1 mixture of diastereomers as indicated by ¹H NMR and HPLC analysis. Demethylation then afforded the di-substituted phenylacetamide **27** in good yield. Unfortunately,

attempts to thionate **26** with Lawesson's reagent failed; however, treatment of the unsubstituted phenylacetamide **25** with Lawesson's reagent did give thioamide **28**, in albeit low yield. Repeated attempts using other reagents (P_2S_5 and $POC_{13}/(TMS)_2S$) or higher temperatures failed to provide thioamide **26** or to improve the yield of **28**. Increased steric hindrance about the C=O bond is thought to be responsible for the lack of reactivity of the CF₃-substituted *N*-phenyl acetamides towards O–S replacement, as lower yields for thionation were also observed for the CF₃-containing benzamides (Scheme 2).

Molecular conformation

As was noted in Figure 1, bibenzyl motifs in ER ligands are well represented as both syn and anti conformations. Since N-aryl-benzamides share a similar bibenzyl-like two-atom connection between two aromatic rings, we were interested in their conformational preference to establish which structural motif they might be mimicking. From early ¹H NMR studies of N-substituted anilides, there is known to be a surprisingly strong preference for the exo-isomer (N-phenyl group trans to carbonyl oxygen).¹⁶ Even simple N-methyl and Nethylformanilides, where steric considerations would place the bulkier alkyl group next to the formyl hydrogen in an endo preference, have a 95% exo preference in solution.¹⁷ This so-called '*exo*-rule' is quite general, and it has been stated that for N-substituted anilides other than formyl, the exo-isomer dominates to the exclusion



Scheme 4. Attempted synthesis of N-benzyl-benzamides.



Scheme 5. Synthesis of CF₃-containing N-alkyl-phenylacetamides.

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of the *endo*-isomer.¹⁶ The *endo*-isomer is detectable only when large *ortho* substituents are present on the *N*-phenyl ring or when R'=H.



Molecular modeling studies on both carboxamide 16a and thiocarboxamide 17a that we have performed using molecular mechanics (MM2) were in agreement with the reported NMR observations, and indicated the exo or s-cis conformer as being energetically more favorable (Fig. 4). In the case of carboxamide 16a, a substantial ΔH of 5.3 kcal/mol between the global minimum energy cis and trans isomers was found, the major energetic difference being due to additional electrostatic contributions in the s-trans conformation. Thiocarboxamide 17a has a smaller ΔH of 0.9 kcal/mol between the global minimum energy cis and trans isomers (not shown). This result is in accord with destabilization of the s-cis (exo) thioamide conformer relative to the s-trans (endo) conformer because of additional steric repulsions between sulfur and the N-alkyl substituent that arise from the larger covalent radii of sulfur versus oxygen (1.4 Å for sulfur, 0.74 Å for oxygen) and its longer bond length (C=S 1.64 Å; C=0 1.24 Å).

Direct structural studies of *N*-substituted thioanilides, let alone hindered systems containing a CF_3 -substituent, have not been reported, as far as we have found. For this reason and because of the relatively small energy difference found from molecular modeling for **17a**, we obtained an X-ray crystallographic structure of the methyl ether derivative of **17a** to firmly establish its preferred stereochemistry about the amide bond. In addition, for comparison, a crystallographic structure was determined on the less hindered *N*-(*i*-propyl)-carboxamide **10b**. ORTEP representations of these molecules are shown in Figure 5.

As can be seen in Figure 5, both the *i*-propyl carboxamide and CF₃-substituted thioamide were found to crystallize in the lower energy s-cis conformation. Prominent torsion angles are indicated in Table 1. Comparison of the amide torsional bonds (torsion A) reveals greater deviation of the N-aryl ring from the amide plane in thiocarboxamide 17a compared to 10b (18° versus 9.7°), consistent with the greater steric hindrance anticipated in the amide plane of the thioamide system. The greater distortion in the thioamide bond planarity is also evident from its improper torsion (torsion \mathbf{E}) and its N-alkyl torsion **B**. Both structures also show a twisting of the phenyl rings out of the amide plane (C and **D**), with the thioamide being more planar. This difference appears in part to be due to additional nearby nonbonding interactions with the CF₃-group in thioamide 17a.

Solution VT ¹H NMR experiments (Fig. 6) were also conducted, and the results were found to be consistent with a single amide conformer. Shown in Figure 6 is the ¹H NMR spectrum of the aromatic region for thioamide 17a at -60, +23, and $+55 \,^{\circ}\text{C}$ in MeOD. The spectra at -60 °C and room temperature are very similar. Interestingly, there is a slow rotation about the N-C bond of the N-phenyl group, which results in distinct signals for the ortho and meta ring protons of the N-phenyl ring. Assignments are based on COSY cross couplings. Upon heating to +55 °C, no new signals are observed for either the CH₃ group (not shown), which remains a doublet at all three temperatures, or the aroyl protons. However, the ortho and meta signals of the N-phenyl ring coalesce. This type of slow N-C rotation is well known in N-phenyl carboxamides and is analogous to hindered bi-phenyl rotations.16 The lack of



Figure 4. Capped-stick models of lowest energy *c-cis* and *s-trans* conformers of caboxamide 16a. This *s-cis* conformer (left) has lower energy $(\Delta H = 5.3 \text{ kcal/mol})$.



Figure 5. Molecular structures for thiocarboxamide 17a (left) and carboxamide 10b (right) (ORTEP; ellipsoids drawn at the 35% probability level).

 Table 1. Prominent torsion angles found in carboxamide 10b and thiocarboxamide 17a

Torsion	Compound 17a thioamide 1–4 atoms	Angle	Compound 10b carboxamide 1–4 atoms	Angle
A	C(2)–C(9)	18°	C(2)-C(8)	9.7°
В	C(16) - S(1)	7.8°	C(14) - O(1)	3.2°
С	C(1) - C(10)	76°	C(1) - C(9)	80°
D	C(3) - N(1)	50°	C(3) - N(1)	62°
E	N-improper torsion	5.4°	N-improper torsion	4.6°

additional signals indicates that there is a single amide conformer in solution, which we presume, on the basis of the computational and X-ray structures, to be the *s*-*cis*. Furthermore, for thioamides, it is worthy to note that the barner for *cis*-*trans* inter-conversion is generally 2– 5 kcal greater than it is for the analogous carboxamides. The traditional 'resonance model' used to rationalize the origin of the higher *cis*-*trans* barriers in thioamides has recently been challenged¹⁸ and is currently under debate.^{19–22} Nevertheless, from these studies we believe that it is reasonable to conclude that sterically hindered *N*-aryl *thio*benzamides also have a strongly preferred *s*-*cis* conformation, similar to *N*-substituted *oxo*anilides. Therefore, these systems can be considered as mimics of the *syn*-bibenzyl substructural motif.

Receptor binding affinities and structure-binding affinity relationships

Binding affinities for the estrogen receptor and octanolwater partition coefficients (Log $P^{o/w}$) determinations of the benzamides we have prepared are shown in Table 2 and are organized according to the substituent on the amide nitrogen (either *N*-alkyl/aryl or *N*-CF₃ containing alkyl/aryl). Binding affinities and Log $P^{o/w}$ determinations for phenylacetamides are presented in Table 3. Binding affinities were obtained from a competitive radiometric binding assay, using [³H]estradiol as tracer and lamb uterine cytosol as a source of ER, and they are expressed as relative binding affinities (RBA), with estradiol having an affinity of 100%. Additional binding



Figure 6. VT-¹H NMR for 17a, aromatic region.

assays of select compounds for receptor subtypes ER α and ER β , as well as ER α and ER β transcriptional activation profiles in human endometrial cancer (HEC-1) cells using an estrogen-responsive reporter gene construct, are also shown in Table 4. Log $P^{o/w}$ determinations were obtained according to the method of Minick,

Table 2. Estrogen receptor relative binding affinity RBA (estradiol = 100) and log octanol water partition coefficient (log $P^{\sigma w}$) data for *N*-phenyl-benzamides

Compound	Core structure	R group	RBA ^a	Log Pov
5a 5b 5c	-CSO ₂ R-	Et <i>n-</i> Bu Bn	0.23 0.13 0.053	2.76 3.66 4.62
10a 10b 10c 10d 10e 10f	-CONR-	Et <i>i-</i> Pr <i>n-</i> Bu <i>i-</i> Bu CH ₂ Ph CH(C ₆ H ₅)CH ₃	$\begin{array}{c} 0.062 \\ 0.009 \\ 0.042 \\ 0.020 \\ 0.089 \\ 0.45 \end{array}$	2.40 2.31
11a 11b 11c 11d 11e 11f 11g	-CSNR-	Et <i>i</i> -Pr <i>n</i> -Bu <i>i</i> -Bu CH ₂ Ph CH(C ₆ H ₅)CH ₃ CH(CH ₃)(<i>i</i> -Pr)	0.62 0.83 1.3 1.5 0.15 2.5 4.6	2.80 2.97 3.99 3.98 4.31

^aRelative binding affinities are the average of duplicate determinations (CV 0.3), measured in lamb uterine cytosol. For estradiol. RBA = 100. For details, see Experimental.

Table 3. Estrogen receptor relative binding affinity RBA (estradiol=100) and log octanol-water partition coefficient (log $P^{\sigma w}$) data for *N*-CF₃-substituted benzamides

Compound	х	R^1	R ²	R ³	RBA ^a	Log Po'u
16a	0	НО	CH ₃	p-HO	0.49	3.29
16b	0	НО	C_6H_5	p-HO	9.0	4.51
16c	0	НО	C_6H_5	́ Н	0.063	4.46
16d	0	НО	C_6H_5	<i>m</i> -HO	0.45	
16e	Ο	HO	C ₆ H ₅	o-HO	0.068	
16f	Ο	Н	C_6H_5	p-HO	8.9	4.97
16g	0	Н	m-HOC ₆ H ₄	p-HO	15	4.63
17a	S	НО	CH ₃	p-HO	7.5	3.99
17b	S	HO	C_6H_5	p-HO	14	5.40

^aRelative binding affinities are the average of duplicate determinations (CV 0.3), measured in lamb uterine cytosol. For estradiol, RBA = 100. For details, see Experimental.

using a standardized Chromegabond MC8 reverse phase HPLC system.²³

Binding affinity of *N*-phenyl benzamides. The receptor binding affinities for *N*-phenyl benzamides are shown in Table 2. The sulfonamides and carboxamides all have rather low affinity, the highest being carboxamide 10f. Within the carboxamide series (10a–f), it is interesting to compare the effect of branching near the amide core. For example, the ethyl (10a) and benzyl (10e) compounds have similarly low affinities, and likewise the branched *i*-propyl amide 10b has very low affinity. However, amide 10f, which contains both an α -phenyl and methyl substituent, shows a significant, 5- to 7-fold increase in affinity compared to either monosubstituted amide 10a or 10e. Overall, the analogous *N*-alkyl

Table 4. Estrogen receptor relative binding affinity RBA (estradiol = 100) and log octanol-water partition coefficient (log $P^{o/w}$) data for *N*-phenyl phenylacetamides

Compound	Х	R	R ²	RBA ^a	Log Po/w
21a	0	CH ₂ CH ₃	Н	0.006	2.68
22a	0	CH ₂ CH ₃	CH ₂ CH ₃	0.004	3.52
27	0	$CH(CH_3)(CF_3)$	CH ₃	0.010	_
28	S	$CH(CH_3)(CF_3)$	Н	0.66	
21b	S	CH ₂ CH ₃	Н	0.20	3.40
22b	S	CH ₂ CH ₃	CH ₂ CH ₃	0.089	4.27
22c	S	CH_2CH_3	$C\tilde{H_2}Ph$	0.34	—

^aRelative binding affinities are the average of duplicate determinations (CV 0.3), measured in lamb uterine cytosol. For estradiol, RBA = 100. For details, see Experimental.

thioamides (**11a–g**) bind with affinity 2- to 90-fold greater than their carboxamide counterparts, which is presumably due to their greater hydrophobic character, although there are not great differences in the Log $P^{o/w}$ values, where comparisons can be made. A related trend is also observed in other non-steroidal ligands containing heteroatoms, such as benzo[b]furans and benzo-[b]thiophenes, with benzo[b]thiophenes displaying higher affinity than benzo[b]furans.^{24,25}

The beneficial effect of placing more sterically encumbered groups near the amide core on receptor affinity is also evident in the thioamide series, because the same relative affinity increases which occurred for carboxamides (10a < 10e < 10f) are also reflected in the thioamides (11e < 11a < 11f). Replacement of the phenyl substituent in 11f with an *i*-propyl group to afford thioamide 11g results in even greater crowding near the amide plane, enhancing the relative affinity to 4.6%, the highest observed for any simple *N*-phenyl benzamide.

Binding affinity of CF₃-containing *N*-phenyl benzamides. The RBA data for the CF₃-containing benzamides are presented in Table 3. Immediately apparent is the relatively high binding affinity for thioamide 17b and carboxamide 16g, and to a lesser extent amides 16b, 16f and 17a. Compound 16a, reported by Hartmann and coworkers, is included for comparison.⁶ In general, all the CF₃-substituted benzamides show affinity enhancements relative to their non-fluorinated amide counterparts (Table 2).

Shown in Figure 7 are four direct comparisons of RBA and Log $P^{O/W}$ data for benzamides from Tables 2 and 3. These four pairs of compounds clearly show the relationship between core structure (*O* and *S* substitution) and peripheral group presentation (methyl to phenyl, and methyl to trifluoromethyl) and how these components alter ER affinity.

Comparison of the relative RBAs for these four pairs of amides, starting from the *i*-propyl pair (upper left) to the CF₃-containing phenethyl pair (lower right), shows a diminishing effect of sulfur–oxygen replacement as binding affinity increases and as larger substituents are added near the amide core. The sulfur–oxygen



Figure 7. Comparison of the relative binding affinities of the principal carboxamides and thiocarboxamides.

enhancement is greatest in the parent *i*-propyl system (10b versus 11b), where the overall affinity is lowest, and is least when the substituents are phenyl and trifluoromethyl (16b versus 17b), where the overall affinity is highest. In addition, there is a general increase in binding affinity with increasing lipophilicity. The observed parallel affinity—lipophilicity increase within the series is not too surprising in light of what is known about the hydrophobic character of the ER ligand binding pocket.²⁶

CF₃ effect: lipophilic, steric or electrostatic effects? The effect of adding a CF₃ group in place of methyl on the RBA may be the result of several factors, including lipophilic, steric and/or electrostatic effects. In terms of steric effects, a recent review by O'Hagan and Rzepa²⁷ cites several studies which place an upper size limit of a CF₃-group being close to that of *i*-propyl. The greatest increases in RBA for CH₃ to CF₃ substitution are found in the carboxamide class (20- to 50-fold, versus 5- to 8fold for thioamide), with the highest being the *i*-propyl compounds (Fig. 7, 16a versus 10b). In order to test whether or not RBA increases associated with CF₃replacement are the result of its steric or its electronic property, the branched $CH(CH_3)(i-Pr)$ thioamide 11g was prepared to compare with the CF_3 analogue 17a. Based on the above-mentioned studies, if steric effects alone operate, compound 11g would be expected to have an RBA close to that of the CF3-substituted thioamide 17a (7.5%). The observed RBA, however, for compound 11g was only 4.6%. This indicates that an additional effect may be contributing to the binding energy, either lipophilic or electrostatic. The fact that the Log $P^{o/w}$ values for **11g** and **17a** are quite similar, with **11g** being somewhat higher, suggests that the difference in binding affinity is probably not due to a lipophilic component. Thus, based on this data it would seem that the additional component has, in addition to its steric effect, an electronic effect as well.

Optimization of binding orientation and hydroxylation pattern. For unsymmetrical non-steroidal ER ligands, a common approach to determining which phenol is imitating the crucial A-ring phenol of estradiol is to prepare and test the corresponding mono-phenolic derivatives.²⁸ The highest affinity mono-phenolic analogue is then presumed to be the A-ring mimic, because the hydroxyl substituent at this position is known to be very important in binding to the ER. Many nonsteroidal ER ligands benefit from a second hydroxyl hydrogen bonding partner and appear to do so in an anti-bibenzyl motif, which places the hydroxyl oxygenoxygen interatomic distances approx. 10-12 Å. Thus, it was apparent to us that the rigid s-cis benzamide conformation (syn-bibenzyl motif) in our amides was likely not benefiting from a second hydrogen bonding interaction, because its hydroxyl oxygen-oxygen interatomic distance was only 7.4 A. The importance for this second hydrogen bond is evident in the recent non-steroidal diethylstilbestrol and raloxifene ERa LBD crystal structures.^{26,29} Both of these complexes have an expected A-ring phenol mimic and a second phenol imitating the D-ring 17β-hydroxyl group of estradiol. In both cases, the second phenol uses the same hydrogen bond interaction with His_{524} as does the 17 β -hydroxy1 group of estradiol.

To obtain an optimal hydroxylation pattern and discern which ring phenol was imitating the A-ring phenol of estradiol, we prepared benzamides 16c-g. The results of these studies (Table 3) clearly show a pattern indicative of a preferred A-ring phenol mimic. For example, removal of the 4-hydroxyl phenol on the benzamide ring and retention of the 4-hydroxyl of the N-phenyl in compound 16c lead to a nearly complete loss in binding affinity compared to the parent di-hydroxy compound **16b** (RBA = 0.063% versus 9.0%). In contrast, removal of N-phenyl hydroxyl and retention of the benzamide 4-hydroxyl result in an analogue 16f, which retains a binding affinity equivalent to that of the parent compound. These findings clearly implicate the benzoyl ring phenol as the A-ring mimic of estradiol. In addition, as might be expected, positioning the hydroxyls elsewhere on the benzoyl ring is not well tolerated, as both the ortho- and meta-hydroxy analogues 16d and 16e have little or no affinity for the receptor.

Having discerned which phenol in the benzamide system was important for binding, we wished to explore alternative positions for a second hydroxyl group in order to benefit from the second hydrogen bonding interaction which is found in the raloxifene and DES ER binding pockets. Depicted in Figure 8 is a schematic representation of the predicted binding mode and phenol interactions for benzamide 16f in the binding pocket of ER. This rudimentary model was derived from the interactions that were described to be present in the estradiol (E_2) X-ray crystal structure in the ER α LBD, the coordinates of which, at the time of our consideration, were not released.²⁶ Based on this view, we proposed that a hydroxyl substituent on the phenethyl ring might be positioned in the binding subpocket near His524.

The choice of positioning the second OH at the *meta* position on the phenethyl substituent seemed somewhat obvious, since modeling showed that alternative OH substitution on the N-phenyl ring fails to provide O·O interatomic distances much greater than in the parasubstituted system 16b. Furthermore, the binding mode depicted in Figure 8 actually places the N-phenyl group more closely in the 11ß subpocket of the ligand binding site, rather than the D-ring subpocket. In choosing the ring-substitution for the hydroxyl on the phenethyl ring, we considered both the para and meta isomers. However, in light of the problems encountered with quinone methide elimination in the N-benzyl benzamide series bearing a *para*-methoxy protecting group, we opted not to prepare the *para*-hydroxy analogue of **16f**, since the electron withdrawing CF₃ group would presumably exacerbate quinone methide elimination.

With the recent release of the $E_2 \ ER\alpha \ LBD$ crystal structure coordinates, we performed molecular docking studies of the proposed *meta*-hydroxybenzamide (**16g**) using TRIPOS' Flexidock routine (see Experimental), to see whether or not the proposed binding mode in Figure 8 was reasonable and whether a *meta*-hydroxy derivative could participate in a hydrogen bonding interaction with His₅₂₄.

The final docked and minimized model, using the *S*enantiomer of **16g**, is depicted in crossed-stereo in Figure 9. The *R*-enantiomer (not shown) was also modeled in the same manner but resulted in a higher energy complex. The final *S*-enantiomer model converged to an RMS of < 0.05 kcal/mol·Å with a reasonably favorable binding energy. In this conformation, the benzoyl phenol is comfortably accommodated in the A-ring pocket, and the *N*-phenyl group projects into what

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Figure 8. Putative binding mode for N-phenyl benzamide template with A-ring mimic interactions and a proposed site for second HO-group.





Figure 10. Depiction of solvent accessible surfaces for 16g in ER α binding pocket.

would resemble the 11 β pocket of the ER α -E₂ structure. Interestingly, the amide bond occupies a region of the binding pocket similar to that of the isosteric ethylene unit of DES in the recently solved ER α -DES crystal structure.²⁹

Using the above model, we generated Connolly solvent accessible surfaces for the ligand and binding pocket to visualize close contacts (Fig. 10). This view reveals mostly favorable interactions within acceptable van der Waals radii. The ligand itself has a volume of 296 Å³, which is 51 Å³ greater than E_2 , but is well under the total volume of 450 Å³ for the binding pocket as reported in the ER α -E₂ crystal structure.²⁶

We were grateful to find that the bis-(phenolic) benzamide **16g** had an RBA of 15%, the highest of any carboxamide ER ligand. This value represents a 1.6-fold increase over the affinity of the corresponding monophenolic benzamide **16f** and is close to what one might expect: in E_2 and other non-steroidal ER ligand systems, removal of the 17 β hydroxyl, as in E_2 , or the second D-ring hydroxyl mimic, as in raloxifene, result in a 0.6 kcal/mol reduction in binding energy, which corresponds to a 3-fold drop in affinity.³⁰ The fact that the addition of the second hydroxyl does not give a full 3fold increase may be due to a somewhat unfavorable hydrogen bond trajectory in the binding pocket. From our model, the distance between the *m*-hydroxyl substituent and His₅₂₄ (3.69 Å) is sufficiently close for some hydrogen bonding interaction, but is not ideal either in terms of distance or geometry. A second more likely explanation may have to do with stereochemical issues: This compound (l6g) was tested as a racemate, so only one enantiomer is likely to be able to benefit from the second hydrogen bonding interaction. The ER α and $ER\beta$ RBAs and the transcriptional activation profile for *rac*-16g in HEC-l cells are discussed below.

N-Phenyl acetamides as ER ligands. The RBA and Log $P^{o/w}$ data for *N*-alkyl- and CF₃-containing acetamides are shown in Table 4. Disappointingly, none of these compounds were found to show appreciable affinity for the ER. Several direct comparisons can be made to other series in this investigation, which suggest that acetamides may not provide viable scaffolds for further combinatorial consideration, despite their potential to support greater structural diversity than the benzamides. Even the CF₃-containing acetamide **27** and thioacetamide **28** have affinities less that 1%, which are 50- and 11-fold lower, respectively, than their benzamide counterparts **16a** and **17a**.

One possible explanation for the low affinity in the acetamide class may lie in their conformation. Monte Carlo conformational studies of compound **27** and the unsubstituted analogue **27b**, using molecular mechanics methods, show a distinct preference for a *syn* conformation, as shown in Figure 11. The *syn*-conformer for **27** is predicted to be 5 kcal/mol lower in energy than the *anti*-conformer; even without the α -methyl substituent, which eliminates possible alkyl-*N*-phenyl steric repulsions, the *syn*-conformer is predicted to be favored by 2.5 kcal/mol (not shown). It is also noted that the barrier to rotation about the acetyl bond was estimated



Figure 11. Conformational analysis of N-penyl-penylacetamide 27.

to be between 4 and 5 kcal/mol for **27**, based on MM2 dihedral drive calculations.

As in the benzamides, in the acetyl conformations of 27 the *exo*-rule is upheld; this places the *N*-phenyl group *exo* with respect to the amide C=O, making the amide configuration *s*-*cis*. This *exo*-effect may also be operative for the α -phenyl substituent as well, since it also prefers to be essentially *exo* to the amide C=O rather than *endo*. Since the hydroxyl O·O interatomic distances for many estrogens requires a minimum of approximately 10.5 Å, the *syn*-conformation is excluded as a structure that is likely to fit in the ER binding pocket. Based on these studies and the RBA data available at this time, further investigations of the acetamide series did not appear to be warranted.

$ER\alpha/ER\beta$ binding affinity, transcriptional activity and enantioselectivity

Shown in Table 5 are the RBA values of select benzamides for purified ER α and ER β , along with their transcriptional activity in HEC-1 cells using an estrogenresponsive reporter gene construct ((ERE)₃-pS2-CAT) with expression vectors for either $ER\alpha$ or $ER\beta$ and benzamide at 10⁻⁶ M. It is apparent that binding affinities show selectivity for ERa, mostly in the range of 2- to 8-fold; compound 16g has the highest selectivity for ERa of 21-fold (27% versus 1.3%). The enantiomers of 17b, which were separated using chiral HPLC (ChiralPak AD column), also showed interesting RBA differences between the two receptor subtypes. For example, enantiomer ent₂-17b was found to have greater selectivity for ER α than enantiomer *ent*₁-17b by more than 5-fold, indicating that the individual binding pockets for ER α and ER β are uniquely sensitive to subtle changes in ligand shape.

In the transcription assays,³¹ agonism is measured with 1 μ M of compound alone and is expressed as the percent of transcriptional activity with 1 nM E₂ (high values indicate agonist activity); antagonism is measured with 1 μ M compound together with 1nM E₂ (low values indicate antagonist activity). All of the benzamides tested were full or nearly full agonists through ER α ,



Figure 12. Transcription activation by ER α and ER β in response to benzamide **16g**. Human endometrial cancer (HEC-1) cells were transfected with expression vectors for ER α or ER β and an (ERE)₃-pS2-CAT reporter gene and were treated with the indicated concentrations of benzamide **16g** for 24 h. CAT activity was normalized for β -galactosidase activity from an internal control plasmid. Values are the mean \pm SD for three or more separate experiments, and are expressed as a percent of the ER α or ER β response with 1 nM E₂.

whereas some were effective partial antagonists through ER β . Compound **16g** was a full agonist on ER α and ER β .

Because of the high activity of amide 16g at 10^{-6} M on both ER α and ER β , a full dose-response curve was determined to see whether any potency selectivity existed for either ER α or ER β . As shown in Figure 12, benzamide 16g behaves as an ER α potency selective agonist, displaying a 500-fold higher potency for activation of ER α over ER β . It is of note that this high potency selectivity for ER α is greater than the 21-fold ER α affinity selectivity of this compound (cf. Table 5).

Conclusions

Through an analysis of potent non-steroidal estrogens, we have identified common substructural bibenzyl and homobibenzyl motifs in the structure of ER ligands. Based on this simple pharmacophore model, we have

Table 5. Benzamide $ER\alpha$ and $ER\beta$ binding affinities and transciptional activity

Compound	RBA (%) (uterine)	RBA (%) ^a		Agonism ^b % CAT Act.		Antagonism ^b % CAT Act. + E ₂		
		ERα	ERβ	α/β ratio	ERα	ERβ	ERα	ERβ
11g	4.6	14	2.4	5.8	65	20	90	40
17a	7.5	23	15	1.5	60	20	65	45
16b	9.0	17	2.7	6.3	75	15	90	35
rac-17b	14	25	8.9	2.8	80	30	83	35
ent ₁ -17b	10	18	11	1.6	85	30	80	48
ent17b	22	40	4.8	8.3	80	10	90	40
16g	15	27	1.3	21	99	99	100	97

^aRelative binding affinities are the average of duplicate determinations (CV 0.3), determined with purifed RD α and ER β preparations. For estradiol, RBA = 100. For details, see Experimental.

^bHEC-1 cells were transfected with either RE α or ER β expression vectors and an (ERE)₃-pS2-CAT reporter gene. CAT activity was normalized for β -galactosidase activity from an internal control plasmid and is expressed as a percent of the ER α or ER β response with 1 nM E₂. Benzamides were tested at 10⁻⁶ M concentration to test for agonism and at 10⁻⁶ M with 10⁻⁹ M E₂ to test for antagonism. These assays are generally reproductive with a CV of 30% relative.

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made a substantial exploration of the amide functionality as a core replacement that might support compound libraries for developing novel SERMs. Using this approach, we prepared and tested representative examples of bis-phenolic amides which displayed both simple N-alkyl substituents and CF₃-containing N-alkyl substituents. Of the two classes examined, benzamides and acetamides, the benzamides displayed the highest affinity, in particular the CF₃-containing benzamides. The recent X-ray crystal structure of E₂ bound to ERa and structure-binding affinity relationships have allowed us to design a bis-phenolic benzamide 16g, which has the highest affinity for any carboxamide ligand for the ER. This compound is also an ER α potency-selective agonist in cell-based assay of ER-mediated transcriptional activity. This work illustrates that ER ligands based on simple amide core structures can be readily prepared. High affinity ER binding, however, requires an appropriate distribution of bulk and functionality. The strong conformational preference of the basic anilide function in all of these ligands defines the scope of further structural and functional expansion of these series.

Experimental

General

Melting points were determined on a Thomas-Hoover UniMelt capillary apparatus and are uncorrected. All reagents and solvents were obtained from Aldrich, Fisher or Mallinckrodt. Tetrahvdrofuran was distilled from sodium/benzophenone. Dimethylformamide was vacuum-distilled prior to use, and stored over 4Å molecular sieves. n-Butyllithium was titrated with Npivaloyl-o-toluidine. All reactions were performed under a dry N₂ atmosphere unless otherwise specified. Reaction progress was monitored by analytical thinlayer chromatography using GF silica plates purchased from Analtech. Visualization was achieved by shortwave UV light (254 nm) or potassium permanganate. Radial preparative-layer chromatography was performed on a Chromatotron instrument (Harrison Research, Inc., Palo Alto, CA) using EM Science silica gel Kieselgel 60 PF₂₅₄ as adsorbent. Flash column chromatography was performed using Woelm 32-63 µm silica gel packing.

Logarithms of octanol-water partition coefficients (Log $P^{o/w}$) were determined using a standardized reverse phase HPLC Chromegabond MC 8 column.²³ HPLC separation of the enantiomers for 17b was performed on an analytical ChiralPak AD (4.6×25 cm) column from Chiral Technologies using 10% i-propanol/hexane as solvent. ¹H and ¹³C NMR spectra were recorded on either a General Electric QE-300 (300 MHz), Varian Unity 400 or 500 MHz spectrometers using CDCl₃, MeOD or $(CD_3)_2SO$ as solvent. Chemical shifts were reported as parts per million downfield from an internal tetramethylsilane standard ($\delta = 0.0$ for ¹H) or from solvent references. NMR coupling constants are reported in Hertz. ¹³C NMR were determined using either the Attached Proton Test (APT) experiment or standard ¹³C pulse sequence parameters. Low resolution and high

resolution electron impact mass spectra were obtained on Finnigan MAT CH-5 or 70-VSE spectrometers. Elemental analyses were performed by the Microanalytical Service Laboratory of the University of Illinois.

Unless otherwise stated, a standard procedure for product isolation was used; this involved quenching by addition of water or an aqueous solution, exhaustive extraction with an organic solvent, washing the extracts, drying over Na₂SO₄, and solvent evaporation under reduced pressure. Quenching media, extraction solvents, and aqueous washes used are noted parenthetically after the phrase 'product isolation'.

Biological procedures

Relative binding affinity assay. Ligand binding affinities (RBAs) using lamb uterine cytosol as a receptor source were determined by a competitive radiometric binding assay using 10 nM [³H]estradiol as tracer and dextrancoated charcoal as an adsorbant for free ligand.³² Purified ER α and ER β binding affinities were determined using a competitive radiometric binding assay using 10 nM [³H]estradiol as tracer, commercially available ER α and ER β preparations (PanVera Inc., Madison, WI), and hydroxylapatite (HAP) to adsorb bound receptor-ligand complex.³³ HAP was prepared following the recommendations of Williams and Gorski.³⁴ All incubations were done at 0°C for 18-24 h. Binding affinities are expressed relative to estradiol on a percent scale (i.e. for estradiol, RBA = 100%). All essays were run in separate, duplicate experiments which were reproducible with a coefficient of variation of less than 30% relative.

Transcriptional activation assay. Human endometrial cancer (HEC-l) cells were maintained in culture and transfected as described previously.³¹ Transfection of HEC-1 cells in 60 mm dishes used 0.4 mL of a calcium phosphate precipitate containing 2.5 μg of pCMVβGal as internal control, 2 μg of the reporter gene plasmid, 100 ng of the ER expression vector, and carrier DNA to a total of 5 μg DNA. CAT activity, normalized for the internal control β-galactosidase activity, was assayed as previously described.³¹

Molecular modeling

Small molecule modeling. Dihedral drive and Monte Carlo conformational searches for *N*-phenyl benzamides 16a and 17a and *N*-phenyl acetamide 27 were conducted using the MM2 force field as implemented in Macromodel v.5.5 with CHCl₃ as a solvent model. All generated conformers from Monte Carlo searches underwent full matrix assisted minimization using the PRCG function with a convergence criteria of 0.001 kcal/mol. Dihedral drives were conducted in 5° steps with a convergence criteria of 0.05 kcal/mol using 1000 minimization steps. Analysis of the energy versus torsion angle provided the estimated rotational barrier.

Receptor docking studies. The starting conformation used for docking studies for 16g was obtained from a

random conformational search using the TRIPOS' force field (MAXIMIN) as implemented in the program SYBYL, version 6.5.2 (Tripos Inc., St. Louis, MO). The cis global minima obtained for benzamide 16g was then overlaid with E_2 in the E_2 -ER α LBD crystal structure²⁶ using a least squares multifitting of five atoms: atoms C(3), C(10), C(9), C(11), and C(12) of E_2 were fitted with the 1 and 4 carbons of the benzamide phenyl ring. the C=O carbon, the nitrogen and the C(2) carbon of the phenethyl group, respectively. The pre-positioned benzamide was then optimally docked in the ER α binding pocket using TRIPOS' Flexidock. Both hydrogenbond donors and acceptors within the pocket surrounding the ligand (Glu₃₅₃, Arg₃₉₄ and His₅₂₄) and the ligand itself, in addition to select torsional bonds were defined. The best docked receptor ligand complex from Flexidock then underwent a three-step minimization: first non-ring torsional bonds of the ligand were minimized in the context of the receptor using the torsmin command, followed by minimization of the sidechain residues within 8 Å of the ligand while holding the backbone and residues Glu₃₅₃ and Arg₃₉₄ fixed. A final third minimization of both the ligand and receptor was conducted using the anneal function (hot radius 8A, interesting radius 16Å) while holding residues Glu_{353} and Arg₃₉₄ fixed to afford the final model. Minimizations were done using the TRIPOS' Forcefield (MAXI-MIN) with the Powell gradient method and default settings (final RMS ≤ 0.05 kcal/mol·A).

X-Ray crystallography

Cystallography details for 17a and 10b. Crystals of 17a were obtained by slow evaporation from ether at 0° C. Crystals were mounted on glass fibers with Paratone-N oil (Exxon) and immediately cooled to -75° C in a cold nitrogen gas stream on the diffractometer. Standard peak search and indexing procedures gave rough cell dimensions, and least squares refinement using 5344 reflections yielded the cell dimensions as given in Table 6.

Data were collected with an area detector by using the measurement parameters listed in Table 6. No systematic absences are noted for the P-1 space group. The measured intensities were reduced to structure factor amplitudes and their esds by correction for background, scan speed, and Lorentz and polarization effects. While corrections for crystal decay were unnecessary (the data were corrected for crystal decay), a Ψ -scan absorption correction was applied, the maximum and minimum transmission factors being 0.991 and 0.876. Systematically absent reflections were averaged to yield the set of unique data. All 5344 data were used in the least squares refinement.

The structure was solved using direct methods (SHELXTL). The correct positions for the C, N, O, S, and F atoms were deduced from an E-map. Subsequent least-squares refinement and difference Fourier calculations revealed the positions of the remaining nonhydrogen atoms. The quantity minimized by the leastsquares program was $\Sigma w (F_0^2 - F_c^2)^2$, where $w = \{[\sigma(F_0^2)]^2 + (0.883P)^2\}^{-1}$ and $P = (F_0^2 + 2F_c^2)/3$. The analytical approximations to the scattering factors were used, and all structure factors were corrected for both real and imaginary components of anomalous dispersion. In the final cycle of least squares, independent anisotropic displacement factors were refined for the non-hydrogen atoms and the aromatic, methyl, and methine hydrogen atoms were fixed in 'idealized' positions with C-H=0.95 Å for the aromatic hydrogens, C-H=0.98 Å for the methyl hydrogens, and C-H=1.00 Å for the methine hydrogen. Successful convergence was indicated by the maximum shift/error of 0.001 for the last cycle. Final refinement parameters are given in Table 6. The largest peak in the final Fourier difference map $(0.795 \text{ e} \text{\AA}^{-3})$ was located 1.64 Å from C(18). A final analysis of variance between observed and calculated structure factors showed no apparent errors.

Single crystals of **10b** were grown from MeOH by slow evaporation at 23 °C in a partially sealed scintillation

Table 6.	 Crystal data and 	structure refinement for	thiocarboxamide	17a and ca	arboxamide 10b
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Complex	17a	
Empirical formula	$C_{18}H_{18}F_{3}NO_{2}S$	$C_{16}H_{17}NO_3$
fw	369.39	271.31
Temperature	198(2) K	198(2) K
Wavelength	0.71073 Å	1.54178 Å
Cryst syst	triclinic	monoclinic
Space group	P-1	$P2_1/n$
Unit cell dimensions	a = 9.5099(7) Å	a = 18.248(4) Å
	b = 10.5351(8) Å	b = 9.054(2) Å
	c = 11.1127(8) Å	c = 18.248(4) Å
Volume	$868.07(11) A^3$	2916.4(10) Å ³
Z	2	8
Density, caled	$1.413 \mathrm{Mg}\mathrm{m}^3$	$1.236 \mathrm{Mg/m^3}$
Abs coeff	$0.228 \mathrm{mm}^{-1}$	0.695 mm ¹
No. of indep refins	5344 [R(int) = 0.0293]	4796 (R(int) = 0.0556)
Refinement method	full matrix least-squares on F^2	full matrix least-squares on F^2
No. of data/restraints/params	3727/0/226	2727/0/362
Goodness-of-fit (GooF) on F^2	0.949	1.062
Final R indices $(I \ge 2\sigma(I))$	$R1 \approx 0.0502$, $wR2 = 0.1393$	$R_1 = 0.0624, wR_2 = 0.1662$
R indices (all data)	R1 = 0.0892, WR2 = 0.1533	R1 = 0.0767, WR2 = 0.1881

vial. Crystals were mounted on glass fibers with Paratone-N oil (Exxon) and immediately cooled to -75 °C in a cold nitrogen gas stream on the diffractometer. Standard peak search and indexing procedures gave rough cell dimensions. Initial inspection of the data suggested a C-centered orthorhombic cell, but no successful solution could be found in any orthorhombic space group. Eventually, a successful solution was found in a primitive monoclinic cell of half the volume in the space group $P2_1/n$. The refinement stalled at $wR_2 = 0.63$, despite having located all atoms and refining the nonhydrogen atoms anisotropically. The unusual equality in the lengths of the a and c axes, combined with the observation that $F_{obs} \gg F_{calc}$ for all the reflections in the 'most disagreeable reflections' list, suggested that the crystal was twinned. A twin law involving mirror reflection through the (101) plane was assumed, and a refinable parameter describing the relative volume fractions of the two twin individuals was added to the model. The intensities were calculated from the relation $F_{TOT}^2 = f$ $F_{hkl}^2 + (1-f) F_{h'k'l'}^2$, where f is the volume fraction of twin individual one and F_{hkl}^2 and $F_{h'k'l'}^2$ are the contributions from the two twins (the twin law specifies that h'k'l' = lkh). The scale factor refined to 0.500(3), and the wR_2 factor dropped to an acceptible value of 0.185.

In the final cycle of least squares, independent anisotropic displacement factors were refined for the nonhydrogen atoms and the aromatic, methyl, and methine hydrogen atoms were fixed in 'idealized' positions with C-H=0.95 Å for the aromatic hydrogens, C-H=0.98 Å for the methyl hydrogens, and C-H=1.00 Å for the methine hydrogen. Successful convergence was indicated by the maximum shift/error of 0.001 for the last cycle. Final refinement parameters are given in Table 6. The largest peak in the final Fourier difference map (0.795 cÅ^{-3}) was located 1.64 Å from C(18). A final analysis of variance between observed and calculated structure factors showed no apparent errors.

Chemical syntheses

4-Methoxyphenyl-4-methoxybenzene sulfonamide (3).⁸ To a mixture of 4-methoxybenzenesulfonylchloride (1.0 equiv) and pyridine (1.1 equiv) in CH₂Cl₂ (0.5 M) at 0 °C was added dropwise to a solution of *p*-anisidine (1.5–5.0 equiv) in CH₂Cl₂ (1.5 M) over 30 min. The reaction mixture was warmed to rt and stirred until completion of reaction as indicated by TLC. Product isolation (H₂O, 2 N HCl) and recrystallized from EtOH afforded product as light red crystals (91%): mp 91–92.5 °C, mp⁸ 93 °C.

General procedure for phase transfer *N*-alkylation of sulfonamides.⁹ To a CH_2Cl_2 (0.06 M) solution of sulfonamide (1.0 equiv) was added an equal volume solution of NaOH (5 equiv) and $(n-Bu)_4NSO_4H$ (0.1 equiv) in H₂O. Alkylhalide (2.0 equiv) was then added directly to the bi-phasic solution and stirred at rt. After 12 h the layers were separated. Product isolation (Na₂CO₃, brine) followed by flash chromatography afforded the pure *N*-alkyl benzene sulfonamide.

N-(4-Methoxyphenyl)-*N*-(ethyl)-4-methoxybenzene sulfonamide (4a).⁸ Purification by flash chromatography (20% EtOAc:hexane) afforded off-white powder (86%): mp 95–7 °C, mp⁸ 100–104 °C; ¹H NMR (CDCl₃) δ 1.06 (t, 311, *J* = 7.1 Hz), 3.52 (q, 2H, *J* = 7.0 Hz), 3.81 (s, 3H), 3.84 (s, 3H), 6.82 (d, 211, *J* = 8.8 Hz), 6.94 (d, 2H, *J* = 8.0 Hz), 6.96 (d, 2H, *J* = 8.0 Hz), 7.54 (d, 2H, *J* = 8.8 Hz).

N-(4-Methoxyphenyl)-*N*-(butyl)-4-methoxybenzene sulfonamide (4b). Purification by flash chromatography (20% EtOAc:hexane) afforded transparent oil (81%): ¹H NMR (CDCl₃) δ 0.85 (t, 3H, *J*=7.2 Hz), 1.34 (m, 4H), 3.46 (t, 2H, *J*=6.8 Hz), 3.80 (s, 3H), 3.86 (s, 3H), 6.81 (d, 2H, *J*=9.0 Hz), 6.92 (app t, 4H, *J*=9.1 Hz), 7.51 (d, 2H, *J*=9.0 Hz); MS (EI, 70 eV) *m*/*z* 349 (M⁺); HRMS calcd for (C₁₈H₂₃NO₄S, 349.1347, found 349.1348).

N-(4-Methoxyphenyl)-*N*-(benzyl)-4-methoxybenzene sulfonamide (4c). Recrystallization from 20% EtOAc/hexane afforded flocculent white crystals (93%): mp 141–142 °C; ¹H NMR (CDCl₃) δ 3.72 (s, 3H), 3.87 (s, 3H), 4.67 (s, 2H), 6.89 (d, 2H, *J*=11 Hz), 6.86 (d, 2H, *J*=11.5 Hz), 6.94 (d, 2H, *J*=11 Hz), 7.21 (m, 5H), 7.59 (d, 2H, *J*=11 Hz); ¹³C NMR (CDCl₃) δ 55.1, 55.4, 55.8, 114.2, 127.7, 128.5, 128.7, 129.9, 130.4, 130.4, 131.7, 136.3, 158.9, 163.0; HRMS calcd for C₂₁H₂₁NSO₄, 383.1191, found 383.118.

General demethylating procedure using BBr₃. To a solution of methyl ether compound in CH_2Cl_2 at -78 °C was added a 1.0 M BBr₃ in CH_2Cl_2 (3 equiv) dropwise over 15 mm. The reaction was allowed to reach rt and stir overnight or until disappearance of starting material as indicated by TLC. The mixture was then cooled to 0 °C. Product isolation (H₂O, Et₂O, brine) and purification via radial or flash chromatography or recrystallization from an appropriate solvent afforded the desired phenolic compounds.

General demethylating procedure using EtSH. To a stirred DMF solution of NaH (6.2 equiv of a 60% w/w dispersion) at 0°C was added dropwise 6.0 equiv of EtSH. After 30 min the mixture was warmed to rt and a solution of the methyl ether protected compound in 5 mL DMF was added dropwise. The reaction mixture was heated to reflux for 4 h; then cooled in an ice-bath and acidified with 2 N HCl. Remaining product isolation (EtOAc, brine) and purification afforded the phenolic compounds.

N-(4-Hydroxyphenyl)-*N*-(ethyl)-4-hydroxybenzene sulfonamide (5a). Deprotected according to BBr₃ procedure. SiO₂ plug (30% EtOAc/hexane) to afford off-white crystals (62%): ¹H NMR ((CD₃)₂SO) 6 0.98 (t, 3H, J=7.2 Hz), 3.43 (q, 2H, J=6.8 Hz), 6.67 (d, 2H, J=8.7 Hz), 6.76 (d, 2H, J=8.7 Hz), 6.86 (d, 2H, J=8.5 Hz), 7.34 (d, 2H, J=8.7 Hz); ¹³C NMR ((CD₃)₂SO) δ 14.1 (CH₃), 45.2 (CH₂), 115.7 (ArCH), 115.8 (ArCH), 122.6 (ArC), 128.1 (ArC), 129.9 (ArCH), 130.1 (ArCH), 157.1 (ArC), 161.6 (ArC); MS (EI, 70 eV) *m*/z 293 (M⁺); HRMS calcd for C₁₄H₁₅NSO₄, 293.0721, found 293.0722. **N-(4-Hydroxyphenyl)-***N*-(**butyl)-4-hydroxybenzene sulfonamide (5b).** Deprotected according to BBr₃ procedure to afford tan crystals, recrystallized from EtOAc/hexane (71%): mp 137–138 °C; ¹H NMR (CDCl₃) δ 0.86 (t, 3H, *J*=7.0 Hz), 1.35 (m, 4H), 3.48 (t, 2H, *J*=7.0 Hz), 6.42 (d, 2H, *J*=8.8 Hz), 6.87 (app t, 4H, *J*=8.9 Hz), 7.47 (d, 2H, *J*=9.0 Hz) ¹³C NMR (CD₃)₂SO) δ 13.8 (CH₃), 19.2 (CH₂), 29.9 (CH₂), 49.7 (CH₂), 115.6 (ArCH), 115.7 (ArCH), 122.6 (ArCH), 128.0 (ArC), 129.9 (ArCH), 130.0 (ArC), 157.0 (ArC), 161.6 (ArC); MS (EI, 70 eV) *m*/*z* 321 (M⁺); HRMS calcd for C₁₆H₁₉NsO₄, 321.1034, found 321.1035. Anal. (C₁₆H₁₉N·0.3 H₂O): C, 58.81; H, 6.05; N, 4.29. Found: C, 58.60; H, 6.03; N, 4.13.

N-(4-Hydroxyphenyl)-*N*-(benzyl)-4-hydroxybenzene sulfonamide (5c). Deprotected according to BBr₃ procedure to afford title compound after purification by radial chromatography (10%): ¹H NMR (CDCl₃) δ 4.65 (s, 2H), 6.66 (d, 2H, *J*=8.5 Hz), 6.80 (d, 2H, *J*=8.5 Hz), 6.96 (d, 2H, 8.8), 7.21 (br s, 5H), 7.59 (d, 2H, *J*= 8.9 Hz); MS (E1, 70 eV) *m*/*z* 355 (M⁺); HRMS calcd for C₁₉H₁₇NSO₄, 355.0878, found 355.0881.

N-(4-Methoxyphenyl)-4-methoxybenzamide (6).³⁵ A CH₂ Cl₂ solution of *p*-anisidine (1.3 equiv) was added dropwise to a stirring CH₂Cl₂ solution (2–5 mL) of pyridine (1.1 equiv) and *p*-anisoyl chloride (1.0 equiv) at 0 °C. Upon complete addition of amine (30 mm) mixture was allowed to reach rt, then re-cooled and vacuum filter to afford the crude carboxamide. Recrystallization from EtOH/benzene afforded the pure carboxamide as off-white crystals (72%): mp 199-200.5 °C, mp³² 202-203 °C.

N-(1-Phenethyl)-4-methoxyaniline (7).⁷ To a 0°C EtOH solution containing *p*-anisidine (1.0 equiv) and catalytic p-toluene sulfonic acid H₂O (0.05 equiv) was added freshly distilled benzaldehyde dropwise. The mixture was allowed to reach rt and complete imine formation occurred within 0.5 h. The crude mixture was concentrated in vacuo, and product isolation followed (EtOAc, brine) to afford a dark brown powder. ¹H NMR confirmed imine which was directly used in the next step without further purification. CH₃ addition: Crude imine was dissolved in toluene and cooled to -78 °C. To this was added MeLi (1.0 M toluene solution, 3 equiv) dropwise over 15 mm. After 60 min at -78 °C, the crude mixture was warmed to rt and stirred for an additional 2h. The reaction was then cooled in an ice-bath and quenched with cold MeOH followed by addition of satd NH₄Cl until neutral pH. Concentration in vacuo followed by remaining workup (EtOAc, H₂O, brine) afforded amine as a red oil. Kügelrohr distillation provided pure product, which crystallized on standing as a red orange powder (75%): ¹H NMR (CDCl₃) δ 1.50 (d, 3H, J = 7.0 Hz), 3.69 (s, 3H), 4.41 (q, 1H, J = 7.0 Hz), 6.47 (AA'BB', 2H, J = 9.0, 3.0 Hz), 6.69 $(AA'BB', 2H, J=9.0, 3.5 Hz), 7.29 (m, 5H); {}^{13}C NMR$ (CDCl₃) & 25.4 (CH₃), 54.5 (CH), 55.9 (CH₃), 114.7 (ArCH), 114.9 (ArCH), 126.1 (ArCH), 127.0 (ArCH), 128.8 (ArCH), 141.7 (ArC), 145.7 (ArC), 152.1 (ArC); MS (El, 70 eV) m/z 227 (M⁺); anal. (C₁₅H₁₇NO): C,

79.26; H, 7.54; N, 6.16. Found: C, 79.31; H, 7.54; N, 6.27.

N-12-(3-Methyl)butanel-4-methoxyaniline (8). To a THF solution at 0°C containing p-anisidine (1 equiv), 3methyl-2-butanone (1.0 equiv), and acetic acid (1.0 equiv) was added solid NaHB(OAc)₃. The mixture was allowed to reach rt and stirred for 15h. The mixture was re-cooled to 0°C; careful workup (H₂O, EtOAc, NaHCO₃, brine, MgSO₄) and concentration in vacuo afforded crude amine. Kügelrohr distillation provided pure product as light vellow oil (71%): bp 100°C (0.3 mm); ¹H NMR (CDCl₃) δ 0.93 (d, 3H, J=6.5 Hz), 1.00 (d, 3H, J = 6.5 Hz), 1.10 (d, 3H, J = 6.0 Hz), 1.86 (m, 1H, J = 6.5 Hz), 3.23 (br s, 1H, ArNH), 3.28 (dg, 1H, J = 7.0, 6.0 Hz), 3.77 (s, 3H), 6.58 (d, 2H, J =7.0 Hz), 6.80 (d, 211, J = 7.0 Hz); ¹³C NMR (CDCl₃) δ 16.5 (CH₃), 17.3 (CH₃), 19.3 (CH₃), 32.1 (CH), 54.5 (CH), 55.8 (CH₃), 114.6 (ArCH), 114.9 (ArCH), 142.1 (ArC), 151.7 (ArC); MS (EI, 70 eV) m/z 193 (M⁺); anal. (C12H19NO): C, 74.57; H, 9.91; N, 7.25. Found: C, 74.46; H, 9.76; N, 7.32.

General procedure for phase transfer *N*-alkylation of carboxamides.^{9,10} Alkylhalide (3 equiv) in benzene was added dropwise to a stirring mixture of carboxamide (1 equiv), finely powdered potassium hydroxide (2 equiv), and *n*-tetrabutylammonium bromide (0.05 equiv) in benzene (0.2 M) at rt. Reaction mixture was heated to 80 °C and maintained at this temperature until disappearance of starting material was observed by TLC (1–3 h). The reaction mixture was then cooled to rt and the salts removed by filtration. Product isolation (H₂O, brine), followed by flash chromatography (10–25% EtOAc/hexane) afforded product, usually as an oil.

General amidation procedures for anilines 7–8 and CF₃containing anilines 14a–d. A 0.18 M benzene or toluene solution containing 1 equiv of aniline (14a, 14b, 7–8) in addition to 3 equiv of dry powdered K₂CO₃ and 1 equiv of pyridine was treated with 1.5 equiv of the corresponding acid chloride and heated between 80 and 110 °C. The reaction was monitored by TLC and the reaction cooled to rt upon completion. The crude mixture was filtered, concentrated in vacuo and taken up in Et₂O. Product isolation (H₂O, brine) and chromatographic purification afforded the protected carboxamide (15a–e and 9f and 9g).

Benzamides **15f** and **15g** were prepared from anilines **14c** and **14d** and the corresponding acid chloride (1.5 equiv) using 1,2-dichlorobenzene as solvent, 3 equiv of dry powdered K_2CO_3 and heated to 120–150 °C for 16 h or until disappearance of starting aniline. Reaction mixture was cooled to rt and eluted over silica with hexane to remove 1,2-dichlorobenzene. Gradient chromatography with 10, 20, and 30% EtOAc/hexane afforded the pure carboxamides **15f** and **15g** upon concentration.

N-(4-Methoxyphenyl)-*N*-(ethyl)-4-methoxybenzamide (9a). Yellow oil (72%): ¹H NMR (CDCl₃) δ 1.12 (t, 3H, J = 7.2 Hz, CH₃), 3.62 (s, 3H, OCH₃), 3.65 (s, 3H, OCH₃), 3.84 (q, 2H, J = 7.0 Hz, CH₂), 6.57 (d, 2H, J = 8.9 Hz, ArH ortho OCH₃, meta RNCO). 6.68 (d, 2H, J = 8.9 Hz, ArH ortho OCH₃, meta CONR), 6.87 (d, 2H, J = 8.9 Hz, ArH ortho RNCO), 7.20 (d, 2H, J = 8.7 Hz, ArH ortho CONR); ¹³C NMR (CDCl₃) δ 12.7 (CH₃), 45.5 (CH₂), 55.0 (OCH₃), 55.2 (OCH₃), 112.8 (ArCH), 114.2 (ArCH), 128.4 (ArC), 128.9 (ArCH), 130.7 (ArCH), 136.3 (ArC), 157.8 (ArC), 160.2 (ArC), 169.6 (C=O); MS (EI, 70 eV) m/z 285 (M⁺); HRMS calcd for C₁₇H₁₉NO₃, 285.1365, found 285.1364.

N-(4-Methoxyphenyl)-*N*-(2-propyl)-4-methoxybenzamide (9b). Amber oil (43%): ¹H NMR (CDCl₃) δ 1.16 (d, 6H, *J*=6.5 Hz), 3.72 (s, 3H), 3.75 (s, 3H), 5.09 (br s, 1H), 6.63 (d, 2H, *J*=8.5 Hz), 6.74 (d, 2H, *J*=9.0 Hz), 6.93 (d, 2H, *J*=8.5 Hz), 7.21 (d, 2H, *J*=8.5 Hz); ¹³C NMR (CDCl₃) δ 21.2 (CH₃), 55.3 (CH₃), 55.5 (CH₃), 113.0 (ArCH), 113.9 (ArCH), 129.1 (ArC), 130.4 (ArCH), 131.8 (ArCH), 132.5 (ArC), 158.6 (ArC), 160.1 (ArC), 170.5 (C=O); MS (EI, 70 eV) *m*/*z* 299 (M⁺); HRMS calcd for C₁₈H₂₁NO₃ 299.1521, found 299.1521.

N-(4-Methoxyphenyl)-N-(butyl)-4-methoxybenzamide (9c). Yellow oil (81%): ¹H NMR (CDCl₃) δ 0.84 (t, 3H, $J = 7.3 \text{ Hz}, \text{ CH}_3$, 1.29 (sext, 2H, $J = 7.0 \text{ Hz}, \text{ CH}_3\text{CH}_2$). 1.53 (quint, 2H, J = 7.8 Hz, CH₂CH₂CH₂), 3.62 (s, 3H. OCH₃ ortho RNCO), 3.66 (s, 3H, OCH₃ ortho CONR), 3.78 (t, 2H, J = 7.7 Hz, CH₂CH₂N), 6.58 (d, 2H, J =8.8 Hz, ArH ortho OCH₃, meta RNCO), 6.68 (d, 2H, J = 8.9 Hz, ArH ortho OCH₃, and meta CONR), 6.88 (d. 2H, J=8.9 Hz, ArH ortho RNCO), 7.19 (d, 2H, J= 8.9 Hz, ArH ortho CONR); ¹³C NMR (CDCl₃) δ 13.7 (CH₃), 20.0 (CH₂), 29.5 (CH₂), 50.3 (CH₂), 54.9 (OCH₃), 55.1 (OCH₃), 112.7 (ArCH), 114.1 (ArCH), 128.4 (ArC), 128.6 (ArCH), 130.5 (ArCH), 136.5 (ArC), 157.6 (ArC), 160.0 (ArC), 169.6 (ArC); MS (EI, 70 eV) m/z 313 (M⁺); HRMS calcd for C₁₉H₂₃NO₃, 313.7794, found 313.7879.

N-(4-Methoxyphenyl)-*N*-(benzyl)-4-methoxybenzamide (9e). White flakes (82%): mp 104-106 °C; ¹H NMR (CDCl₃) δ 3.72 (s, 3H, OCH₃ para RNCO), 3.74 (s, 3H, OCH₃ para CONR), 5.09 (s, 2H, CH₂Ph), 6.68 (d, 2H, *J*=9.2 Hz, ArH ortho OCH₃, and meta RNCO), 7.31 (m, 9H, Ph and ArH); ¹³C NMR (CDCl₃) δ 54.3 (PhCH₂), 55.3 (OCH₃), 55.5 (OCH₃), 113.1 (ArCH), 114.3 (ArCH), 127.4 (ArCH), 128.6 (ArCH), 128.3 (ArC), 128.7 (ArCH), 128.9 (ArCH), 131.1 (ArCH), 137.0 (ArC), 138.0 (ArC), 158.0 (ArC), 160.6 (ArC), 170.2 (ArC); MS (EI, 70 eV) *m*/*z* 347 (M⁺); HRMS calcd for C₂₂H₂₁NO₃, 347.1521, found 347.1522.

N-(4-Methoxyphenyl)-*N*-(1-phenethyl)-4-methoxybenzamide (9f). Prepared according to general procedure above using aniline 7 and *p*-anisoyl chloride. Purification by flash chromatography (2% (CH₃)₂CO/CH₂Cl₂) afforded product as clear oil (70%): ¹H NMR (CDCl₃) δ 1.48 (d, 3H, *J* = 6.8 Hz), 3.67 (s, 3H), 3.69 (s. 3H), 6.38 (br q, 1H, *J* = 6.8 Hz), 6.56 (br d overlapping br s, 4H), 6.62 (d, 2H, *J* = 8.8 Hz), 7.23 (d, 2H, *J* = 8.5 Hz), 7.29 (m, 5H); ¹³C NMR (CDCl₃) δ 16.9 (CH₃), 53.3 (CH), 55.2 (CH₃), 55.3 (CH₃), 112.9 (ArCH), 113.6 (ArCH), 127.5 (ArCH), 128.2 (ArCH), 128.3 (ArCH), 129.2 (ArC), 130.6 (ArCH), 131.5 (ArCH), 132.7 (ArC), 141.7 (ArC), 158.3 (ArC), 160.2 (ArC), 170.6 (C=O); MS (EI, 70 eV) *m*/*z* 361 (M⁺).

N-(4-Methoxyphenyl)-*N*-(1-methyl-2-methyl-1-propyl)-4methoxybenzamide (9g). Prepared according to general procedure above using aniline **8** and *p*-anisoyl chloride. Purification by flash chromatography (25% EtOAc/ hexane) afforded product as light yellow oil (90%): ¹H NMR (CDCl₃) δ 0.93 (d, 6H, *J*=6.7 Hz), 1.90 (sept, 1H, *J*=7.0 Hz), 3.69 (s, 3H), 3.71 (s, 3H), 3.73 (d, 2H, *J*=7.0 Hz), 6.62 (d, 2H, *J*=8.5 Hz), 6.71 (d, 2H, *J*=8.9 Hz); ¹³C NMR (CDCl₃) δ 20.3 (CH₃), 27.0 (CH), 55.2 (CH₃), 55.4 (CH₃), 57.2 (CH₂), 113.0 (ArCH), 114.3 (ArCH), 128.7 (ArCH), 129.0 (ArC), 130.7 (ArCH), 137.1 (ArC), 157.8 (ArC), 160.3 (ArC), 170.3 (C=O); MS (EI, 70 eV) *m*/*z* 313 (M⁺).

N-(4-Hydroxyphenyl)-*N*-(ethyl)-4-hydroxybenzamide (10a). Deprotected according to BBr₃ procedure to afford offwhite powder, recrystallized from MeOH (70%): mp 208–209 °C; ¹H NMR ((CD₃)₂SO) δ 1.03 (t, 3H, *J*= 6.9 Hz, CH₃), 3.72 (q, 2H, *J*=7.1 Hz, CH₂), 6.52 (d, 2H, *J*=8.6 Hz, ArH meta RNCO), 6.62 (d, 2H, *J*=8.5 Hz, ArH meta CONR), 6.86 (d, 2H, *J*=8.6 Hz, ortho RNCO), 7.06 (d, 2H, *J*=8.5 Hz, ortho CONR), 9.48 (s, 1H, OH), 9.71 (s, 1H, OH); ¹³C NMR ((CD₃)₂SO) δ 12.6 (CH₃), 39.1 (CH₂), 114.3 (ArCH), 115.7 (ArCH), 127.1 (ArC), 129.1 (ArC), 130.5 (ArCH), 134.7 (ArC), 155.7 (ArC), 158.3 (ArC), 168.9 (ArC); MS (EI, 70 eV) *m*/*z* 257 (M⁻¹); HRMS calcd for C₁₅H₁₅NO₃, 257.1052, found 257.1051.

N-(4-Hydroxyphenyl)-*N*-(2-propyl)-4-hydroxybenzamide (l0b). Deprotected according to BBr₃ procedure to afford powder, recrystallized from MeOH (77%): mp 210–212 °C; ¹H NMR (MeOD) δ 1.15 (d, 6H, *J*= 6.8 Hz), 4.94 (br s, 1H), 6.56 (br s, 2H), 6.67 (d, 2H, *J*=8.4 Hz), 6.88; MS (EI, 70 eV) 271 (M⁺); anal. (C₁₆H₁₇NO₃·0.1 H₂O): C, 70.36; H, 6.36; N, 5.13. Found: C, 70.35; H, 6.37; N, 4.90.

N-(4-Hydroxyphenyl)-*N*-(butyl)-4-hydroxybenzamide (10c). Deprotected according to BBr₃ procedure and purified by flash chromatography (25% EtOAc/hexane) to afford off-white crystals (95%): mp 174–175 °C; ¹H NMR ((CD₃)₂SO) δ 0.84 (s, 3H), 1.25 (sext, 2H, *J*=7.6 Hz), 1.43 (quint, 2H, *J*=7.6 Hz), 3.69 (t, 2H, *J*=7.5 Hz), 6.51 (d, 2H, *J*=8.6 Hz), 6.61 (d, 2H, *J*=8.7 Hz), 6.85 (d, 2H, *J*=8.7 Hz), 7.03 (d, 2H, *J*=8.5 Hz); ¹³C NMR ((CD₃)₂SO) δ 14.1 (CH₃), 19.9 (CH₂), 40.2 (CH₂), 114.5 (ArCH), 115.9 (ArCH), 127.4 (ArC), 129.2 (ArCH), 130.6 (ArCH), 135.1 (ArC), 155.8 (ArC), 158.4 (ArC); MS (EI, 70 eV) *m*/*z* 285 (M⁺); HRMS (EI) calcd for C,₇H₁₉NO₃, 285.1365, found 285.1368; anal. (C₁₇H₁₉NO₃·0.5 H₂O): C, 69.37; H, 6.85; N, 4.76. Found: C, 69.30; H, 6.65; N, 4.57.

N-(4-Hydroxyphenyl)-*N*-(2-methyl-1-propyl)-4-hydroxybenzamide (10d). Deprotected according to BBr₃ procedure and purified by flash chromatography (25% EtOAc/hexane) to afford white foam (quant): mp 166 °C dec.; ¹H NMR ((CD₃)₂SO) δ 0.93 (d, 6H, *J*=6.5 Hz), 1.82 (m, 1H, J = 6.5 Hz), 3.66 (d, 2H, J = 7.5 Hz), 6.59 (d, 2H, J = 8.5 Hz), 6.68 (d, 2H, J = 9 Hz), 6.94 (d, 2H, J = 8.5 Hz), 7.10 (d, 2H, J = 8.5 Hz); MS (EI, 70 eV) m/z 285 (M⁺); anal. (C₁₇H₁₉NO₃·l.0 H₂O): C, 67.31; H, 6.98; N, 4.62. Found: C, 67.45; H, 6.74; N, 4.41.

N-(4-Hydroxyphenyl)-N-(benzyl)-4-hydroxybenzamide (10e). Deprotected according to EtSH procedure and purified by flash chromatography (25% EtOAc/hexane), which upon solvent concentration and recrystallization from CHCl₃ afforded title compound as off-white flakes (60%): mp 103 105°C; ¹H NMR ((CD₃)₂SO) δ 4.95 (s, 2H, CH_2), 6.53 (d, 2H, J=6.3 Hz, ArH meta RNCO), 6.56 (d, 211, J = 6.2 Hz, ArH ortho RNCO), 6.77 (d, 2H, J = 8.7 Hz, ArH meta CONR), 7.13 (d, 2H, J=8.7 Hz, AsH ortho CONR), 7.25 (m. 5H, Ph), 9.67 (broad s, 2H, OH); ¹³C NMR ((CD₃)₂SO) δ 53.1 (CH₂), 114.2 (ArCH), 115.5 (ArCH), 122.3 (ArC), 126.6 (ArC), 126.9 (ArCH), 127.9 (ArCH), 128.3 (ArCH), 128.7 (ArCH), 130.6 (ArCH), 134.9 (ArC), 137.9 (ArC), 155.5 (ArC), 158.5 (ArC), 169.4 (ArC); MS (EI, 70 eV) m/z 319 (M⁺); HRMS (EI) calcd for $C_{20}H_{17}N0_3$. 319.1208, found 319.1206; anal. (C₂₀H₁₇NO₃·2.5 H₂O): C, 65.92; H, 6.09; N, 3.84. Found: C, 65.58; H, 5.81; N, 3.53.

N-4-Hydroxyphcnyl-*N*-(1-phenethyl)-4-hydroxybenzamide (10f). Deprotected according to BBr₃ procedure recrystallized from McOH/CH₂Cl₂ to afford white powder (95%): mp 204-206 °C; ¹H NMR (MeOD) δ 1.50 (d, 3H. J = 6.8 Hz), 4.95 (s, under MeOH), 6.24 (br s, 2H), 6.49 (br s, 2H), 6.54 (br s, 2H), 6.54 (d, 2H, J = 8 Hz), 7.10 (d, 2H, J = 8 Hz), 7.28 (m, 5H); ¹³C NMR (MeOD) δ 17.4 (CH₃), 49.0 (CH), 115.3 (ArCH), 115.9 (ArCH), 124.9 (ArCH), 128.6 (ArCH), 128.9 (ArC), 129.1 (ArCH), 129.3 (ArCH), 131.4 (ArCH), 132.4 (ArC), 132.7 (ArCH), 142.5 (ArC), 157.9 (ArC), 159.8 (C=O): MS (EI, 70 eV) m/z 333 (M⁻¹); HRMS (EI) calcd for C₂₁H₁₉NO₃, 333.1365, found 333.1360; anal. (C₂₁H₁₉NO₃): C, 75.66; H, 5.74; N, 4.20. Found: C, 75.51; H, 5.80; N, 4.21.

General thionation procedure¹²

The carboxamide (1.01 mmol) and Lawesson's reagent (0.51 mmol) in 3 mL of HMPA were heated to $80-100 \text{ }^{\circ}\text{C}$ until the carboxamide had been consumed. After disappearance of carboxamide, the reaction mixture was allowed to cool to rt and then poured onto 5 mL of water. Product was extracted repeatedly with Et₂O, dried over Na₂SO₄, rotary evaporated in vacuo and purified via flash chromatography. The protected thioamides were then directly used in the deprotection step with minimal purification.

N-(4-Hydroxyphenyl)-*N*-(ethyl)-4-hydroxythiobenzamide (11a). Deprotected according to BBr₃ procedure and purified by flash chromatography (25% EtOAc/hexane) to afford yellow foam (63%): ¹H NMR ((CD₃)₂SO) δ 1.16 (1, 3H, *J*=6.8 Hz, CH₃), 4.34 (q, 2H, *J*=6.8 Hz, CH₂), 6.44 (d, 2h, *J*=8.4 Hz, ArH *meta* RNCS), 6.59 (d, 2H, *J*=8.9 Hz, ArH *meta* CSNR), 6.87 (d, 2H, *J*=8.7 Hz, ArH *ortho* RNCS), 6.99 (d, 2H, *J*=8.6 Hz, ArH *ortho* CSNR); ¹³C NMR ((CD₃)₂SO) δ 11.1 (CH₃), 52.1 (CH₂), 114.1 (ArH), 115.6 (ArH), 115.7 (ArH), 122.6 (ArH), 128.5 (ArH), 129.8 (ArH), 135.3 (ArC), 136.4 (ArC), 156.3 (ArC), 157.6 (ArC), 201.0 (CS); MS (EI, 70 eV) m/z 273 (M⁺); HRMS (EI) calcd for C₁₅H₁₅NSO₂, 273.0824, found 273.0824; anal. (C₁₅H₅NSO₂·0.3 H₂O): C, 64.63; H, 5.64; N, 5.02. Found: C, 64.51; H, 5.93; N, 4.64.

N-(4-Hydroxyphenyl)-*N*-(*i*-propyl)-4-hydroxythiobenzamide (11b). Deprotected according to BBr₃ procedure and purified by flash chromatography (25% EtOAc/ hexanc) to afford yellow foam (98%): ¹H NMR (MeOD, *i*-propyl rotamers 1:4:1 ratio) δ 1.15 (d, 1H, J=6.5 Hz), 1.20 (d, 4H, J=6.5 Hz), 1.27 (d, 1H, J=6.5 Hz), 6.00 (sept, 1H, J=6.5 Hz), 6.46 (d, 2H, J=8.5 Hz), 6.61 (d, 2H, J=9 Hz), 6.81 (d, 2H, J=8.5 Hz), 6.96 (d, 2H, J=8.5 Hz) additional minor ArCH resonances not listed; ¹³C NMR (MeOD) δ 20.6 (CH₃), 21.9 (CH₃ minor), 54.8 (CH), 114.8 (ArCH), 115.9 (ArCH), 130.1 (ArCH), 131.7 (ArCH), 133.4 (ArC), 137.8 (ArC), 157.9 (ArC), 158.0 (ArC), 204.4 (C=S); MS (EI, 70 eV) *m*/*z* 287 (M⁺); HRMS, C₁₆H₁₇NSO₂, 287.0980; found, 287.0971.

N-(4-Hydroxyphenyl)-*N*-(*n*-butyl)-4-hydroxythiobenzamide (11c). Deprotected according to BBr₃ procedure and purified by radial chromatography (5% MeOH/ CH_2Cl_2) to afford yellow oil (10%): ¹H NMR ((CD_3)₂ SO) δ 0.84 (s, 3H, J=7.5 Hz, CH₃), 1.27 (sext, 2H, $J = 6.8 \text{ Hz}, \text{ CH}_3 \text{CH}_2$), 1.61 (quint, 2H, $J = 8.0 \text{ Hz}, \text{ CH}_3$) CH_2CH_2), 4.30 (t, 2H, J=7.7 Hz, CH_2NCS), 6.43 (d, 2H, J=8.5 Hz, ArH meta RNCS), 6.58 (d, 2H, J=8.2 Hz, ArH meta CSNR), 6.87 (d, 2H, J = 8.2 Hz, ArH ortho RNCS), 6.97 (d, 2H, J=8.5 Hz, ArH ortho CSNR); ¹³C NMR ((CD₃)₂SO) δ 14.2 (CH₃), 19.7 (CH₂), 27.9 (CH₂), 56.8 (CH₂), 114.1 (ArCH), 115.9 (ArCH), 128.5 (ArCH), 129.9 (ArCH), 135.2 (ArC), 136.8 (ArC), 156.2 (ArC), 157.7 (ArC), 201.0 (C=S); MS (EI, 70 eV), m/z 301 (M⁺); HRMS calcd for C₁₇H₁₉NO₂S, 301.1136; found 301.1133.

N-(4-Hydroxyphenyl)-*N*-(*i*-butyl)-4-hydroxythiobenzamide (11d). Deprotected according to BBr₃ procedure and purified by flash chromatography (35% EtOAc/hexane) to afford yellow foam (86%): ¹H NMR (CDCl₃) δ 0.98 (d, 6H, *J*=6.8 Hz), 2.16 (m, 1H, *J*=6.8 Hz), 4.34 (d, 2H, *J*=7.6 Hz), 6.41 (d, 2H, *J*=8.8 Hz), 6.60 (d, 2H, *J*=8.8 Hz), 6.79 (d, 2H, *J*=8.8 Hz), 6.96 (d, 2H, *J*=8.8 Hz); ¹³C NMR (CDCl₃) δ 20.3, 26.9, 27.2, 50.8, 63.9, 114.5, 116.1, 127.9, 129.5, 136.4, 137.7, 155.4, 156.5, 203.0; MS (EI, 70 eV) *m*/*z* 301 (M⁺); HRMS calcd for C₁₇H₁₉NO₂S, 301.1136; found, 301.1144.

N-(4-Hydroxyphenyl)-*N*-(benzyl)-4-hydroxythiobenzamide (11e). Deprotected according to EtSH procedure and purified by flash chromatography (35% EtOAc/hexane) to afford yellow foam (16%): ¹H NMR ((CDCl₃)) δ 5.76 (s, 2H, CH₂), 6.59 (d, 2H, *J* = 8.4 Hz, ArH *meta* RNCS), 6.61 (d, 211, *J* = 8.3 Hz, ArH *meta* CSNR), 6.77 (d, 2H, *J* = 8.8 Hz, ArH *ortho* RNCS), 7.27 (m, 5H, Ph), 7.38 (d, 2H, *J* = 6.4 Hz, ArH *ortho* CSNR); MS (EI, 70 eV) *m/z* 335 (M⁺); HRMS calcd for C₂₀H₁₇NO₂S, 335.0980; found, 335.0984.

N-(4-hydroxyphenyl)-N-(1-phenethyl)-4-hydroxythiobenzamide (11f). Deprotected according to BBr₃ procedure and recrystallized from CHCl₃ to afford light yellow powder (quant): mp 174–175 °C; ¹H NMR (CDCl₃) δ 1.55 (d, 3H, *J*=7.2 Hz); 5.89 (br s, 1H); 6.27 (br s, 1H) 6.44 (d, 2H, *J*=8.8 Hz), 6.5 (br s, 1H), 6.71 (br s, 1H), 6.96 (d, 2H, *J*=8.8 Hz), 7.32 (m, 5H), 7.52 (q, 1H, *J*=6.8 Hz); MS (EI, 70 eV) *m/z* 349 (M⁺); anal. (C₂₁H₁₉NO₂S·0.4 H₂O): C, 70.72; H, 5.60; N, 3.93. Found: C, 70.88; H, 5.42; N, 3.98.

N-(4-Hydroxyphenyl)-*N*-[2-(3-methyl)-butyl]-4-hydroxythiobenzamide (11g). Deprotected according to BBr₃ procedure and purified by flash chromatography (25% EtOAc/hexane) to afford foam (37%, two steps from carboxamide): ¹H NMR (CDCl₃) δ (major methyl rotamer) 0.95 (d, 3H, J=6.6 Hz), 1.14 (d, 3H, J=6.9 Hz), 1.21 (d, 3H, J=6.3 Hz), 1.82 (br m, 1H), 5.82 (quin, 1H, J=7.4 Hz), 6.30 (d, 2H, J=8.5 Hz), 6.60 (br m, 4H), 6.83 (d, 2H, J=8.5 Hz); ¹³C NMR (CDCl₃) δ 17.1 (CH₃), 19.8 (CH₃), 20.9 (CH), 32.5 (CH), 114.7 (ArCH), 115.6 (ArCH), 128.8 (ArCH), 129.5 (ArCH), 133.2 (ArC), 137.2 (ArC), 155.2 (ArC), 155.3 (ArC), 204.4 (C=S); MS (EI, 70 eV) m/z 315 (M⁺); anal. (C₁₈H₂₁NO₂S·0.8 H₂O): C, 65.54; H, 6.91; N, 4.35. Found: C, 65.15; H, 6.67; N, 3.91.

Triphenylphosphine-4-methoxyphenylimine (12a).³⁶ p-Anisidine was diazotized in 50% H₂SO₄ and then added to a NaN₃-sodium acetate buffered solution.¹⁵ The azide was extracted with diethyl ether and the phenolic side-product removed by Na₂CO₃ wash. Concentration in vacuo afforded the crude azide which was directly used in the next step. Ylide formation: to an etheral solution of azide was added an equimolar solution of PPh₃ at rt. After the solution was heated under reflux for 2h and N₂ evolution ceased, the solvent was concentrated and resulting Ph₃P=O oxide removed. The crude imine was passed over SiO₂ plug (ether) to remove additional oxide. Final concentration and tituration with hexanes afforded pure imine as orange crystals $(53\% \text{ from } p\text{-anisidine}): \text{mp } 116-117.5 ^{\circ}\text{C}; \text{mp}^{33}$ 119-120°C.

Triphenylphosphine-phenylimine (12b).^{14,37} A cold aqueous solution of NaNO₂ (1.2 equiv) was added dropwise to a solution of aniline in 10% HCl at 0–5 °C with vigorous stirring. The mixture was kept below 5 °C for 30 min, and then a solution of NaN₃ in water was added dropwise while the temperature was kept below 5 °C. After 1 h the reaction was warmed to rt and extracted with diethyl ether. The extracts were dried over Na₂SO₄ and concentrated to afford the crude azide¹⁴ as an oil, which was directly used in the next step without additional purification. *Ylide formation:* same as described for **12a** to afford light yellow crystals upon solvent concentration (64% from aniline): mp 128–130 °C (mp³⁴ 131–132 °C).

N-(1,1,1-Trifluoro-2-propylidene)-4-methoxyaniline (13a).⁶ Ylide 12a and trifluoroacetone were heated to reflux in C_6H_6 for 12h. The reaction was cooled to rt and concentrated in vacuo. The residue was triturated with Et₂O to remove Ph₃PO and the resulting filtrate concentrated followed by Kügelrohr distillation under

reduced pressure to afford product as light yellow oil (92%): bp 66 °C (0.1 mm); ¹H NMR (CDCl₃) δ 2.06 (s, 3H), 3.81 (s, 3H), 6.79, 6.92 (AA'BB', 4H, *J*=8.8, 4.0 Hz); ¹³C NMR δ 14.6 (CH₃), 55.6 (CH₃), 114.5 (ArCH), 120.5 (q, CF₃, *J*=273 Hz), 121.1 (ArCH), 121.5 (ArCH), 140.5 (ArC), 157.1 (q, CCF₃, *J*=33 Hz), 157.6 (ArC); MS (EI, 70 eV) *m/z* 217 (M⁺); anal. (C₁₀H₁₀NF₃O): C, 55.30; H, 4.64; N, 6.45. Found: C, 55.59; H, 4.99; N, 6.64.

N-(1,1,1-Trifluoro-2-propyl)-4-methoxyaniline (14a).⁶ To a Et₂O solution of LiAlH₄ (0.5 equiv) at 0 °C was added a Et₂O solution of propylidene 13a dropwise over 30 min. Upon complete addition the mixture was refluxed for 2h. The mixture was then cooled to 0°C and quenched using a 1:1:3 work up (H₂O:3M NaOH:H₂O). The resulting mixture was decanted into a separatory funnel, the aqueous layer separated and ether layer washed with brine, dried over MgSO₄ and concentrated. Purification by Kügelrohr distillation provided pure amine as light red oil (64%): bp 80°C (0.2 mm); ¹H NMR (CDCl₃) δ 1.34 (d, 3H, J=7.0 Hz), 3.32 (br s, 1H), 3.72 (s, 3H), 3.87 (m, 1H), 6.61 (AA'BB', 2H, J = 8.8, 4.0 Hz), 6.77 (AA'BB', 2H, J = 8.8, 3.5 Hz); ¹³C NMR (CDCl₃) δ 15.2 (CH₃), 52.7 (q, CHCF₃, J = 30 Hz), 55.8 (CH₃), 115.1 (ArCH), 115.4 (ArCH), 126.5 (q, CF₃, J = 280 Hz), 140.2 (ArC), 153.2 (ArC); MS (EI, 70 eV) m/z 219 (M⁺).

N-(1,1,1-Trifluoro-2-phencthyl)-4-methoxyaniline (14b). Prepared according to procedures outlined above from ylide 12a and trifluoroacetophenone. The imine was isolated then used directly in reduction step to afford the title compound as a light yellow oil upon distillation (69%): bp 110 °C (0.1 mm); ¹H NMR (CDCl₃) δ 3.72 (s, 3H), 4.10 (br s, 1H), 4.81 (q, 1H, *J*=7.0 Hz), 6.61 (AA'BB', 2H, *J*=8.8, 4.0 Hz), 6.75 (AA'BB', 2H, *J*=8.8, 3.5 Hz), 7.40 (m, 5H); ¹³C NMR (CDCl₃) δ 55.8 (CH₃), 62.0 (CHCF₃, q, *J*=30 Hz), 115.0 (ArCH), 115.9 (ArCH), 125.5 (q, CF₃, *J*=280 Hz), 128.2 (ArCH), 129.1 (ArCH), 129.3 (ArCH), 134.5 (ArC), 139.7 (ArC), 153.5 (ArC); MS (EI, 70 eV) *m*/*z* 281 (M⁺); anal. (C₁₅H₁₄NF₃O): C, 64.05; H, 5.02; N, 4.98. Found: C, 63.90; H, 4.84; N, 5.05.

N-(1,1,1-Trifluoro-2-phenethyl)-aniline (14c). Prepared according to procedures outlined above using ylide 12b and trifluoroacetophenone. The imine was isolated then used directly in reduction to afford the title compound as a clear oil upon distillation (59%): bp 100 °C (0.5 mm); ¹H NMR (CDCl₃) δ 4.36 (d, 1H, J=6.5 Hz, NH), 4.96 (quint, 1H, J=7.0 Hz), 6.68 (d, 2H, J= 8.0 Hz), 6.82 (t, 1H, J=7.0 Hz), 7.20 (d, 2H, J=7.0 Hz), 7.42 (m, 3H), 7.50 (d, 2H, J=6.5 Hz), ¹³C NMR (CDCl₃) δ 60.7 (q, CHCF₃, J=29 Hz), 114.1 (ArCH), 119.5 (ArCH), 122.5 (q, CF₃, J=278 Hz), 128.1 (ArCH), 129.2 (ArCH), 129.4 (ArCH), 129.6 (ArCH), 134.3 (ArC), 145.8 (ArC); MS (EI, 70 eV) m/z 251 (M⁺); anal. (C₁₄H₁₂F₃N): C, 66.93; H, 4.81; N, 5.57. Found: C, 66.79; H, 4.80; N, 5.54.

N-[1,1,1-Trifluoro-2-(3-methoxy)-phenethyl]-aniline (14d). 2,2,2-Trifluoro-1-(3-methoxyphenyl)-ethanone precursor).¹³

Mg turnings (1.25 g, 50 mmol), 3-methoxyphenyl bromide (9.35 g, 50 mmol), and anhydrous THF (50 mL) were gingerly heated until a vigorous reaction took place. When all the Mg turnings were dissolved the reaction mixture was cooled to 0 °C. A solution of *N*-trifluoroacetylpiperidine³⁵ (7.52 g, 45 mmol) in anhydrous THF (10 mL) was added to the Grignard reagent dropwise over 0.5 h with stirring at 0 °C. Upon complete addition, ice-bath was removed and the mixture stirred for 2 h. The reaction was quenched by the addition of satd aq NH₄Cl (5 mL), and the precipitates romoved by filtration. The filtrate was dried over MgSO₄, evaporated in vacuo, and the crude ketone distilled to give 6.6 g (72%) of a colorless liquid: bp 50 °C (0.5 mm) (bp⁹ 84 85 °C (12 mm).

The above prepared 2,2,2-trifluoro-1-(3-methoxyphenyl)ethanone was then reacted with vlide 12a as described previously. Imine isolation and LAH reduction afforded the title compound as a light green oil upon distillation (80%): bp 125 °C (0.8 mm); ¹H NMR (CDCl₃) δ 3.79 (s, 3H), 4.31 (br s, 1H), 4.87 (q, 1H, J = 7.2 Hz), 6.63 (d, 2H, J = 7.6 Hz), 6.76 (td, 1H, J = 7.6, 0.8 Hz), 6.89 (dd, 1H, J = 8.5, 2.4 Hz), 6.99 (br s, 1H), 7.03 (d, 1H, J = 7.6 Hz), 7.15 (m, 2H), 7.28 (t, 1H, J = 8.0 Hz); ¹³C NMR (CDCl₃) δ 55.1 (CH₃), 60.3 (q. CHCF₃, J = 30 Hz), 113.8 (ArCH), 114.1 (ArCH), 119.1 (ArCH), 120.1 (ArCH), 124.9 (q, CF_3 , J = 281 Hz), 128.7 (ArCH), 129.2 (ArCH), 129.8 (ArCH), 135.5 (ArC), 145.4 (ArC), 159.8 (ArC); MS (E1, 70 eV) m/z 281 (M⁺); anal. (C₁₅H₁₄F₃NO): C, 64.05; H, 5.02; N, 4.98. Found: C, 64.15; H, 4.99; N, 4.84.

N-(4-Methoxyphenyl)-*N*-(1,1,1-trifluoro-2-propyl)-4-methoxybenzamide (15a).⁶ Prepared according to general procedure for CF₃-containing anilines and purified by flash chromatography (25% EtOAc/hexane) to afford product as white powder (68%): mp 85 ·86 °C: ¹H NMR (CDCl₃) δ 1.21 (d, 3H, J = 7.0 Hz), 3.69 (s, 3H), 3.73 (s, 3H), 5.79 (sept, 1H, J = 6.5 Hz), 6.62 (d, 2H, J = 8.5 Hz), 6.75 (br s, 2H), 7.12 (br s, 2H), 7.22 (d, 2H, J = 8.5 Hz); ¹³C NMR (CDCl₃) δ 12.5 (CH₃), 55.2 (CH₃), 55.4 (CH₃), 113.1 (ArCH), 114.1 (ArCH), 125.2 (q, CF₃, J = 282 Hz), 128.0 (ArC), 130.6 (ArCH). 131.7 (ArC), 159.0 (ArC), 160.6 (ArC), 171.6 (C=O): MS (E1, 70 eV) m/z 353 (M⁺).

N-(4-Methoxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-4methoxybenzamide (I5b). Prepared according to general procedure for CF₃-containing anilines and purified by flash chromatography (25% EtOAc/hexane) to afford product as transparent oil (77%): ¹H NMR (CDCl₃) δ 3.68 (s, 3H), 3.69 (s, 3H), 6.05 (br s, 1H), 6.42 (br s, 1H), 6.61 (d overlapping br s, 3H, J=8.5 Hz) 7.04 (br q, 1H, J_{1HF} =6.5 Hz), 7.25 (m, 8H); ¹³C NMR (CDCl₃) δ 55.1 (CH₃), 55.2 (CH₃), 58.8 (ArCHCF₃), 113.0 (ArCH), 113.6 (ArCH), 125.1 (q, CF₃, J=281 Hz), 127.5 (ArC), 128.5 (ArCH), 129.0 (ArCH), 130.0 (ArCH), 130.7 (ArCH), 131.9 (ArC), 132.0 (ArCH), 132.9 (ArCH), 158.8 (ArC), 160.5 (ArC). 171.4 (C=O); MS (E1, 70 eV) m/z 415 (M⁺); HRMS caled for C₂₃H₂₀F₃NO₃, 415.1395. Found, 415.1387. *N*-(4-Methoxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-benzamide (15c). Prepared according to general procedure for CF₃-containing anilines and purified by flash chromatography (10% EtOAc/hexane) to afford product as light yellow oil (73%): bp 125 °C (0.3 mm); ¹H NMR (CDCl₃) δ 3.61 (s, 3H), 6.01 (br s, 1H), 6.52 (2 overlapping br s, 3H), 7.12 (m, 4H), 7.26 (m, 6H); ¹³C NMR (CDCl₃) δ 55.4 (CH₃), 58.1 (CHCF₃), 113.6 (ArCH), 125.1 (q, CF₃, *J*=282 Hz), 127.9 (ArCH), 128.7 (ArCH), 129.2 (ArCH), 129.6 (ArCH), 130.2 (ArCH), 131.4 (ArC), 131.9 (ArC), 132.4 (ArCH), 135.7 (ArC), 159 (ArC), 172.1 (C=O); MS (EI, 70 eV) *m*/*z* 385 (M⁺); anal. (C₂₂H₁₈F₃NO₂): C, 68.57; H, 4.71; N, 3.68. Found: C, 68.22; H, 5.03; N, 3.68.

N-(4-Methoxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-3methoxybenzamide (15d). Prepared according to general procedure for CF₃-containing anilines and purified by flash chromatography (10% EtOAc/hexane) to afford product as transparent oil (79%): ¹H NMR (CDCl₃) δ 3.66 (2-s, 3H, ~1.2/1 ratio benzoyl rotamers), 3.88 (s, 3H), 6.78 (br d, 1H, *J*=8.5 Hz), 6.8 (br s, 1H), 7.03 (q, 1H, *J*=6.5 Hz, CHCF₃), 7.25 (m, 5H), 7.65 (dd, 1H, *J*=8.5, 2.3 Hz), 7.75 (app dt, 1H, *J*=7.6, 0.8 Hz); MS (EI, 70 eV) *m*/z 415 (M⁺).

N-(4-Methoxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-2methoxybenzamide (15e). Prepared according to general procedure for CF₃-containing anilines and purified by flash chromatography (10% EtOAc/hexane) to afford product as transparent oil (36%, 84% corrected for recovered aniline); ¹H NMR (CDCl₃) δ 3.63 (s, 3H), 3.68 (s, 3H), 6.10 (br s, 1H), 6.43 (br s, 2H), 6.58 (d, 2H, J=8.5 Hz), 6.72 (app t, 2H, J=8.5 Hz), 7.21 (m, 7H); MS (EI, 70 eV) m/z 415 (M⁺).

N-(Phenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-4-methoxybenzamide (15f). Prepared according to the general procedure for CF₃-containing anilines using 1,2dichlorobenzene as solvent and purified by flash chromatography (10% EtOAc/hexane) to afford product as a transparent oil (54%): ¹H NMR (CDCl₃) δ 3.69 (s, 3H), 6.62 (d, 2H, J=8.5 Hz), 7.11 (m, 3H), 7.25 (m, 9H); ¹³C NMR (CDCl₃) δ 55.1 (CH₃), 58.5 (CHCF₃), 113.0 (ArCH), 125.2 (q, CF₃, J=282 Hz), 127.3 (ArC), 127.9 (ArCH), 128.6 (ArCH), 129.0 (ArCH), 129.9 (ArCH), 130.7 (ArCH), 130.9 (ArCH), 131.9 (ArC), 139.5 (ArC), 161.0 (ArC), 171.2 (C=O); MS (EI, 70 eV) m/z 385 (M⁺); anal. (C₂₃H₂₀NO₃F₃·1.1 CHCl₃): C, 52.95; H, 3.89; N, 2.56. Found: C, 52.84; H, 3.90; N, 2.65.

N-(Phenyl)-*N*-[1,1,1-trifluoro-2-(3-methoxy)-phenethyl]-4-methoxybenzamide (15g). Prepared according to general procedure for CF₃-containing anilines using 1,2dichlorobenzene as solvent and purified by flash chromatography (10–25% EtOAc/hexane) to afford product as transparent oil (51%): ¹H NMR (CDCl₃) δ 3.68 (s, 3H), 3.69 (s, 3H), 6.61 (d, 2H, *J*=8.5 Hz), 6.77 (s, 1H), 6.88 (app td, 2H, *J*=8.5, 3.5 Hz), 7.10 (m, 5H), 7.18 (app t, *J*=8.0 Hz), 7.26 (d, 2H, *J*=8.5 Hz); ¹³C NMR (CDCl₃) δ 55.3 (CH₃), 55.4 (CH₃), 58.7 (CHCF₃), 113.1 (ArCH), 115.0 (ArCH), 115.5 (ArCH), 122.5 (ArCH), 125.4 (q, CF₃, J = 281 Hz), 127.5 (ArC), 128.1 (ArCH), 128.7 (ArCH), 129.7 (ArCH), 130.9 (ArCH), 131.1 (ArCH), 133.3 (ArC), 139.7 (ArC), 159.6 (ArC), 160.9 (ArC), 171.4 (C=O); MS (EI, 70 eV) m/z 415 (M⁺).

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-propyl)-4hydroxybenzamide (16a).⁶ Deprotected according to BBr₃ procedure and recrystallized from EtOAc/hexane to afford white powder (62%): mp 203–205 °C (mp⁶ 207–207.5 °C); ¹H NMR (acetone- d_6) δ 1.24 (d. 3H, J = 6.5 Hz), 5.78 (q, 1H, J = 6.5 Hz), 6.62 (d, 2H, J = 8.8 Hz), 6.78 (br s, 2H), 7.05 (br s, 2H), 7.18 (d, 2H, J = 8.8 Hz), 8.61 (s, 1H), 8.70 (s, 1H); ¹³C NMR (acetone- d_6) δ 12.6 (CH₃), 51.3 (q, CH, ³ J_{H-F} = 30 Hz), 114.9 (ArCH), 115.0 (ArCH), 116.2 (ArCH), 126.8 (q, CF₃, ¹ J_{CF} =282 Hz), 128.2 (ArC), 131.4 (ArCH), 131.6 (ArC), 157.7 (ArC), 159.3 (ArC), 171.8 (C=O); MS (EI, 70 eV) m/z 325 (M⁺); anal. (C₁₆H₁₄NO₃F₃): C, 59.08; H 4.34; N, 4.31. Found: C, 59.14; H, 4.37; N, 3.96.

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-4hydroxybenzamide (16b). Deprotected according to BBr₃ procedure and purified by radial chromatography (25% EtOAc/hexane) to afford product as thick oil (53%): ¹H NMR (MeOD) δ 5.91 (br s, 1H) 6.32 (br s, 1H), 6.54 (d, 2H, J=8.8 Hz), 6.62 (br s, 1H), 6.94 (q, 1H, J_{H-F} =9.2 Hz), 7.13 (d, 2H, J=8.8 Hz), 7.23 (m, 6H); ¹³C NMR (MeOD) δ 60.2 (CH), 113.9 (ArCH), 114.5 (ArCH), 125.0 (q, CF₃, J=282 Hz), 125.8 (ArC), 128.0 (ArCH), 128.7 (ArCH), 129.6 (ArCH) 130.0 (ArCH), 130.2 (ArC), 131.3 (ArC), 131.8 (ArC), 156.9 (ArC), 158.8 (ArC), 172.7 (C=O); MS (EI, 70 eV) m/z387 (M⁺); anal. (C₂₁H₁₆F₃NO₃·0.1 H₂O): C, 64.81; H, 4.20; N, 3.60. Found: C, 64.68; H, 4.30; N, 3.45

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)benzamide (16c). Deprotected according to BBr₃ procedure and recrystallized from EtOAc/hexane to afford white powder (83%): mp 158–159 °C, ¹H NMR (CDCl₃) δ 5.91 (br s, 1H), 6.30 (br s, 1H), 6.61 (br s, 1H), 6.74 (s, 1H), 6.95 (q, 1H, *J*=8.8 Hz), 7.24 (m, 9H); MS (EI, 70 eV) *m*/*z* 371 (M⁺); anal. (C₂₁H₁₆F₃NO₂): C, 67.92; H, 4.34; N, 3.77. Found: C, 68.05; H, 4.35; N, 3.63.

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-3hydroxybenzamide (16d). Deprotected according to BBr₃ procedure and recrystallized from EtOAc/hexane to afford off-white powder (77%): mp 163–164 °C; ¹H NMR (CDCl₃) δ 5.92 (br s, 1H), 6.28 (br s, 1H), 6.64 (m, 2H), 6.96 (m, 3H), 7.31 (m, 7H); MS (EI, 70 eV) *m*/*z* 387 (M⁺); anal. (C₂₁H₁₆NO₃F₃·H₂O): C, 62.52; H, 4.48; N, 3.46. Found: C, 62.52; H, 4.17; N, 3.27.

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-2hydroxybenzamide (16e). Deprotected according to BBr₃ procedure and recrystallized from EtOAc/hexane to afford off-white powder (59%): mp 191.5–193 °C; ¹H NMR (MeOD) δ 6.10 (br s, 2H), 6.59 (app q, 2H, J=7.6, 7.2 Hz), 6.96 (m, 3H), 7.32 (m, 6H); MS (EI, 70 eV) m/z 387 (M⁻⁴); anal. (C₂₁H₁₆F₃NO₃·0.7 H₂O): C, 63.06; H, 4.38; N, 3.50. Found: C, 63.04; H, 4.08; N, 3.50. *N*-(Phenyl)-*N*-(1,1,1-trifluoro-2-phenethyl)-4-hydroxybenzamide (16f). Deprotected according to BBr₃ procedure and recrystallized from EtOAc/hexane to afford offwhite powder (quant): mp 114–116 °C; ¹H NMR (CDCl₃) δ 6.48 (d, 2H, *J*=8.4 Hz), 7.0 (m, 4H) 7.15 (d, 2H, *J*=8.4 Hz), 7.26 (d, 6H), 7.72 (s, 2H, exchange with D₂O); ¹³C NMR (MeOD) δ 50.8 (CH), 114.9 (ArCH), 125.0 (q, CF₃, *J*_{CF}=281 Hz), 126.3 (ArC), 128.2 (ArCH), 128.6 (ArCH), 129.3 (ArCH), 130.0 (ArCH), 130.8 (ArCH), 131.0 (ArCH), 131.6 (ArC), 139.2 (ArC), 158.2 (ArC), 172.6 (C=O); MS (EI, 70 eV) *m*/*z* 371 (M⁺); anal. (C₂₁H₁₆F₃NO₂·0.5 H₂O): C, 66.31; H, 4.50; N, 3.68. Found: C, 66.19; H, 4.27; N, 3.57.

N-(Phenyl)-*N*-(1,1,1-trifluoro-2-(3-hydroxyphenethyl))-4hydroxybenzamide (16g). Deprotected according to BBr₃ procedure and recrystallized from CHCl₃ to afford small off-white crystals (95%): mp 163–165 °C; ¹H NMR (CDCl₃) δ 6.43 (d, 2H, *J*=8.5 Hz), 6.67 (d, 1H, *J*= 8.0 Hz), 6.82 (dd, 1H, *J*=8.5, 2 Hz), 6.91 (s, 1H), 7.08 (m, 8H), 7.62 (s, 1H); ¹³C NMR (CDCl₃) 114.9 (ArCH), 116.6 (ArCH), 117.0 (ArCH), 121.2 (ArCH), 124.6 (q, CF₃, *J*=226 Hz), 126.1 (ArC), 128.3 (ArCH), 128.6 (ArCH), 129.8 (ArCH), 130.6 (ArCH), 132.6 (ArC), 138.8 (ArC), 156.3 (ArC), 157.8 (ArC), 173.0 (C=O); MS (EI, 70 eV) *m*/*z* 387 (M⁺); anal. (C₂₁H₁₆F₃NO₃): C, 65.12; H, 4.16; N, 3.62. Found: C, 64.78; H, 4.18; N, 3.38.

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-propyl)-4hydroxythiobenzamide (17a). Deprotected according to BBr₃ procedure and recrystallized from ether to afford yellow powder (93%): ¹H NMR (MeOD) δ 1.33 (d, 3H, *J*=7.5 Hz), 6.47 (d, 2H, *J*=8 Hz), 6.58 (d, 1H, *J*= 6.5 Hz), 6.64 (d, 1H, *J*=6 Hz), 6.79 (d, 1H, *J*=8.5 Hz), 6.96 (d, 1H, *J*=8.0 Hz), 7.01 (d, 3H, *J*=8.5 Hz; ArCH *ortho* C=O and CHCF₃); MS (EI, 70 eV) *m/z* 341 (M⁺); anal. (C₁₆H₁₄F₃NO₂S·0.1 H₂O): C, 56.00; H, 4.17; N, 4.08. Found: C, 55.76; H, 4.13; N, 3.74.

N-(4-Hydroxyphenyl)-N-(1,1,1-trifluoro-2-phenethyl)-4hydroxythiobenzamide (17b). Deprotected according to BBr₃ procedure and purified by chromatography (5% $MeOH/CH_2Cl_2$) to afford title compound as yellow foam (74%): ¹H NMR (CDCl₃) δ 5.47 (br s, 1H, OH), 5.58 (br s, 1H, OH), 5.86 (d, 1H, J=5.5 Hz, ArH ortho to OH on aniline ring), 6.25 (d, 1H, J = 5.5 Hz, ArH ortho to OH on aniline ring), 6.36 (d, 2H, J=8 Hz, ArH ortho to OH on benzamide ring), 6.57 (d, 1H, J = 6 Hz, ArH ortho to N–C=S), 6.93 (d, 2H, J=9 Hz, ArH ortho to \overline{C} =S), 6.98 (d, 1H, J=6Hz, ArH ortho to N–C=S), 8.39 (q, 1H, J_{HF}=9Hz); ¹³C NMR (CDCl₃) δ 63.8 (CHCF₃), 114.6 (ArCH), 115.0 (ArCH), 115.5 (ArCH), $12\overline{5.1}$ (q, J=163 Hz, CF₃), 129.0 (ArCH), 129.3(ArCH), 129.8 (ArCH), 129.9 (ArCH), 131.4 (ArC), 133.4 (ArC), 136.3 (ArC), 155.3 (ArC), 155.7 (ArC), 208.3 (C=S); MS (EI, 70 eV) m/z 403 (M⁺); HRMS calcd for C₂₁H₁₆F₃NO₂S, 403.0853, found 403.0858; anal. (C21H16F3NO2S·0.3 H2O): C, 61.70; H, 4.09; N, 3.43. Found: C, 61.68; H, 3.92; N, 3.25.

N-(4-Methoxyphenyl)-4-methoxyphenylacetamide (18). To a $0 \,^{\circ}$ C CH₂Cl₂ solution of *p*-anisidine (1.3 equiv) and pyridine (1.1 equiv) was added dropwise a CH₂Cl₂

solution (2 5 mL) of 4-methoxyphenylacetyl chloride (1.0 equiv). Upon complete addition (15 min) mixture was allowed to reach rt. Upon completion of the reaction, the mixture was diluted with CH₂Cl₂ followed by product isolation (H₂O, CuSO₄, brine). Recrystallization from EtOAc/hexane afforded the title carboxamide as small white crystals (82%): mp 127-128 °C; ¹H NMR (CDCl₃) & 3.65 (s, 2H), 3.76 (s, 3H), 3.82 (s, 3H), 6.80 (d, 2H, J=9.0 Hz), 6.92 (d, 2H, J=8.5 Hz), 7.07 (s, 1H),7.23 (d, 2H, J = 9.0 Hz), 7.30 (d, 2H, J = 8.5 Hz); ¹³C NMR (CDCl₃) δ 43.7 (CH₂), 55.3 (CH₃), 55.5 (CH₃), 114.1 (ArCH), 114.6 (ArCH), 121.8 (ArCH), 126.5 (ArC), 130.7 (ArCH), 130.8 (ArC), 156.5 (ArC), 159.0 (ArC), 169.7 (C=O); MS (E1, 70 eV) m/z 271 (M⁺); anal. (C₁₆H₁₇NO₃): C, 70.83; H, 6.32; N, 5.16. Found: C, 70.87; H, 6.29; N, 5.29.

N-(4-Methoxyphenyl)-*N*-(ethyl)-4-methoxyphenylacetamide (19). Prepared according to phase transfer catalysis conditions described above for carboxamides. Purification by either Kügelrohr distillation or flash chromatography (25% EtOAc/hexane) afforded product as redorange oil (quant): bp 150°C (0.2 mm); ¹H NMR $(CDCl_3) \delta 1.08$ (t, 3H, J = 7.0 Hz), 3.33 (s, 2H), 3.70 (q, 2H, J = 7.1 Hz), 3.78 (s, 3H), 3.84 (s, 3H), 6.77 (d, 2H, J = 8.5 Hz, 6.79 (d, 2H, J = 8.5 Hz), 6.97 (d overlapping, 2H, J = 8.5 Hz), 6.98 (d overlapping, 2H, J = 8.5 Hz); ¹³C NMR (CDCl₃) δ 13.2 (CH₃), 40.5 (CH₂), 44.4 (CH₂), 55.4 (CH₃O), 55.7 (CH₃O), 113.9 (ArCH), 114.8 (ArCH), 127.9 (ArC), 129.9 (ArCH), 130.2 (ArCH), 135.2 (ArC), 158.4 (ArC), 159.2 (ArC), 171.2 (C=O); MS (EI, 70 eV) m/z 299 (M⁺); anal. (C₁₈H₂₁NO₃·0.1 CHCl₃): C, 69.83; H, 6.83; N, 4.50. Found: C, 69.82; H, 6.98; N, 3.98.

N-(4-Methoxyphenyl)-N-(ethyl)- α -ethyl-4-methoxyphenylacetamide (20a). *n*-BuLi in hexane (1.56 M, 2.0 equiv) was added dropwise to a -78 °C THF (i-propyl)₂NH (2.2 equiv) solution, then warmed to 0 °C. After 0.5 h at 0° C the mixture was re-cooled to -78° C and a THF solution of acetamide 19 (1.0 equiv) added dropwise. After 20 min EtI (1.3 equiv) was added in one portion and the mixture allowed to reach rt. Product isolation (H_2O, Et_2O, Na_2SO_4) and purification by flash chromatography (25% EtOAc/hexane) as light yellow oil (95%): ¹H NMR (CDCl₃) δ 0.78 (t, 3H, J = 7.5 Hz, CH₃ α -ethylacetamide), 1.05 (t, 3H, J = 7.0 Hz, CH₃ N-ethyl), 1.60 (dquint, 1H, J = 14.0, 7.0 Hz, α -CH₂), 2.04 (dquint, 1H, J = 14.0, 7.0 Hz, α -CH₂), 3.27 (dd, 1H, J = 8.5, 7.0 Hz, CH α-benzylic), 3.67 (m, 2H, -NCH₂CH₃), 3.77 (s, 3H), 3.85 (s, 3H), 6.75 (d, 2H, J=8.8 Hz), 6.87 (br s, 3H)4H), 6.98 (d, 2H, J = 8.8 Hz); ¹³C NMR (CDCl₃) δ 12.5 (CH₃), 13.1 (CH₃), 28.5 (CH₂), 44.3 (CH₂), 55.2 (CH₃), 55.6 (CH₃), 113.7 (ArCH), 114.5 (ArCH), 129.2 (ArCH), 132.9 (ArC), 135.0 (ArC), 158.4 (ArC), 159.0 (ArC), 173.5 (C=O); MS (EI, 70 eV) m/z 327 (M⁺); anal. (C₂₀H₂₅NO₃): C, 73.37; H, 7.70; N, 4.28. Found: C, 72.84; H, 7.56; N, 3.59.

N-(4-Methoxyphenyl)-*N*-(ethyl)- α -benzyl-4-methoxyphenylacetamide (20b). Preparation as described in 20a and purification by flash chromatography (15% EtOAc/ hexane) afforded title compound as an oil (86%): ¹H NMR (CDCl₃) δ 0.94 (t, 3H, J = 7.2 Hz, NCH₂CH₃), 2.75 (dd, 1H, J = 12.8, 4.8 Hz, methine <u>H</u>), 3.43 (m, 2H, NCH₂CH₃), 3.58 (dd, 1H, J = 10.2, 5.1 Hz), 3.70 (dd, 1H, J = 13.3, 7.1 Hz) 3.76 (s, 3H), 3.79 (s, 3H), 6.15 (br s, 1H), 6.75 (d, 2H, J = 8.3 Hz), 6.78 (d, 2H, J = 6.7 Hz), 7.09 (m, 4H), 7.24 (m, 3H); ¹³C NMR (CDCl₃) δ 12.8 (CH₃), 41.5 (CH₂), 44.2 (CH₂), 50.3 (CH), 55.2 (OCH₃), 55.4 (OCH₃), 113.7 (ArCH), 114.3 (ArCH), 126.2 (ArCH), 128.2 (ArCH), 129.1 (ArCH), 129.4 (ArCH), 130.0 (ArCH), 132.4 (ArC), 134.6 (ArC), 140.1 (ArC), 158.5 (ArC), 158.9 (ArC), 172.5 (C=O); MS (EI, 70 eV) m/z 389 (M⁺).

N-(4-Hydroxyphenyl)-*N*-(ethyl)-4-hydroxyphenylacetamide (21a). Deprotected according to BBr₃ procedure and purified by chromatography (50% EtOAc/hexane) to afford tan foam (quant): mp 75 °C dec.; ¹H NMR ((CD₃)₂SO) δ 0.96 (t, 3H, *J*=7.5 Hz), 3.16 (s, 2H), 3.36 (q, 2H, *J*=7.5 Hz), 6.59 (d, 2H, *J*=8.5 Hz), 6.78 (m, 4H), 6.97 (d, 2H, *J*=8.5 Hz), 9.18 (br s, 1H), 9.62 (br s, 1H); MS (EI, 70 eV) *m*/*z* 271 (M⁺); anal. (C₁₆H₁₇ NO₃·0.9 H₂O): C, 66.84; H, 6.59; N, 4.87. Found: C, 66.95; H, 6.20; N, 4.77.

N-(4-Hydroxyphenyl)-*N*-(ethyl)-4-hydroxyphenylthioacetamide (21b). Deprotected according to BBr₃ procedure and purified by chromatography (50% EtOAc/hexane) to afford tan powder (84%): mp 156–157 °C; ¹H NMR ((CD₃)₂SO) δ 1.08 (t, 3H, *J*=7.0 Hz), 3.67 (s, 2H), 4.14 (q, 2H, *J*=2 Hz), 6.54 (d, 2H, *J*=8.5 Hz), 6.75 (app t, 4H, *J*=8.0 Hz), 6.87 (d, 2H, *J*=8.5 Hz), 9.17 (s, OH), 9.78 (s, OH): MS (EI, 70 eV) *m*/*z* 287 (M⁺⁺); HRMS calcd for C₁₆H₁₇NO₂S, 287.0980; found, 287.0977; anal. (C₁₆H₁₇NO₂S·0.2H₂O): C, 66.04; H, 6.03; N, 4.81. Found: C, 65.97; H, 6.03; N, 4.78.

N-(4-Hydroxyphenyl)-*N*-(ethyl)-α-ethyl-4-hydroxyphenylacetamide (22a). Deprotected according to BBr₃ procedure and purified by chromatography (50% EtOAc/ hexane) to afford white foam (70%): ¹H NMR (CDCl₃) δ 0.76 (t, 3H. J=7.5 Hz), 1.04 (t, 3H, J=7.0 Hz), 1.54 (m. 1H, CH₂ β-amide), 1.94 (m, 1H, CH₂ β-amide), 3.27 (t, 1H, J=7.0 Hz, CH α-amide), 3.62 (m, 2H, -NCH₂CH₃), 6.63 (d, 2H, J=8.8 Hz), 6.79 (d, overlapping br s, 2H, J=8.8 Hz), 7.05 (br s, 4H); ¹³C NMR (CDCl₃) δ 12.5 (CH₃), 13.0 (CH₃), 28.2 (CH₂), 44.8 (CH₂), 50.7 (CH), 116.1 (ArCH), 128.6 (ArCH), 129.3 (ArCH), 131.7 (ArCH), 133.7 (ArC), 155.4 (ArC), 156.8 (ArC), 174.7 (C=O); MS (EI, 70 eV) *m*/*z* 299 (M⁺); HRMS calcd for C₁₈H₂₁NO₃, 299.1521, found 299.1520.

N-(4-Hydroxyphenyl)-*N*-(ethyl)- α -ethyl-4-hydroxyphenylthioacetamide (22b). Deprotected according to BBr₃ procedure and purified by chromatography (25% EtOAc/hexane) to afford yellow residue (95%): ¹H NMR (MeOD) δ 0.73 (t, 3H, *J*=7.3 Hz), 1.14 (t, 3H, *J*=7.2 Hz), 1.76 (m, 1H, CH₂ β -amide), 2.17 (m, 1H, CH₂ β -amide), 3.62 (dd, 1H, *J*=7.8, 7.0 Hz, CH α amide). 4.16 (m, 1H, CH₂ α -NCO), 4.30 (m, 1H, CH₂ α -NCO), 6.56 (dd, 1H, *J*=8.4, 2.6 Hz), 6.56 (AA'BB', 2H, *J*=8.7, 2.3 Hz), 6.75 (dd, 1H, *J*=8.4, 2.8 Hz), 6.91 (dd, 1H, *J*=8.6, 2.7 Hz), 6.94 (AA'BB', 2H, *J*=8.5, 1.9 Hz), 7.0 (dd, 1H, J=8.7, 2.8 Hz); ¹³C NMR (CDCl₃) δ 11.4 (CH₃), 12.6 (CH₃), 13.9 (CH₂), 52.3 (CH₂), 56.2 (CH), 115.0 (ArCH), 116.3 (ArCH), 128.5 (ArCH), 129.1 (ArCH), 129.9 (ArCH), 133.5 (ArC), 136.5 (ArC), 154.6 (ArC), 155.9 (ArC), 207.5 (C=S); MS (EI, 70 eV) m/z 315 (M⁺); HRMS calcd for C₁₈H₂₁NO₂S, 315.1293; found, 315.1301.

N-(4-Hydroxyphenyl)-*N*-(ethyl)-α-benzyl-4-hydroxyphenylthioacetamide (22c). Deprotected according to BBr₃ procedure and purified by chromatography (15% EtOAc/hexane) to afford yellow residue (47%): ¹H NMR (MeOD) δ 1.05 (t, 3H, J=7.2 Hz), 2.90 (dd, 1H, J=12.9, 4.9 Hz, PhCH₂), 3.66 (dd, 1H, J=12.9, 9.5 Hz, PhCH₂), 3.95 (m, 1H, CH₃CH₂), 4.05 (dd, 1H, J=9.3, 5.2 Hz, PhCH₂CH-), 4.30 (m, 1H, CH₃CH₂), 6.02 (dd, 1H, J=8.4, 2.7 Hz), 6.56 (d, 2H, J=8.3 Hz), 6.66 (overlapping dds, 2H, J=9.2, 3.1 Hz), 6.75 (dd, 1H, J=8.8, 3.0 Hz), 7.00 (d, 2H, J=8.7 Hz), 7.10 (m, 5H); HRMS calcd for C₂₃H₂₃NSO₂, 377.1449; found, 377.1443.

N-(4-Methoxyphenyl)-N-(1,1,1-trifluoro-2-propyl)-4-methoxyphenylacetamide (25). Prepared according to general amidation procedure for CF₃-containing anilines; product isolation and purification via flash chromatography (25% EtOAc/hexane) afforded title compound as light yellow oil (40%): ¹H NMR (CDCl₃) δ 1.16 (d, 3H, J=7 Hz), 3.32 (s, 2H), 3.76 (s, 3H), 3.84 (s, 3H), 5.61 (sept, 1H, J = 7.6 Hz), 6.77 (d, 2H, J = 8.9 Hz), 6.90 (m, 3H), 6.92 (d, 2H, J = 8.9 Hz), 7.06 (m, 1H); ¹³C NMR (CDCl₃) δ 12.5 (CH₃), 40.7 (CH₂), 50.1 (q, CHCF₃, J=31 Hz), 55.3 (CH₃O), 55.6 (CH₃O), 113.9 (ArCH), 114.3 (ArCH), 114.6 (ArCH), 125.1 (q, CF_3 , J = 283 Hz), 127.0 (ArC), 129.9 (ArC), 130.2 (ArCH), 132.0 (ArCH), 158.6 (ArC), 159.9 (ArC), 172.8 (C=O); HRMS calcd for $C_{19}H_{20}NF_3O_3$. 367.1395, found, 367.1394; anal. (C₁₉H₂₀NF₃O₃): C, 62.12; H, 5.49; N, 3.81. Found: C, 61.76; H, 5.58; N, 3.66.

N-(4-Methoxyphenyl)-*N*-(1,1,1-trifluoro-2-propyl)- α -methyl-4-methoxyphenylacetamide (26). n-BuLi in hexane (1.56 M, 2.0 equiv) was added dropwise to a -78 °C THF (*i*-propyl)₂NH (2.2 equiv) solution, then warmed to 0°C. After 0.5 h at 0°C the mixture was re-cooled to -78 °C and a THF solution of N-(1,1,1-trifluoro-2-propyl) substituted acetamide 25 (1.0 equiv) was added dropwise. After 20 min MeI (1.3 equiv) was added in one portion and the mixture allowed to reach rt. Crude product isolation (H₂O, Et₂O, Na₂SO₄) afforded an amber oil (83%) which was determined by both ¹H NMR and HPLC to be an approx. 1:1 ratio of diastereomers for the expected product: ¹H NMR (CDCl₃) δ 1.05 (2-ds, 3H, J = 7.5 Hz), 1.28 (2-ds, 3H, J = 7.5 Hz), 3.42 (overlapping qs, 1H, J = 7.5 Hz), 5.58 (m, 1H, $CHCF_3$), 6.44 (2-dds, 1H, J = 8.8, 2.8 Hz), 6.68 (dd, 0.5 H, J = 8.8, 2.8 Hz), 6.70 (m, 2.5H), 6.81 (m, 2H), 7.01 (2-dds, 1H, J=8.4, 2.3 Hz), 7.20 (2-dds, 1H, J=8.4,2.3 Hz); HPLC (254 nm) 25% EtOAc/hexane, Micropac SiO₂ column: 5.82 min. (46.2%), 7.19 min. (53.7%), $sd\pm5\%$.

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-propyl)- α -methyl-4-hydroxyphenylacetamide (27). Deprotection using BBr₃ procedure and chromatography (35% EtOAc/ hexane) afforded title compound as a white powder (95%) with a 1:1 ratio of diastercomers as found by ¹H NMR: mp 165–170 °C; ¹H NMR (MeOD) δ 1.10 (overlapping ds, 3H, *J*=7.5 Hz), 1.25 (overlapping ds, 3H, *J*=7.5 Hz), 3.49 (overlapping qs, 1H, *J*=7.5 Hz), 5.52 (m, 1H, CHCF₃), 6.43 (2-dds, 1H, *J*=8.8, 2.8 Hz), 6.57 (dd, 0.5H, *J*=8.8, 2.8 Hz), 6.64 (m, 2.5H), 6.72 (m, 2.5H), 6.86 (app td, 1H, *J*=8.4, 2.3 Hz), 7.07 (2-dds, 1H, *J*=8.4, 2.3 Hz); MS (EI, 70 eV) *m*/*z* 353 (M⁺); HRMS calcd for C₁₈H₁₈NF₃O₃, 353.1238; found, 353.1235.

N-(4-Hydroxyphenyl)-*N*-(1,1,1-trifluoro-2-propyl)-4-hydroxyphenylthioacetamide (28). Carboxamide 25 (0.3 mmol, 100 mg) was treated with Lawesson's reagent as outlined in general procedure. Product isolation (H₂O, Et₂O, Na₂SO₄) and radial chromatography (20% EtOAc/hexane) afforded 5 mg of protected thiocarboxamide in low yield (5%). Subsequent deprotection using BBr₃ procedure afforded the title compound as a light yellow residue upon SiO₂ purification (30% EtOAc/ hexane, quant): ¹H NMR (MeOD) δ 1.24 (d and m, 3H, *J*=7.8 Hz), 3.85 (ABq, 2H, *J*=13 Hz), 5.56 (sept, 1H, *J*=7.8 Hz), 6.72 (m, 4H), 6.85 (m, 4H); HPLC: Whatman C18 column, 90%MeOH:10%H₂O R₁= 3.123 > 98% pure at 254 nm.

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Solid-Phase Synthesis of Tetrasubstituted Pyrazoles, Novel Ligands for the Estrogen Receptor

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Most ligands for the estrogen receptor (ER) are not well suited for synthesis by combinatorial means, because their construction involves a series of carbon-carbon bond forming reactions that are not uniformly high yielding. In previous work directed to overcoming this limitation, we surveyed various phenol-substituted five-membered heterocycles, hoping to find a system that would afford both high ER binding affinity and whose synthesis could be adapted to solid-phase methods (Fink et al. *Chem. Biol.* **1999**, *6*, 206-219.) In this report, we have developed a reliable and efficient solid-phase method to prepare the best of these heterocycles, the tetrasubstituted pyrazoles, and we have used this methodology to produce small, discrete libraries of these novel ER ligands. We used a combination of FT-IR and nanoprobe ¹H NMR-MAS to characterize intermediates leading up to the final pyrazole products directly on the bead. We also developed a scavenging resin, which enabled us to obtain products free from inorganic contaminants. We prepared a 12-member test library, and then a 96-member library, and in both cases we determined product purity and ER binding affinity of all of the library members. Several interesting binding affinity patterns have emerged from these studies, and they have provided us with new directions for further exploration, which has led to pyrazoles having high affinity and potency as agonists and antagonists toward the ER α subtype.

Introduction

In recent years, combinatorial chemistry, with its ability to generate a large set of structurally related analogues, has become a bona fide tool for increasing productivity in the functional assessment of compound libraries and the rapid development of structure-activity relationships.¹ Combinatorial techniques for preparing peptide libraries are well established, but methods for the combinatorial synthesis of small-molecule libraries for the development of useful pharmaceuticals remains a formidable challenge. Small molecules are generally not oligomeric, and a diverse range of chemical transformations may be required for their synthesis. Therefore, the development of an appropriate parallel synthesis format is not always straightforward. This limitation is increasingly encountered in the quest for chemical diversity, because the fidelity of a given library member can be compromised when the building block components are not uniformly reactive. Consequently, there has been an active quest to develop the sort of truly general, high viclding transformations necessary to maintain high quality in the preparation of small-molecule libraries.

Novel estrogens having tissue selective action suitable for menopausal hormone replacement or the treatment and prevention of breast cancer are actively being sought by the pharmaceutical industry.² These agents, often referred to as selective estrogen receptor modifiers or SERMs, generally consist of a nonsteroidal core structure onto which is



Figure 1. Estrogen receptor- α selective pyrazole discovered from a solution-phase study. RBA is relative binding affinity for the estrogen receptor; for estradiol, RBA = 100%.

appended a basic or polar function.³ Most ligands for the estrogen receptor (ER), however, are not well suited for synthesis by combinatorial means, because their construction generally involves a series of carbon—carbon bond forming reactions that are not uniformly high yielding, nor necessarily easily adaptable to solid-phase synthesis methods. Two small combinatorial libraries of nonsteroidal ER ligands have been reported,^{4,5} but these efforts produced either low affinity ligands or ones of limited structural diversity.

To circumvent these current limitations and to expand the possible combinatorial approaches to ER ligands, we have proposed a novel modular ER pharmacophore consisting of a *variable* core structure, onto which are linked four independent substituents of defined variability. Using this paradigm, we have investigated core structures consisting of five-membered heterocycles, as well as other functionalitics.⁶ Among the five-membered systems that we have studied to date, we found that 1,3,5-triaryl-4-alkyl-pyrazoles, such as compound 1 (Figure 1), showed high binding affinity for the estrogen receptor, and intriguingly this compound had 100-fold higher potency as an agonist in a cell-based transcription assay through ER α than through ER β .⁷ Com-

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Scheme 1. Routes for the Preparation of the 12-Member and the 96-Member Pyrazole Libraries



pounds that show high ER subtype selectivity are in great demand as agents to define the biological roles of these subtypes⁸ and as potential tissue-selective estrogens for the uses noted above.

Because of these interesting initial findings, we wished to use a combinatorial approach to explore this pyrazole template further, to gain a better understanding of its structure—binding affinity pattern with the ER and to discover additional compounds with ER subtype selectivity. In particular, we hoped that such an expanded study would enable us to determine the preferred binding orientation for these pyrazoles in the ER, an issue that was not settled by our initial investigations.⁶ We required this information to guide our future placement of pharmacophore elements in the design of selective antiestrogens (Stauffer, S. R., Huang, Y., Aron, Z., Coletta, C. J., Sun, J., Katzenellenbogen, B. S., Katzenellenbogen, J. A., unpublished).

As we had noted in their original conception,⁶ the tetrasubstituted pyrazoles offer an attractive template for the combinatorial development of ligands for the estrogen receptor. Their heterocyclic nucleus provides a core upon which several common substructural motifs found in high affinity ER ligands can adequately be displayed in several possible configurations, and their synthesis involves simple condensation reactions, some of which have already been demonstrated on solid phase, although mostly for trisubstituted pyrazole derivatives.^{9–12}

This work describes our efforts in the design and preparation of libraries of tetrasubstituted pyrazoles as ER ligands and our evaluation of the binding affinities of the library members. These studies have led us to a new series of high affinity compounds that display very high ER α agonist selectivity in cell-based transactivation assays, the details of which will be reported elsewhere (Stauffer, S. R., Sun, J., Katzenellenbogen, B. S., Coletta, C. J., Katzenellenbogen, J. A., unpublished).

Results and Discussion

Library Strategy. It was our intention to prepare discrete analogues (96-member libraries) of our tetrasubstituted pyrazole pharmacophore, noted above, in a parallel splitsplit format to produce material for ER binding affinity assays. Synthetically, the classical 1,3-dione-hydrazine condensation route (illustrated in Scheme 1) seemed attractive, because it is well precedented and can be accomplished under conditions that are potentially adaptable to solid-phase synthesis.⁶ Also, the β -diketone component can be readily generated from a crossed-Claisen condensation reaction.

Using this route, we can introduce molecular diversity in the target pyrazole in either the alkylphenone, the ester, or the hydrazine component. Our choice to introduce the C-4 alkyl group early on within the ketone, rather than by alkylating the β -diketone intermediate, was based on solution feasibility studies; in contrast to the pyrazole synthesis strategy reported by Marzinzik and Felder,⁹ we found that alkylation was not general for the preparation of 1,3-diarylpropane-1,3-diones. Our approach is also attractive because many of the 4'-methoxy-alkylphenone precursors are commercially available or can be produced in one step via Friedel–Crafts acylation reactions.

Linker. We selected the phenol component as the site for attachment of the pyrazole to the polymer support because it is convenient and it is the one functional group that is present in all of the pyrazoles. Merrifield's resin was used as the solid support, so the tethered phenol became attached to the resin as a p-substituted benzyl ether.

Our overall linker strategy is advantageous for two reasons: First, we knew from solution studies for tetrasubstituted pyrazoles of this type that a robust linker would be required to withstand the conditions for pyrazole formation (DMF/THF 120 °C).⁶ Thus, we expected that the stable benzyl ether link would be more satisfactory than other more labile linkers. Second, we anticipated that the strong acidic 320 Journal of Combinatorial Chemistry, 2000, Vol. 2, No. 4



Figure 2. Components for 12-member library.

conditions required for product cleavage from the resin (i.e., BBr₃) could be adjusted to cleave, as well, any methyl ethers protecting other phenolic components. This approach is similar to that reported by Green for the preparation of lavendustin analogues,¹³ but different in that our linker is a benzyl ether rather than a benzyl ester.

Method Development. The adaptation of the pyrazole synthesis from solution phase to solid phase required extensive optimization and later modification. Our initial route is shown in Scheme 1, and it was used for the preparation a small 12-member library, which was intended to serve as a standard for later, larger library design. In addition, this smaller library was developed to evaluate the generality of the reaction conditions by using various electronically demanding building blocks (Figure 2). The conditions described below were initially developed using our original lead compound 1 as a model.

Few starting alkylphenones (2, Scheme 1) are available commercially, but they were readily prepared via Friedel– Crafts acylation, using anisole and the desired acid chloride. Thus, the protected *p*-methoxy-alkylphenone 2 was obtained and then demethylated using AlCl₃ to afford 3, which was subsequently loaded onto Merrifield's resin (Novabiochem Inc., 1.39 mmol/g), according to the method of Ellman and co-workers.¹⁴ Formation of the resin-bound ketone 4 was ascertained using standard techniques (FT-IR, null %Cl combustion analysis), and a portion was cleaved with BF₃– SMe₂ to determine the loading capacity based on mass recovery. After minimal workup, the expected ketone was cleanly observed by ¹H NMR and HPLC (purity >99%), and the loading capacity was found to be between 0.9 and 1.1 mmol/g.

Following ketone immobilization, we used a crossed-Claisen condensation reaction to form the requisite β -diketones (6). This transformation proceeds satisfactorily using excess lithium hexamethyldisilazide (LHMDS) and activated *p*-nitrophenyl esters (**5a**,**b**,**d**) in THF. Other bases, such as NaNH₂ or KOt-Bu, were less effective, particularly with higher alkylphenones (e.g., propiophenone and higher). In addition, *p*-nitrophenyl activated esters proved to be most general in providing high yields of the dione. In most cases, the active ester could be present in the reaction vessel during the addition of base. However, esters with electron with-drawing groups, such as compound **5d** (Figure 2), required preformation of the resin-bound enolate prior to the addition of the ester, because of competing nucleophilic addition of the disilamide anion to the activated ester.

We found nanoprobe ¹H NMR-MAS to be an indispensable tool for following β -dione formation on solid-phase and for optimizing reaction conditions. Although FT-IR was useful for monitoring ketone immobilization, we did not find it satisfactory for evaluating the progress of dione formation. We had expected that a shift in the C=O stretch of the alkylphenone at 1675 cm⁻¹ would occur as the dione formed and equilibrated with its intramolecularly hydrogen bonded enol tautomer. We were surprised, however, when the ¹H NMR spectrum of 1,3-bis(4-methoxyphenyl)-2-ethyl-1,3propanedione showed that these α -substituted β -diketones exist exclusively in the diketo tautomeric form in solution. We presume that this is due to the A-strain present in these substituted molecules, because ¹H NMR analysis shows that the less hindered 1,3-diaryl-1,3-propanediones, which do not bear a 2-alkyl substituent, exist as mixtures of both enol and diketo tautomers. In most cases FT-IR revealed a C=O doublet for the dione. However, the intensity of this signal was often weak and variable, and thus was not a reliable indicator of reaction progress. In contrast, by using ¹H NMR-MAS and nanoprobe technology, we were able to conveniently and quickly ascertain the level of β -diketone formation.

Shown in Figure 3 panel A (top) is the spectrum for the starting resin-bound 4-hydroxy-butryophenone, and in panel



Figure 3. Panel A (top): ¹H NMR-MAS of polymer-bound butryophenone derivative **11**. Panel B (bottom): ¹H NMR-MAS of β -diketone product **12b** after Claisen condensation reaction.

B (bottom) is the spectrum for the expected dione product after Claisen condensation with 4-methoxy-(4'-nitrophenyl)benzoate. Because we are using a short linker, which holds these molecular segments close to the rigid polystyrene backbone, the line widths for these signals remain fairly large (20 Hz), and as a result, coupling information is unavailable. Despite this limitation, however, we can readily discern diagnostic chemical shift differences between the product and starting material, so that we confidently assess the success or failure of the reaction.

To form the tetrasubstituted pyrazole on solid phase, we first tried conditions that had been successfully used for the solution-phase synthesis (heating the dione and hydrazine hydrochloride salt to 110-120 °C in DMF/THF solutions for up to 16 h), as well as conditions developed my Marzinzik and Felder for solid-phase pyrazole synthesis (DMA as a solvent at 80 °C).9 Unfortunately, even with extended heating, neither of these methods gave the desired heterocycle. When we used triethylamine (TEA) to neutralize the hydrazine hydrochloride and DMF as solvent, we obtained low yields of the desired product, but purity after cleavage/ deprotection was less than 35%. Ultimately, we found that toluene as solvent with the addition of TEA was effective. affording the pyrazole product 1 in >90% purity after cleavage/deprotection. One should note that subsequently we found that these reaction conditions, which involved heating toluene to 80 °C for 9 h, were not compatible with our 96well reaction plate (Polyfiltronics 96-well Unifilter plate). Nevertheless, at the time we used these conditions to prepare

 Table 1. HPLC Purity Determination and Estrogen Receptor

 Binding Affinities (RBA Values) of C(4) *i*-Butyl Pyrazole

 Library



^a RP-HPLC, 80% MeOH:H₂O, flow rate of 1.0 mL/min, detection at 254 nm (performed before purification by radial chromatography). ^b Relative Binding Affinity (RBA) values determined by a modification of our standard competitive binding assay using only three concentrations of ligand; all compounds tested were at >80% purity. ^c Independent assays performed on the two individual regioisomers.

a 12-member library of pyrazoles and then later modified the pyrazole-forming step to make it compatible with our 96-well reaction plate (see below).

The diones (6) needed for this (and the larger library) were prepared in a single batch, using a homemade reaction block capable of holding 16–20 mL sealed conical polypropylene tubes as reaction vessels. The block was rotated in an oven at 40 °C for 4 h, using a modified rotary evaporator motor. After washing the resins and drying them overnight, we verified dione formation by nanoprobe ¹H NMR-MAS. Each dione resin was then split into the appropriate number of portions and reacted with the appropriate hydrazine, the cyclization products then being cleaved/deprotected with BBr₃. For this smaller 12-member library, the cleaved material was collected, treated to a minimal workup (MeOH, passage through a SiO₂ plug), and analyzed for purity by HPLC.

Shown in Table 1 are the HPLC-evaluated purities and the ER binding affinities for the C-(4)-*iso*-butyl pyrazoles (**8a**-**I**) in the 12-membered library. The HPLC purity values listed were obtained on the pyrazoles after only minimal workup, yet some of these are quite high (>90%). The pyrazoles derived from *tert*-butylhydrazine were of the lowest quality. The major byproduct in those having lower purity was the corresponding β -diketone intermediate, and in some cases small amounts of the starting ketone. Prior to binding affinity determination, all compounds were purified by radial chromatography, so that their purities were at least 80%. The molecular ions of all purified pyrazoles in the C(4) *i*-butyl series were also verified by ES-MS.

The binding affinities of these pyrazoles were determined in a competitive radiometric binding assay (for details see Experimental Section), using [³H]estradiol as tracer and lamb uterine cytosol as a source of ER, and they are expressed as relative binding affinity values (RBA), with estradiol having an RBA of 100%. In the past, we have found that repeat determinations of binding affinities by this assay have a coefficient of variation of 0.3.

We were concerned that even small amounts of contaminants in the purified pyrazoles might be affecting the RBA determinations. To test this, we examined the effect of dihydroxyphenyl dione 9 on the affinity determination of pyrazole 1. This dione (9) itself binds to the ER only very weakly (RBA = 0.012%), and when it was added to pyrazole 1 at 10, 20, and 30%, we noted no effect on the RBA value obtained for pyrazole 1. Thus, we believe that the binding affinity values we have determined for all of these pyrazoles (Table 1) are valid.



Overall, the *i*-butyl pyrazoles bind to ER with reasonable affinity (Table 1), and even within this small set of compounds some structure—affinity trends are apparent. Clearly, there is a primary preference for hydroxy substitution at R_1 , as in most cases, and the pyrazoles in the **8e**-**h** series bind better than those in the **8a**-**d** and **8i**-**l** series. Bromine substitution (series **8i**-**l**) is unusual for nonsteroidal ligands, but pyrazole **8j** (1.4:1 ratio of regioisomers) has a reasonable affinity of 3.3%. The reasonable affinity of the *tert*-butyl pyrazole **8h** suggests that bulky substituents other than phenyl are tolerated at R_2 . The highest affinity members of this small pyrazole library, **8b** and **8f**, contain two and three hydroxyl substituents, respectively. The similar but lower affinity diphenolic pyrazole **8e** suggests that two distinct binding orientations may exist for **8e** and **8b**.

In those cases where more than one regioisomer was expected and they could be separated by HPLC, two RBA values are indicated. In three cases where we could do this, one regioisomer showed a higher affinity: In both the monohydroxy pyrazole **8a** and dihydroxy pyrazole **8b**, a 3-10-fold preference was found for one regioisomer, and in the case of pyrazole **8i**, this preference is much greater.

At this point, we could not make a definitive assignment of the structure of these regioisomers, but these results suggest that only one of them is providing an effective mimic of the A-ring of estradiol. In other work, we have carried out independent, regioselective syntheses of single pyrazole regioisomers in a related series, and we have conducted molecular graphics modeling studies of how these isomers might fit into the ligand binding domain of ER (Stauffer, S. R., Huang, Y., Coletta, C. J., Katzenellenbogen, J. A., unpublished). These studies suggest that there are indeed preferred orientations of these regioisomers within the ER binding pocket. However, it appears that the preferred orientation may switch, depending on the substituent display.

Adaptation to a 96-Well Format. To prepare the 96member library, we used the Polyfiltronics 96-well Unifilter plate. The Unifilter plate contains a single underlying membrane which is designed to hold back most organic solvents, except when a vacuum is applied. Unfortunately, the Polyfiltronics plate did not withstand our conditions for pyrazole formation, using prolonged heating with toluene at 80 °C.

In our search for alternate conditions for pyrazole formation which the Polyfiltronics plate would withstand, we used pyrazole 1 as a model, and we explored a greater number of solvents and solvent mixtures for pyrazole formation (THF, BuCN, HC(OMe)₃, CH₃NO₂, and alcohols) with and without various additives (TiCl₄, Na₂SO₄, and 4 Å molecular sieves), at room temperature and at elevated temperatures. Ultimately, only alcohol solvents proved to be effective in forming the pyrazole product and were compatible with the Unifilter plate.

In this larger library, we also wanted to develop an expedited method for product isolation, whereby we could quench the cleavage/deprotection reagent BBr₃ and neutralize the HBr generated without introducing water, thereby avoiding cumbersome liquid—liquid phase extractions. Our approach was to develop a scavenger resin to neutralize the HBr and remove the inorganic contaminants by sequestration. Our initial efforts were based on work by Cardillo and coworkers,¹⁵ involving the use of macroreticular carbonate resin as a reagent in the hydrolysis of alkyl halides.

Resin **10a** (Scheme 2), which is readily prepared from the chloride form of the ion exchange Amberlyst A-26 resin, was effective in quenching the BBr₃, but not surprisingly the final product was contaminated with NaBr. However, by modifying the resin to a sodium-free, bicarbonate form, we could still neutralize the HBr, but now with the liberation of only CO₂ and H₂O. Moreover, the polymer-bound quaternary ammonium groups sequestered any bromide ions from solution. Thus, by treating the crude cleaved/deprotected product with this bicarbonate resin, we could remove all of the inorganic contaminants, leaving the desired pyrazole remaining in solution together with only MeOH (bp 65 °C), B(OMe)₃ (bp 58 °C), and a small amount of H₂O, so that solvent removal gave the product free from any reagent contamination.

Two methods were used to prepare the bicarbonate form of A-26 resin (Scheme 2). Both resins (10b and 10c) afford the product pyrazole in reasonable purity, but the bicarbonate resin which was prepared using $NaHCO_3$ (10b) still contained small amounts of Na ions, risking product contamination with NaBr. The preferred method to prepare Na ion-free resin was first to convert the chloride form of the A-26 resin to the hydroxide form and then generate the bicarbonate resin by bubbling CO_2 through an aqueous suspension of the resin. This ensured that there was no Na ion contamination, and this material (10c) afforded the pyrazole product with somewhat better purity. Shown in Scheme 1 is the overall optimized solid-phase synthesis route to the 96-member pyrazole library. The notable modifications from the original route used to prepare the 12-member library are the conditions for the pyrazole-forming step and final workup procedure involving the bicarbonate resin 10c.

The individual components chosen for the 96-member library are shown in Figure 4. The components used to 1



Figure 4. Components for 96-member pyrazole library.

Scheme 2. Development and Use of Bicarbonate Resin for HBr Neutralization



10c (resin contains 0.02 % Na)
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Figure 5. HPLC purities for 96-member ER library.

prepare the *i*-butyl library (Figure 2, Table 1) were again included as standards to measure the success and reproducibility of the synthesis and to verify the RBA assays. The progress of pyrazole formation for suspected "worst-case" combinations, such as CF₃-phenyl and *tert*-butyl hydrazines reacting with halogen-substituted diones, was monitored using FT-IR. The disappearance of the C=O signal at 1670 cm⁻¹ was a reliable marker for determining the progress of individual reactions. Unfortunately, for several CF3-substituted hydrazines we were unable to drive the reaction to completion, even after subjecting these resins to fresh reagent and heating for an additional 20 h. In any event, cleavage/ deprotection with BBr3 followed, and reactions were carefully quenched with MeOH and then incubated with bicarbonate resin 10c for 1 h at 50-60 °C to ensure complete HBr neutralization and bromide ion sequestration. After being cooled, the pyrazole products were collected and concentrated, then reconstituted in 1 mL McOH and analyzed using a standard, steep gradient elution on a high throughput reverse-phase HPLC column. Because the pyrazoles are uniformly fluorescent, we used fluorescence detection in tandem with UV detection to identify which of the eluted peaks was the pyrazole. This proved to be a robust approach to characterizing product purity. After obtaining the purity for each member, we determined the ER binding affinity in a simplified, three-point assay (see Experimental Section for dctails).

Purity and ER Binding Affinity Relationships. A summary of the HPLC purities for the final pyrazoles is shown in Figure 5, according to gray scale coded ranges. The average purity for the library was $50\% (\pm 15\%)$. This is not an unreasonable level of purity when you consider that this library included components, such as *tert*-butyl hydrazine hydrochloride, which we have found to be less reactive than the aromatic hydrazines. As before, the principal impurities could be identified as the unreacted dione precursors, which we had previously shown did not affect the results of the binding assay (see above). On the basis of product yields,

p-CF₃-substituted phenyl hydrazines appear to be even less reactive than *t*-butyl hydrazine hydrochloride, because this group of pyrazoles had the lowest overall purity of the whole library. Where regioisomers were expected within this series (see above), it was difficult to assign the product peaks, even with the help of fluorescence as a marker. In these cases, we assumed the purity to be 30%.

Shown in Figure 6 are the relative binding affinity (RBA) values for the 96-member library, indicated as ranges according to a gray scale coded legend. Several members of the library showed appreciable affinity for the ER. Particularly gratifying was the fact that most members of the 12-member *i*-butyl control library (Table 1) had RBA values which were reproduced quite well in the larger library (i.e., within 30%, relative), with the exception of **8f** (7.6%) vs **B-8** (23%). A repeat assay **8f** showed that its determination in the original 12-member library was low. The binding affinity of 16 additional select members was also re-tested after chromatographic purification (>80%), and the RBA values for these members also agreed quite well with the original determinations on the crude isolated products.

The use of an affinity array chart in Figure 6 permits a rapid, visual assessment of binding affinity patterns. For example, it is readily apparent that, for pyrazoles in both the R_1 ethyl and *i*-butyl series (column 2 and 8), those with HO substituents at R_2 have, overall, the highest affinity. Within these two series (columns 2 and 8), a number of substituents are tolerated at R_3 , the best being p-HO-C₆H₄, for which the two highest affinity pyrazoles are represented, pyrazole **B-2** (14%) and **B-8** (23%). For pyrazoles with R_3 = p-HO-C₆H₄ (row B), a number of substituents are tolerated to varying degrees; those with fluorine substituents on R₂ for both the ethyl and *i*-butyl series bind moderately well (RBA = 1-5%). For members with $R_1 = i$ -butyl and $R_3 = p-HO-C_6H_4$, even more polar substituents appear to bind well. Particularly noteworthy is the *m*-HO analogue **B-9**, which has a relative binding affinity of 6.8%, and the fluoro derivative B-11. By contrast, few other R₃ substituents are



Figure 6. ER binding affinity for 96-member pyrazole library. (In unsymmetrical systems, i.e., $R_2 \neq p$ -OH, the average regioisomer ratio was 1.4 to 1.)

well tolerated by the ER, one exception, however, being the *tert*-butyl pyrazoles **H-2** and **H-8**, which each have affinities of 6–7%. As was the case with pyrazole **8j** from the 12-member *i*-butyl library, several pyrazoles with *p*-bromo substituents at R_2 have reasonable affinity ($\sim 2-3\%$). However, these affinities are much lower when $R_1 = ethyl$ (RBA <1%), than when $R_1 = i$ -butyl.

In investigations to be reported elsewhere (Stauffer, S. R., Coletta, C. J., Sun, J., Katzenellenbogen, B. S., Katzenellenbogen, J. A., unpublished), we have made other analogues based on the high affinity 1,3,5-*p*-hydroxyphenyl-pyrazole template, and we have found a compound with an RBA of 30% that shows very high affinity and potency selectivity for ER α . We have also prepared other pyrazoles that act as ER α potency selective antagonists (Stauffer, S. R., Huang, Y., Aron, Z. D., Coletta, C. J., Sun, J., Katzenellenbogen, B. S., Katzenellenbogen, J. A., upublished).

Conclusions

We have developed a consistent and efficient method to prepare and isolate tetrasubstituted pyrazoles using a solidphase strategy, and we have used this approach for the production of small, discrete libraries of estrogen receptor (ER) ligands. A combination of FT-IR and nanoprobe ¹H NMR-MAS allowed us to characterize intermediates leading up to the final pyrazole products directly on the bead. We also developed a scavenging resin to afford the cleaved/ deprotected products free from inorganic contaminants. Using this approach, we prepared a 12-member test library, and then a more extensive 96-member library, and we determined product purity and ER binding affinity of all the library members. Several interesting binding affinity patterns emerged from these studies, and they have provided us with clear directions for further exploration of these tetrasubstituted pyrazoles, through which we have found a high affinity and high potency agonist with excellent selectivity for ER α and pyrazoles that selectivity antagonize ER α .

Experimental Methods

General. Melting points were determined on a capillary apparatus and are uncorrected. All reagents and solvents were obtained from Aldrich, Fisher, or Mallinckrodt. Tetrahydrofuran was freshly distilled from sodium/benzophenone. Dimethylformamide was vacuum distilled prior to use and was stored over 4 Å molecular sieves. Et₃N was stirred with phenylisocyanate, filtered, distilled, and stored over 4 Å molecular sieves. All reactions were performed under a dry N₂ atmosphere unless otherwise specified. Radial preparativelayer chromatography was performed on a Chromatotron instrument (Harrison Research, Inc., Palo Alto, CA) using EM Science silica gel Kieselgel 60 PF254 as adsorbent. Flash column chromatography¹⁶ was performed using Woelm 32-63 μ m silica gel packing. Resin bound ketone 11 was prepared according to procedure outlined in Scheme 1 from commercially available 4'-methoxybutyrophenone or from anisole and the acid chloride.

¹H and ¹³C NMR spectra were recorded on 400 or 500 MHz spectrometers, using CDCl₃ or MeOD as solvent. Chemical shifts were reported as parts per million downfield from an internal tetramethylsilane standard ($\delta = 0.0$ for ¹H) or from solvent references. NMR coupling constants are reported in hertz. ¹³C NMR data were determined using either the Attached Proton Test (APT) experiment or standard ¹³C pulse settings. Low resolution electrospray mass spectrometry was performed using a VG Quattro (quadrupole-hexapole-quadrupole, QHQ) mass spectrometer system (Fisons Instruments, VG Analytical; Manchester, U.K.).

FT-IR Analysis and Nanoprobe ¹**H NMR-MAS Spectra.** FT-IR analysis was performed on an FTIR spectrometer, and absorption bands are reported in cm⁻¹. Infrared analysis was accomplished by placing approximately 1–2 mg of resin between two NaCl plates, swelling the beads in CHCl₃, and immediately recording an FT-IR spectrum. Solid-phase NMR spectra were obtained on a Varian 500 MHz wide-bore spectrometer using a nanoprobe. The polystyrene-bound

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dionc intermediates 12a-1 were characterized using 2 mg or less of resin in a pre-swollen CDCl₃ solution (nanotube, approximately 40 μ L total volume). The sample was spun about the magic angle (54.7 °C) at 2.3–2.5 kHz, and four scans were collected for each ¹H experiment, using a pulse width of 15 μ s and delay time of 4 ms. Tetramethylsilane was used as an internal reference.

Relative Binding Affinity Assay. Ligand binding affinities (RBAs) using lamb uterine cytosol as a receptor source were determined by a competitive radiometric binding assay. This assay is conducted in 96-well microtiter plates in a total volume of 70 μ L, using ca. 1 nM of estrogen receptor binding equivalents, 10 nM [3H]estradiol as tracer, and dextran-coated charcoal as an adsorbent for free ligand.¹⁷ On the basis of interspecies comparison, lamb uterus is thought to contain predominantly estrogen receptor subtype α .^{18,19} All incubations were done at 0 °C for 18-24 h. Binding affinities are expressed relative to estradiol (RBA = 100%) and are reproducible with a coefficient of variation of 0.3. RBA determinations for the 96-member library were performed using a three-point assay, at ligand concentrations of 10^{-4} , 10^{-6} , and 10^{-8} M and were corrected for individual purity level.

Chemical Syntheses. 4'-Methoxy-4-methyl-valerylophenone (2). To a stirred solution of AlCl₃ (6.4 g, 48 mmol) in 1,2-dichloroethane at 0 °C was added 4-methyl-valeryl chloride (5.3 g, 39.0 mmol) dropwise over approximately 10 min. The resulting solution was warmed to room temperature, stirred for 0.5 h, then re-cooled to 0 °C, and a 1,2-dichloroethane solution of anisole (5.1 mL, 46 mmol in 20 mL) was added dropwise. The reaction was stirred at room temperature for 9 h, then cooled to 0 $^{\circ}$ C, quenched with H₂O, and extracted with CH₂Cl₂. The organic layer was washed with saturated NaHCO₃ and brine, then dried over MgSO₄, and concentrated to afford product as a pale yellow oil (6.0 g, 75%): ¹H NMR (CDCl₃) δ 0.94 (d, 6H, J = 6.0), 1.60 (m, 3H, overlapping methine and β -CH₂), 2.90 (t, 2H, J =6.8), 3.87 (s, 3H), 6.92 (d, 2H, J = 8.8), 7.90 (d, 2H, J =8.8); ¹³C NMR (CDCl₃) δ 22.6, 28.5, 33.9, 36.4, 55.2, 113.5, 130.2, 163.6, 198.7; MS (EI, 70 eV) m/z 206.1 (M⁺). Anal. (C₁₃H₁₈O₂): C, 75.69; H, 8.80. Found: C, 75.63; H, 8.59.

4'-Hydroxy-4-methyl-valerylophenone (3). To a stirred solution of AlCl₃ (19.4 g, 145 mmol) in benzene (300 mL) at 0 °C was added a benzene solution (60 mL) of ketone **2** over 20 min. The reaction mixture was then brought to reflux for 2 h. After cooling to room temperature the mixture was poured over 400 mL of water, and then the organic layer washed with saturated NaHCO₃ and brine. The organic layer was then dried over NaSO₄ and concentrated to afford the phenolic product as a tan solid which was then used directly in the next step to functionalize Merrifield's resin (8.97 g, 96%): ¹H NMR (CDCl₃) δ 0.94 (d, 6H, J = 6.0) 1.27 (m, 3H, overlapping methine and β -CH₂), 2.90 (t, 2H, J = 8), 6.87 (d, 2H, J = 8.5), 7.91 (d, 2H, J = 9.0).

Alkylphenone Loading onto Merrifield Resin (4, 11). To a solution of the phenolic ketone (28.0 mmol, either 3 or 4'-hydroxy-butyrophenone) in DMF (144 mL) at 0 °C was added NaH as a 60% oil dispersion (1.05 g, 26.4 mmol). The reaction mixture was returned to room temperature;

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Merrifield resin (5.74 g, 1.39 mmol/g Novabiochem, Inc.) was added, and the reaction was heated to 55 °C for 22 h. The solution was then cooled to room temperature and quenched by addition of methanol (150 mL). The resin-bound product was then isolated via vacuum filtration and rinsed with methanol/DMF 1:1 (1 \times 300 mL), DMF (3 \times 350 mL), CH_2Cl_2 (3 × 350 mL), and methanol (3 × 350 mL). The resulting salmon-colored resin was then dried in vacuo for 18 h to afford >6 g of immobilized ketone. FT-IR analysis revealed the C=O stretch at 1670 cm^{-1} , which verified that the resin-bound ketone was produced. In addition, %Cl analysis was performed to verify the absence of halogen, indicating that most of the chloromethyl sites were substituted. Loading capacity, which ranged from 0.9 to 1.1 mmol/ g, was determined from the mass recovery of expected product by cleaving a small portion of the ketone resin using BF₃-SMe₂.

General Procedure for Preparation of *p*-Nitrophenyl Benzoate Esters (5a-f). To a mechanically stirred solution of the substituted benzoic acid (52.6 mmol), *p*-nitrophenol (22.2 g, 157.8 mmol), and DMAP (3.2 g, 26.3 mmol) in 100 mL CH₂Cl₂ at 0 °C was added diisopropylcarbodiimide (57.9 mmol) dissolved in CH₂Cl₂ (100 mL) dropwise over 10 min. The resulting solution was allowed to stir for 18 h at room temperature. The reaction mixture was filtered, and the filtrate was concentrated under reduced pressure. The crude esters were recrystallized from EtOH or EtOAc to afford the desired ester (55-95%). Products were judged pure by comparison to literature melting points²⁰ and then used in the next step of the synthesis.

2-Ethyl-1,3-bis-(4-hydroxyphenyl)-propane-1,4-dione (9). The methoxy-protected derivative of 1 (prepared in 95%) yield from 4-methoxybutyrophenone and the p-nitrophenylbenzoate ester using LHMDS, under the same conditions used for the solid-phase synthesis; see 12a-l, and purified by flash chromatography (ethyl acetate/hexanes)) was treated with 5 equiv of 1.0 M BBr₃ in CH₂Cl₂ at 0 °C. The mixture was allowed to reach room temperature and stir for 5 h. The reaction was then re-cooled to 0 °C and carefully quenched with water and repeatedly extracted with Et₂O. The combined organic layers were dried over Na2SO4, and upon solvent removal the title compound was afforded as a tan foam: ¹H NMR (MeOD) δ 0.99 (t, 3H, J = 7.5), 2.01 (q, 2H, J = 7.5) 5.32 (t, 1H, J = 1.5), 6.81 (AA'BB', 2H, J = 7.0, 3.0), 7.90 (AA'BB', 2H, J = 6.5, 3.0); ¹³C NMR (MeOD) δ 11.4, 23.0, 56.6, 115.1, 128.0, 130.9, 162.8, 195.9; MS (EI, 70 eV) m/z 284 (M⁺). Anal. (C₁₂H₁₆O₄): C, 71.28; H, 5.67. Found: C, 70.90; H, 5.57.

Preparation of Bicarbonate Form of Amberlyst A-26 Resin Using NaOH/CO₂ (10c). A medium sintered glass filter containing 15 g of Amberlyst A-26 resin in the chloride form (average capacity \sim 3.7 mequiv/g) was treated portionwise with 1 L of an aqueous 1.0 M NaOH solution. The resin in the hydroxide form was then dried under aspirator vacuum and suspended in a 150 mL water solution. A flow of CO₂ gas was bubbled into the aqueous solution for approximately 30 min with gentle mixing. The resin was then washed sequentially with MeOH, acetone, and ether, and then extensively dried in vacuo. Elemental analysis indicated essentially no Na ion contamination (0.02%).

Solid-Phase Conditions Used To Prepare 12-Member Pyrazole Library (8a-1). A THF suspension for each of the six dione resins (50 mg, loading capacity 0.92-0.98 mmol/g, prepared according to procedure below), one of the four appropriate hydrazine hydrochlorides (15 equiv), and triethylamine (15 equiv) in toluene (2 mL) were heated to 80 °C for 9 h in a glass cell culture tube with a Teflon-lined screwed cap. Mixing was achieved using a rotating reaction block via a modified rotary evaporator motor. The reaction mixtures were cooled to room temperature and then isolated via vacuum filtration in coarse sintered glass frits. The product resins were rinsed with methanol/DMF 1:1 (2 \times 5 mL), DMF (3×5 mL), CH₂Cl₂ (3×10 mL), and methanol $(2 \times 5 \text{ mL})$. The resulting resins were dried in vacuo for 18 h and then pre-swollen in a 2 mL CH₂Cl₂ solution under N₂ at -78 °C, using 10 mL conical vials, each containing a magnetic stir bar. Pyrazole cleavage/deprotection was followed by addition of 1 M BBr₃ (5 equiv, 0.7 mL) in CH₂-Cl₂. The reaction was allowed to warm to room temperature and stir for 6 h and was quenched at 0 °C by the addition of CH₃OH (2 mL). The resins were isolated by vacuum filtration and rinsed twice with 5 mL portions of CH₃OH. The individual filtrates were concentrated in vacuo, dissolved in MeOH, and concentrated once again. This procedure was repeated twice, prior to RP-HPLC analysis. Pyrazoles 8a-l were then purified to >80% HPLC purity by radial chromatography (10-20% MeOH/CH2Cl2) and submitted for ES-MS and RBA analysis. The expected molecular ion pattern by electrospray was observed in all cases: (8a) ES-MS calcd for $C_{25}H_{24}N_2$ 368.4, found MH^{2+} 370.1 (individual isomers separately measured); (8b) ES-MS calcd for C₂₅H₂₄N₂O₂ 384.5, found MH⁺ 385.2 (individual isomers separately measured); (8c) ES-MS calcd for C₂₅H₂₃FN₂O 386.5, found MH⁺ 387.2; (8d) ES-MS calcd for C₂₃H₂₈N₂O 348.5, found MH⁺ 349.1; (8e) ES-MS cacld for C₂₅H₂₄N₂O₂ 384.5, found MH²⁺ 386.2; (8f) ES-MS calcd for C₂₅H₂₄N₂O₃ 400.5, found MH⁺ 401.1; (8g) ES-MS calcd for C₂₅H₂₃FN₂O₂ 402.5, found MH⁺ 403.1; (8h) ES-MS calcd for $C_{23}H_{28}N_2O_2$ 364.5, found MH⁺ 365.2; (8i) ES-MS calcd for C₂₅H₂₃BrN₂O 446.1, found MH⁺ 447.0 (individual isomers separately measured); (8) ES-MS calcd for $C_{25}H_{23}BrN_2O_2$ 462.1, found MH⁺ 463.1; (8k) ES-MS calcd for C₂₅H₂₂BrFN₂O 464.1, found MH⁺ 465.1; (81) ES-MS calcd for C₂₃H₂₇BrN₂O 426.1, found MH⁺ 427.2.

General Procedure for Batch Dione Synthesis (12a– I). Twelve conical polypropylene cell culture tubes (20 mL) were charged with 400 mg of ketone resin (4 and 11, approximately 0.38 mmol, six tubes per ketone), *p*-nitrophenyl esters (5.76 mmol, 5a-c), and 15 mL of THF. Each tube was then purged with a nitrogen atmosphere using a 14/20 septa and then chilled in an ice bath. A 1.0 M THF solution of LHMDS (3.84 mmol) was added dropwise via syringe to each tube; the reaction was then sealed with a screw cap and shaken vigorously. Esters 5d-f could not be present during enolate formation due to their reactivity toward LHMDS and thus were added in one portion 30 min after addition of LHMDS. All 12 reaction tubes were then loaded into a homemade reaction block, which was slowly rotated within a 40 °C oven using a modified rotary evaporator motor. After 5 h, the resins were transferred to coarse sintered glass filters using CH₂Cl₂. A rinse procedure followed involving methanol/DMF 1:1 (1 \times 30 mL), DMF $(3 \times 20 \text{ mL})$, CH₂Cl₂ $(3 \times 20 \text{ mL})$, and methanol $(2 \times 20 \text{ mL})$ mL). The resins were then dried in vacuo, and the level of product formation was determined by nanoprobe ¹H NMR-MAS. In all 12 cases, disappearance of the ketone α -protons occurred, and a new signal at approximately δ 5-5.3 was observed for the α/α' methine proton, indicating successful dione formation. Diagnostic chemical shifts are reported below (note: aromatic resonances from dione which overlap with polystyrene backbone in addition to the resin backbone signals themselves are not listed; for purposes of naming, polystyrene is abbreviated "PS"; integration values are approximately $\pm 15\%$).

1-PS-(4-Benzyloxyphenyl)-2-ethyl-3-phenyl-propane-1,3-dione (12a): δ 1.02 (3H, CH₃CH₂), 2.15 (2H, CH₃CH₂), 4.95 and 5.05 (two overlapping s, 3H, PS-ArCH₂O and α/α'-CH), 7.95 (4H, ArCH ortho-C=O).

1-PS-(4-Benzyloxyphenyl)-2-ethyl-3-(4-methoxyphenyl)propane-1,3-dione (12b): δ 1.01 (3H, CH₃CH₂), 2.15 (2H, CH₃CH₂), 3.74 (3H, OCH₃), 4.98 (s, 3H, ArCH₂ and α/α'-CH), 7.95 (4H, ArCH ortho-C=O).

1-PS-(4-Benzyloxyphenyl)-2-ethyl-3-(3-methoxyphenyl)propane-1,3-dione (12c): δ 1.02 (3H, CH₃CH₂), 2.14 (2H, CH₃CH₂), 3.72 (3H, OCH₃), 5.05 (two overlapping s, 3H, PS-ArCH₂O and α/α' -CH), 7.51 (2H, ArCH), 7.96 (2H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2-ethyl-3-(4-bromophenyl)propane-1,3-dione (12d): δ 1.01 (3H, CH₃CH₂), 2.13 (2H, CH₃CH₂), 4.97 (s, PS-ArCH₂O and α/α'-CH); 7.50 (ArCH), 7.79 (ArCH), 7.95 (ArCH).

1-PS-(4-Benzyloxyphenyl)-2-ethyl-3-(4-fluorophenyl)propane-1,3-dione (12e): δ 1.01 (3H, CH₃CH₂), 2.14 (2H, CH₃CH₂), 4.97 (s, PS-ArCH₂O and α/α' -CH), 7.97 (4H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2-ethyl-3-(3-fluorophenyl)propane-1,3-dione (12f): δ 1.01 (3H, CH₃CH₂), 2.13 (2H, CH₃CH₂), 4.97 (s, PS-ArCH₂O and α/α' -CH), 7.64, 7.69, 7.95 (three s, 4H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2-*iso*-butyl-3-phenyl-propane-**1,3-dione (12g)**: δ 0.96 (6H, (CH₃)₂), 2.01 (β-CH₂), 4.95 (2H, PS-ArCH₂O), 5.24 (1H, α/α' -CH), 7.39 (3H, ArCH), 7.48 (ArCH), 7.97 (3H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2*iso***-butyl-3-(4-methoxyphenyl)-propane-1,3-dione (12h)**: δ 0.95 (6H, (CH₃)2), 2.02 (β-CH₂), 3.76 (3H, OCH₃), 4.93 (2H, PS-ArCH₂O), 5.17 (1H, α/α' -CH), 7.97 (4H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2-*iso***-butyl-3-(3-methoxyphenyl)-propane-1,3-dione** (**12i**): δ 0.96 (6H, (CH₃)₂), 2.00 (β-CH₂), 3.74 (3H, OCH₃), 4.95 (2H, PS-ArCH₂O), 5.23 (1H, α/α' -CH), 7.29 (ArCH), 7.52 (2H, ArCH), 7.97 (2H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2-*iso*-butyl-3-(4-bromophenyl)-propane-1,3-dione (12j): δ 0.95 (6H, (CH₃)₂), 1.97 and 2.05 (overlapping s, β-CH₂), 4.95 (2H, PS-ArCH₂O), 5.14 (1H, α/α'-CH), 7.51 (2H, ArCH), 7.81 (2H, ArCH), 7.96 (3H, ArCH).

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1-PS-(4-Benzyloxyphenyl)-2-*iso***-butyl-3-(4-fluorophenyl)propane-1,3-dione (12k)**: δ 0.95 (6H, (CH₃)₂), 1.99 and 2.05 (overlapping s, β-CH₂), 4.95 (2H, PS-ArCH₂O), 5.16 (1H, α/α'-CH), 7.98 (4H, ArCH).

1-PS-(4-Benzyloxyphenyl)-2-*iso***-butyl-3-(3-fluorophenyl)propane-1,3-dione (12l)**: δ 0.95 (6H, (CH₃)₂), 1.95 and 2.07 (overlapping s, β-CH₂), 4.97 (2H, PS-ArCH₂O), 5.18 (1H, α/α' -CH), 7.66 (ArCH), 7.73 (ArCH), 7.97 (4H, ArCH).

General Solid-Phase Conditions Used To Prepare 96-Member Pyrazole Library. Pyrazole Formation: Each of the above diones (12a-1) were divided into eight approximately 50 mg portions and distributed across each row of an 8×12 Polyfiltronics Unifilter plate. After charging each well with the appropriate dione, the 12 commercially available hydrazine components (Figure 4) were carefully added as preweighed solids to the appropriate wells while carefully blocking off neighboring wells in order to avoid cross-contamination. The bottom of the plate was then secured in a reaction clamp (Polyfiltronics Combiclamp) with two Viton gaskets. To each well was added 1.5 mL of EtOH and approximately $100 \,\mu\text{L}$ of TEA. The top of the plate was scaled and the assembly agitated and heated to 65 °C in an oscillating cell culture oven. The reaction was monitored by FT-IR using several suspected "worst-case" scenarios (H-2, F-2, and E-5 and our lead compound, A-2) by observing the diminishing intensity of the C=O signal at 1670 cm⁻¹. After 38 h, the plate was cooled to room temperature, and the crude mixtures were filtered within the reaction block by vacuum aspiration (Polyfiltronics filter/collection vacuum manifold). The resin-bound pyrazoles were then rinsed with methanol/DMF 1:1 (2 \times 2 mL), DMF (3 \times 2 mL), CH₂Cl₂ $(3 \times 2 \text{ mL})$, and methanol $(3 \times 2 \text{ mL})$ and then dried in vacuo overnight in a desiccator.

Deprotection/Cleavage: In a N₂-charged glovebag at room temperature, the resins were suspended in 0.5 mL of CH₂Cl₂ and treated dropwise with a 1.0 M CH₂Cl₂ solution of BBr₃ (500 μ L). The reaction block was sealed as before and allowed to stand at room temperature with occasional mixing. After 8 h, the resins were carefully quenched with cold MeOH (0.5 mL), and then approximately 100 mg of bicarbonate resin 10c was added to each well. After 20 min, when most of the CO_2 evolution ceased, the reaction top was resealed and the block heated to 50-60 °C for 1 h. Upon cooling to room temperature, the released library members were collected in a 96-well 2 mL collection plate, and the solvent evaporated to dryness (a 96-pin manifold capable of providing a gentle overhead stream of N₂ to each well in addition to gentle warming using a hot plate allowed for the removal of most of the McOH; the remaining residual volatiles were removed in vacuo). Each member was then reconstituted in 1 mL of McOH and analyzed by RP-HPLC.

Characterization of 96-Member Pyrazole Library. RP-HPLC analysis was performed on an analytical HPLC system with an autosampler. UV detection was performed at 254 nm and a flow rate of 1.0 mL/min using a C-18 CombiScreen analytical column (YMC, Inc., 4.6 mm \times 50 mm) with a McOH:H₂O gradient solvent system (40–90% McOH, 0–5 min, total run time = 10 min). A fluorometer HPLC flow

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detector was used to aid in identification of the expected pyrazole peak. An excitation wavelength of 306 nm and emission wavelength of 364 nm were used, based on the fluorescence properties of pyrazole 1. After RP-HPLC analysis, the library members were transferred from auto sampler vials to tared 7×40 mm flat bottom vials (8 mm crimp tops), and the solvent was then removed using a Savant vacuum centrifuge. After further drying in vacuo the sample weight was recorded, and a modified three-point competitive binding assay was then performed. Authentic HPLC traces obtained for 8a-1 (Table 1) were used to help confirm their later synthesis in the 96-member library. These controls represent 13% of the total library members produced. The RBA values of these "controls" from this library were equivalent to those obtained for the same compounds in the 12-membered library, within the statistical limits of the binding assay (coefficient of variation is 0.3).

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Pyrazole Ligands: Structure–Affinity/Activity Relationships and Estrogen Receptor-α-Selective Agonists

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Articles

Pyrazole Ligands: Structure–Affinity/Activity Relationships and Estrogen Receptor-α-Selective Agonists

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We have found that certain tetrasubstituted pyrazoles are high-affinity ligands for the estrogen receptor (ER) (Fink et al. Chem. Biol. 1999, 6, 205-219) and that one pyrazole is considerably more potent as an agonist on the ER α than on the ER β subtype (Sun et al. *Endocrinology* **1999**, 140, 800–804). To investigate what substituent pattern provides optimal ER binding affinity and the greatest enhancement of potency as an ER α -selective agonist, we prepared a number of tetrasubstituted pyrazole analogues with defined variations at certain substituent positions. Analysis of their binding affinity pattern shows that a C(4)-propyl substituent is optimal and that a p-hydroxyl group on the N(1)-phenyl group also enhances affinity and selectivity for ERa. The best compound in this series, a propylpyrazole triol (PPT, compound 4g), binds to ER α with high affinity (ca. 50% that of estradiol), and it has a 410-fold binding affinity preference for ER α . It also activates gene transcription only through ER α . Thus, this compound represents the first $ER\alpha$ -specific agonist. We investigated the molecular basis for the exceptional ER α binding affinity and potency selectivity of pyrazole 4g by a further study of structure-affinity relationships in this series and by molecular modeling. These investigations suggest that the pyrazole triols prefer to bind to ER α with their C(3)-phenol in the estradiol A-ring binding pocket and that binding selectivity results from differences in the interaction of the pyrazole core and C(4)-propyl group with portions of the receptor where ER α has a smaller residue than ER β . These ER subtype-specific interactions and the ER subtype-selective ligands that can be derived from them should prove useful in defining those biological activities in estrogen target cells that can be selectively activated through ERa.

Introduction

The estrogen receptor (ER) displays a remarkable capacity for binding nonsteroidal ligands with high affinity.¹ Many of these ligands have been developed into hormonal agents having mixed agonist—antagonist and tissue-selective activities that are useful in menopausal hormone replacement, in fertility regulation, and in the prevention and treatment of breast cancer. Because of their unusual pharmacology, some of these agents have been termed selective estrogen receptor modulators (SERMs).² To understand the molecular basis of the tissue selectivity of these SERMs, it would be helpful to know in detail how they are interacting with the ER. However, short of performing X-ray crystallographic analysis of ER complexes with each ligand, it can be a challenge to obtain such information.

If the nonsteroidal ligand bears a reasonable structural relationship with steroidal estrogens, it is gener-

* Address correspondence to: John A. Katzenellenbogen, Dept. of Chemistry, University of Illinois, 600 S. Mathews Ave., Urbana, IL 61801. Tel: 217-333-6310. Fax: 217-333-7325. E-mail: jkatzene@ uiuc.edu. ally quite easy to imagine the orientation that this ligand is likely to adopt when it is bound by ER.^{3,4} ER mutagenesis studies,⁵ and the recent X-ray crystal-lographic structures of ER complexed with both estradiol and three nonsteroidal ligands (raloxifene, hydroxy-tamoxifen, and diethylstilbestrol), provide additional guidance in the selection of reasonable binding orientations for ligands of this type.^{4,6} However, when the nonsteroidal estrogens have structures that are more divergent from those of steroidal estrogens, it becomes a greater challenge to predict ligand-binding orientation.⁷

The recent characterization of a second ER gene, encoding ER β , places a further premium on our understanding of the details of ligand—receptor interaction,^{8,9} because it would be especially interesting to have ligands that could activate or inhibit each of the ER subtypes with high selectivity. Such ligands would be valuable tools to define the biological effects that are mediated by ER α and ER β . So far, however, there have been only a few reports of ER subtype-selective estrogens, and in many cases, the selectivity has been relatively modest.¹⁰

Recently, we investigated various heterocyclic diazole structures as core elements for nonsteroidal estrogens

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Figure 1. Pyrazole core optimized for high-affinity ER binding (X = H, OH; R = alkyl).

of novel design.^{11,12} Our aim was to identify systems that would be amenable to combinatorial assembly and library synthesis. From among the several systems we studied that included imidazoles, oxazoles, thiazoles, and isoxazoles, we found that high-affinity ER ligands could be obtained by appropriate substitution on a pyrazole core. Subsequently, we used a parallel, solidphase synthesis approach to prepare some combinatorial pyrazole libraries of moderate size.¹³ In the cases we investigated, high-affinity binding required tetrasubstitution of the pyrazole core and an appropriate display of aromatic, phenolic, and aliphatic groups.^{11,12} An example of this optimal pattern of substitution that we have thus far delineated for pyrazoles includes three aromatic groups at the 1-, 3-, and 5-positions, specifically phenols at the 3- and 5-positions, and an alkyl group at the 4-position (Figure 1). Initial studies on one of these compounds (Figure 1; X = H, R = Et) showed that it acted as an agonist on both ER subtypes, but it was considerably more potent on ER α than on ER β .¹⁴ Thus, this pyrazole was termed an ERa potencyselective agonist.¹⁴

In this report, we have investigated structure– activity relationships in these tetrasubstituted pyrazoles to delineate two aspects of their behavior: (1) the C(4)alkyl substituent and phenol hydroxyl pattern that provide optimal ER subtype selectivity and (2) the extent to which the ER α -selective binding affinity and potency of these pyrazoles can be understood in the context of crystal structures of the ER α ligand-binding domain (LBD) and models for ER β derived from these structures. Our studies provide new information on these issues, and in the process, we have identified a pyrazole that shows complete ER α selectivity in transcription activation by the receptor.

Results and Discussion

Chemical Syntheses. The compounds we have studied have various C(4)-alkyl substituents and have either a phenyl or 4-hydroxylphenyl substituent on N(1). Shown in Scheme 1 is the route used for the synthesis of the pyrazoles 4a-i. The starting alkylphenones 1a-f were readily prepared in good yields by Friedel-Crafts acylation of anisole. The requisite β -diketones $2\mathbf{a}-\mathbf{e}$ were then produced in moderate to good yields by acylation of the corresponding lithium enolates with 4-nitrophenyl 4-methoxybenzoate. Because the pyrazoles are sterically crowded, rather harsh conditions were required for their formation (>16 h at reflux at 110-120 °C in DMF/THF solution). Nonetheless, these conditions served quite well for pyrazole formation in solution. Several of the pyrazole intermediates protected as methyl ethers were isolated in good yield. However, with the trimethoxy pyrazoles 3g-i, it was difficult to

Scheme 1. Synthesis of C(4)-Alkylpyrazole Analogues



separate the protected product from traces of the diketone precursor by chromatography. So, in these cases the crude material was passed through a short silica gel column and, without further purification, was then deprotected using BBr₃. Pyrazoles **3d**,**e** were also treated in this manner. The final phenolic pyrazoles **4a**-i were isolated in 30–98% yield after purification by recrystallization and/or chromatography.

In addition to the final pyrazole products 4a-i, we wished to prepare the isopropyl analogue derived from ketone 1f. However, we encountered difficulties in the preparation of this hindered dione because of significant O-acylation that occurred during the Claisen condensation. Separation of the O-acylated byproduct from the desired dione proved to be difficult. Moreover, once the dione was isolated, the pyrazole condensation failed using the conditions described above, which had been optimized for solution-phase synthesis. To avoid these problems, we utilized a solid-phase synthesis according to the methodology that we had previously developed,¹³ and in this manner we were able to obtain the isopropylpyrazole 6 starting from the resin-bound ketone 5, as shown in Scheme 2, although the overall yield was relatively low.

To prepare a complete series of all of the possible diphenol and monophenol analogues of the C(4)-ethyl triphenol **4f**, we also synthesized monophenols $7\mathbf{a}-\mathbf{c}$ and the remaining diphenols **8a,b** (all mono- and diphenols are shown in Scheme 3, top). Monophenol **7a** was prepared from dibenzoylmethane (**9**) and 4-methoxyphenylhydrazine by the usual sequence¹¹ (Scheme 3, middle). The synthesis of the remaining two monophenols (**7b,c**) by a regioselective route has been described elsewhere (not shown).¹⁵ The two remaining

Scheme 2. Synthesis of C(4)-Isopropylpyrazole 6 by Solid-Phase Methodology¹³



6 (11%, three steps from ketone)

diphenols (**8a**,**b**) were also prepared by this regioselective method, as shown in Scheme 3 (bottom). The two chalcones (**12a**,**b**) are treated with *p*-methoxyphenylhydrazine under anerobic conditions to give the pyrazolines **13a**,**b**. The corresponding anions, generated by treatment with LDA, were ethylated, and the C(4)ethylpyrazolines **14a**,**b** were then oxidized to the pyrazoles **15a**,**b** and deprotected to the desired bisphenols **8a**,**b**.

ER Binding Affinity of Tetrasubstituted Pyrazoles. The ER binding affinity of pyrazoles $4\mathbf{a}-\mathbf{i}$, 6, $7\mathbf{a}-\mathbf{c}$, and $8\mathbf{a}$, **b** was determined in a competitive radiometric binding assay using purified full-length human ER α and ER β , as previously described.^{16,17} The affinities are expressed as relative binding affinity (RBA) values and are presented in two tables: Table 1, which is discussed here, covers the effect of the nature of the C(4)-alkyl substituent on the binding affinity and ER subtype selectivity of pyrazoles in the triphenol and the principal 3,5-diphenol series; the data in Table 2, which compares the affinity of the three possible mono- and diphenols with the triphenol in the C(4)-ethylpyrazole series, relates to the issue of ligand orientation and is discussed in a later section.

Two interesting trends are notable in the binding affinity data presented in Table 1. In both the diphenol series (X = H, 4a-e, 6) and the triphenol series (X = OH, 4f-i), optimal binding affinity for ER α requires a C(4)-alkyl substituent that is not too long (*n*-Bu: 4e,i) or too short (Me: 4a), the highest affinity being found with the intermediate size substituents: Et (4b,f), *n*-Pr (4c,g), and *i*-Bu (4d,h). Thus, it appears that the subpocket that is accommodating this group has a limited size and relatively narrow shape. A more significant trend evident in this series has to do with the effect of the phenolic substituent on the N(1)-phenyl group. In each case where a comparison can be made (namely, Et, Pr, *i*-Bu, or *n*-Bu), the triphenols (X = OH, **4f**-**i**) have higher affinity on ER α and/or lower affinity on ER β than do the diphenols (X = H, **4a**-**e**). As a result, the triphenols consistently have higher ER α affinity selectivity than do the corresponding 3,5-diphenols.¹⁸ In fact, some of the members show a remarkable selectivity in their binding affinity for ER α vs ER β , which is 30-40 for some of the pyrazole diols (**4c**,**d**) and as high as 200-400 for some of the pyrazole triols (**4f**,**g**).

The increased ER α binding affinity of the triphenols is not expected, because additional polar substituents are generally poorly tolerated in the center of the ligandbinding pocket of ER, at least in most nonsteroidal ligand systems that have been examined, such as the benzo[b]thiophenes¹⁹ and 2,3-diarylindans.⁷ One exception is found with certain triphenylacrylonitriles, where addition of a third hydroxyl increases binding affinity.²⁰

We have previously reported binding affinities for one of these pyrazoles (**4b**), using receptor preparations containing only the *LBDs* of human ER α and ER β , rather than *full-length* human ER α and ER β .¹⁴ The values obtained previously were higher but less ER α selective (60 ± 16 for ER α and 18 ± 4 for ER β). At this point, the reasons for these affinity differences between full-length ER and the ER LBD are not clear, although they have been seen, as well, in a structurally different class of ER ligands.¹⁷ However, it is of note that the higher ER α /ER β affinity ratio that we obtain for pyrazole **4b** with the full-length ERs is more consistent with the 120-fold potency ratio we described in transcription assays than was the 3-fold ER α /ER β affinity ratio obtained with ERs containing only the LBDs.¹⁴

Transcriptional Activity and Potency of Tetra-

Scheme 3. Synthesis of the Remaining Mono- and Diphenolic C(4)-Ethylpyrazoles



substituted Pyrazoles. For investigation of transcription activation ability, we selected two compounds: the pyrazole having the highest ER α /ER β affinity selectivity (410-fold, the propylpyrazole triol 4g, called PPT for convenience) and the corresponding pyrazole in the diol series (4c, called propylpyrazole diol or PPD). These two pyrazoles were assayed for estrogen agonist activity in transactivation assays using human endometrial cancer (HEC-1) cells transfected with expression plasmids for ER α and ER β and an estrogen-responsive reporter gene plasmid. The dose-response curves for these compounds are shown in Figure 2.

Both compounds are potent in activating gene transcription through ER α , but the PPD (4c) is weak in transcriptional activation through ER β and PPT (4g) is completely inactive in stimulating transcription via ER β . Thus, PPT (4g) is an ER α -specific agonist. This pharmacological profile is reminiscent of the behavior of the C(4)-ethyl analogue of pyrazole 4c (namely, pyrazole 4b), that we have previously described as an agonist that is more potent on ER α than on ER β . This compound (**4b**) had an EC₅₀ of ca. 1 nM on ER α and showed a 120-fold potency selectivity for this ER subtype.¹⁴ However, in cell transfection assays both PPT and PPD are nearly 10-fold more potent than the original pyrazole **4b** on ER α , and they are much more ER α -selective.

As we had noted in our earlier study on pyrazole 4b,¹⁴ and is clearly evident here as well, the ER α selectivity of these pyrazoles in terms of their *potency* in transcription assays is substantially greater than their *affinity* selectivity in binding assays. For example, the ER α affinity selectivity of PPD (4c) is 32, yet its potency selectivity is ca. 1000 (cf. Figure 2). Likewise, the ER α affinity selectivity of PPT (4g) is 410, and its potency selectivity, which although is difficult to accurately evaluate (cf. Figure 2), is probably greater than 10000. Such a discordance between affinity and potency might be explained in the context of "tripartite receptor pharmacology", a concept that we advanced some time

 Table 1. Relative Binding Affinity Data for C(4)-Alkyl

 Triphenol and 3,5-Diphenol Pyrazole Analogues^a



					ERα/ERβ				
compd	R	Х	RBA ER α^b	RBA ER β^b	RBA ratio				
3,5-Diphenols									
4a	Me	н	0.76 ± 0.18	0.28 ± 0.16	2.7				
4b	\mathbf{Et}	Н	31 ± 15	1.1 ± 0.2	28				
6	i-Pr	Н	5.6 ± 2	0.86 ± 0.11	6.5				
4c (PPD)	n-Pr	Н	16.8 ± 0.3	0.52 ± 0.03	32				
4d	<i>i-</i> Bu	Η	56 ± 6	1.4 ± 0	40				
4e	n-Bu	Η	8.7 ± 2.0	0.47 ± 0.1	19				
1,3,5-Triphenols									
4f	Et	OH	36 ± 6	0.15 ± 0.014	240				
4g (PPT)	n-Pr	OH	49 ± 12	0.12 ± 0.04	410				
4h	<i>i-</i> Bu	OH	75 ± 6	0.89 ± 0.06	84				
4i	n-Bu	OH	14 ± 4	0.18 ± 0.09	77				

^{*a*} Relative binding affinity (RBA), where estradiol is 100%. Values are the mean of at least 2 and more typically 3 or more independent determinations (±SD).^{33 b} Competitive radiometric binding assays were done with purified full-length human ER α and ER β (PanVera Inc.), using 10 nM [³H]E₂ as tracer and HAP for adsorption of the receptor-tracer complex.^{16,17}

ago.²¹ Binding measured in vitro with purified ERs involves only the interaction between ligand and receptor, whereas transcription measured in cells involves the additional interaction of the ligand-receptor complex with coactivators and other cellular components. Thus, compared to their relative affinities for ER α and ER β , the relative potency of two ligands can be modulated by differences in the strength with which their respective ligand-receptor complexes bind to coactivators or are modified by other cellular elements, although other factors, as well, might be involved. In this case, it appears that the pyrazole complexes with ER α are better able to bind coactivators than are the ER β pyrazole complexes, as we have documented in a recent study.²²

A Model for the Binding of Pyrazoles with the ER Subtypes. The high affinity, and particularly the high selectivity, with which these pyrazoles bind to ER α raises the important issue of what molecular features underlie the differences in their interaction with ER α vs ER β . Without crystal structures available for the comparison of any of these pyrazoles complexed with both ER α and ER β , we are currently limited to investigating this issue by molecular modeling.

1. Structure-Affinity Relationships and the Orientation of Pyrazoles in the Ligand-Binding Pocket of ER α . Because the pyrazoles we have studied here are rather symmetrical and are polyphenolic, it is not obvious which orientation these ligands might prefer to adopt within the ligand-binding pocket of ER. In principle, the triphenolic pyrazoles could adopt six orientations (see Figure 4): Each of the three phenolic rings could play the role of the A-ring of estradiol, and in each case the remainder of the pyrazole could adopt two orientations about the bond connecting the A-ring phenol mimic to the pyrazole core (darkened bond).

These six binding modes are given the designations: N(1), N(1)', C(3), C(3)', C(5), and C(5)'.

A classical approach that has been used to try to ascertain which orientation di- and triphenolic compounds adopt within the ER binding pocket has been to systematically delete the phenolic hydroxyl group and examine the effect of this deletion on the binding affinity.⁷ Because the hydroxy group on the phenol that corresponds to the A-ring of estradiol is thought to make the major contribution to ligand-binding affinity, the greatest reduction in binding should occur when this hydroxyl group is the one deleted.³ We have undertaken such an analysis by comparing the affinities of pyrazoles in which the three phenols in the triphenol **4f** were each singly deleted (diphenol set: 4b, 8a,b) or were deleted in three pairs (monophenol set: 7a-c). The affinities of these three diphenols and three monophenols, together with the triphenol parent 4f, are listed in Table 2. Because we had prepared the most pyrazoles in the C(4)-ethyl series, we chose to do this structure-affinity study in this series.

The simplest analysis we can make of the structurebinding affinity pattern shown in Table 2 is the following: In the monophenol set (7a-c), the N(1)- and C(3)phenols (7a, b) have comparable affinities, but the C(5)phenol (7c) binds ca. 100-fold less well. Because of the very low affinity of the C(5)-monophenol (7c), we believe that the C(5)-phenol cannot function as the mimic of the A-ring of estradiol, in essence, eliminating orientations C(5) and C(5)' as possibilities. (It is of note that these two "eliminated" binding modes project bulky groups simultaneously in directions that correspond to C(7) and C(11) of estradiol; cf. Figure 3.) Elimination of the C(5)-phenol leaves the N(1)- and C(3)-phenols as potential estradiol A-ring mimics, but the very similar affinities of both of these monophenols suggest that each one may function as the A-ring mimic.

Analysis of the binding affinity pattern of the bisphenols is somewhat more ambiguous, but suggestive, nevertheless: Deletion of the N(1)-phenol from the triphenol has only a minor effect on the binding affinity (4b vs triphenol 4f), which means that in the context of the triphenol in ER α , the N(1)-phenol is not contributing significantly to binding affinity. Thus, one would expect that the C(3)-phenol is likely to be the more important of the two and thus that the orientations C(3) and C(3)'are more likely than the orientations N(1) and N(1)'. Deletion of either the C(3)-phenol or the C(5)-phenol results in a 4–5-fold decrease in binding (8a,b vs 4f). The fact that the N(1),C(3)-bisphenol (8b) still binds quite well is not surprising, because the C(3)-phenol is still available to be the estradiol A-ring mimic. However, the fact that the N(1),C(5)-phenol (8a) also binds quite well might seem surprising, because this diphenol lacks the important C(3)-hydroxyl.

We believe that the reasonably good affinity of diphenol **8a** suggests that this pyrazole can bind with the N(1) group as the estradiol A-ring mimic, meaning that the N(1) and N(1)' ligand orientations are also possible, though, based on the minimal effect of the N(1)-phenol deletion, they are probably not significantly populated when a C(3)-phenol is available. (It seems unlikely that diphenol **8a** would bind with the C(5)-phenol in the A-ring pocket, because of the very low affinity of the



Figure 2. Transcription activation by ER α (left) and ER β (right) in response to pyrazoles **4c** (PPD) and **4g** (PPT). Human endometrial cancer (HEC-1) cells were transfected with expression vectors for ER α or ER β and an (ERE)3-pS2-CAT reporter gene and were treated with the indicated concentrations of ligand for 24 h. CAT activity was normalized for β -galactosidase activity from an internal control plasmid. Values are expressed as a percent of the ER α or ER β response with 1 nM E₂, which is set at 100%.³²



Figure 3. Potential pyrazole binding relative to estradiol. Both modes A and A' have the C(3)-phenol oriented to project in to the estradiol A-ring binding pocket. Modes A and A' are related to one another by rotation of the pyrazole core around the darkened bond.

C(5)-monophenol **7c**.) Thus, the phenol deletion/binding affinity approach has only reduced the number of possible ligand orientations in the binding pocket from six to four, with a suggested preference for two orientations, C(3) and C(3)'. As described in the section below, we have tried to narrow down the possible ligand orientations even further by molecular modeling.

The conclusions thus far that the C(3)-phenol is the preferred A-ring mimic but that the N(1)-phenol is a possible alternative are consistent with two other studies that we have recently completed. In a related but more limited investigation of the orientation of ligand binding in two isomeric pyrazole series, we concluded that the orientation with the C(3)-phenol as the A-ring mimic was most likely in the pyrazole series that is the same as the one studied here and that in the isomeric series the phenol that occupied topologically congruent substituent position also functioned as the A-ring mimic.²³ In another investigation in which we worked

to convert these pyrazoles from agonists to antagonists by appending a basic side chain onto the four substituents on these pyrazoles, we found that high affinity was retained only when this group was attached to the C(5)-hydroxyl.²⁴ On the basis of this structure–affinity pattern, as well as some molecular modeling and consideration of the structure of the ER LBD complexes with antagonists, we concluded in that study that the most likely orientation of these basic side chainsubstituted pyrazoles was N(1).²⁴ In this orientation, the basic side chain could project outward through the 11 β channel that forms in the ER LBD complexes with antagonists.^{4,6}

2. Molecular Modeling of the Orientation of Pyrazole Triols in the Ligand-Binding Pocket of the ERs. We have previously done modeling of the pyrazole ligands in the ER LBDs, first in our initial study on these ligands¹¹ and, more recently, in connection with a study of the orientation of these ligands in



Figure 4. Crossed-stereoview of propylpyrazole triol (PPT, **4g**) docked and minimized in ER α LBD pocket according to binding mode A, showing selected residues close to the ligand. The pyrazole ligand is shown with standard atom colors. The residues in ER α are identified and shown in yellow. At the two positions where the ER β sequence differs from ER α , the ER β residues are shown in red (Met in ER β in place of ER α Leu384 and Ile in ER β in place of ER α Met421).

Table 2. Relative Binding Affinity Data for C(4)-Ethylpyrazole Tri-, Di-, and Monophenols^{*a*}



^{*a*} Relative binding affinity (RBA), where estradiol is 100%. Values are the mean of at least 2 and more typically 3 or more independent determinations (±SD).^{33 b} Competitive radiometric binding assays were done with purified full-length human ER α and ER β (PanVera Inc.), using 10 nM [³H]E₂ as tracer and HAP for adsorption of the receptor-tracer complex.^{16,17}

the binding pocket.²³ As a result of this latter study,²³ we have come to appreciate that there are some ambiguities in how polyphenolic ligands such as these pyrazoles might be accommodated by ER, at least at the level at which we are currently able to model. Such ambiguity is echoed in the results of the structure– binding affinity considerations discussed above. Therefore, for the purposes of this paper, we have chosen to illustrate a binding mode for these pyrazoles in which the C(3)-phenyl group is placed in the ER LBD pocket that normally binds the A-ring of estradiol (Figure 3, see C(3) and C(3)'), as this appeared to be the most likely orientation for these polyphenolic systems (see above and ref 23). We have, however, also examined the other four possible orientations.

We conducted a ligand docking/binding minimization routine by placing PPT (4g) in ER α in both orientations: mode C(3) and mode C(3)' (Flexidock routine within Sybyl; see Experimental Section, Molecular

Modeling). During the course of the ligand docking routine, the PPT that began in orientation mode C(3)'became reoriented into mode C(3), whereas the one that began in mode C(3) remained in mode C(3). Therefore, mode C(3)' was not considered further. After the docking/minimization routine, the PPT in mode C(3) was nicely accommodated in this orientation, with only minimal changes in the side chain conformations of some of the residues that line the ligand-binding pocket. We also modeled the four other possible orientations of this pyrazole using the same routine (not shown), but we found that none of them gave a fit as good as that of the pyrazole orientation illustrated as mode C(3) in Figure 3. The fact that the N(1) and N(1)' modes did not score well in our modeling, although they seemed possible on the basis of the structure-affinity considerations above, suggests either limitations in our modeling method or the fact that these modes are not important in the triphenolic pyrazoles and only become important when certain phenols are missing (see above) or a basic side chain is appended.²⁴

In Figure 4, we show a crossed-stereoview skeletal model illustrating PPT (4g) in the ER α structure, oriented in binding mode C(3), after the ligand docking/ minimization routine. For clarity and for the ER α and $ER\beta$ comparison below, only selected residues from ER are shown. In this orientation, the N(1)-phenyl group of PPT (4g) projects into a region of the binding pocket which represents the C/D-subpocket that is normally occupied by portions of the C- and D-rings and the 18methyl substituent on estradiol (E_2) . Despite the importance of the N(1)-phenol in enhancing ER α binding selectivity (see above and Table 1), there are no obvious sites for its interaction in this subpocket. The N(1)- and C(5)-aryl groups of PPT project somewhat more deeply into the C/D-subpocket than do the corresponding regions of estradiol. However, this region is known to tolerate a variety of substituents,³ and certain residues have been repositioned by the docking/minimization routine to make additional space for the pyrazole ligand. It is of note that these residue conformational changes, which allow for PPT binding in this mode, involve relatively minimal alteration in the total protein energy.

3. Analysis of Interactions Between Pyrazole Substituents and Amino Acid Residues in ER α and ER β . We have used the C(3) orientation mode



Figure 5. Crossed-stereoview of ligand skeletal structures and receptor molecular surfaces for the ER α - and ER β -propylpyrazole triol (PPT, **4g**) complexes. The PPT in ER α is shown with standard atom colors, and the ER α receptor surface is shown in yellow; in the ER β -PPT complex, both the ligand and receptor surface are shown in red. For further details, see text.

model derived above to try to understand the origins of the high ER α affinity selectivity of PPT (4g), and to do this, we have constructed a model for $ER\beta$ that is derived from ERa LBD-diethylstilbestrol (DES) structure (see Experimental Section for details).⁶ Although the LBDs of ER α and ER β have only 56% amino acid identity, the residues that line the ligand-binding pocket are nearly identical.²⁵ In fact, of the 22-24 residues that are considered to be in contact with the ligand (i.e., within 4 Å), all but two are identical in ER α and ER β .²⁵ The only differences are at ER α position 384, where ER α has a leucine but ER β has a methionine, and at ER α position 421, where ER α has a methionine but ER β has an isoleucine. It has been speculated that these sequence differences, in particular, may underlie the differences in the ER α and ER β binding affinities of ligands that have been studied so far.²⁵ For clarity, we have distinguished these residues in Figure 4 by color: The residues common to ER α and ER β are illustrated in yellow, but at the two positions where the $ER\beta$ residues are different, the ER β residues, colored in red, are superimposed over the ERa residues in yellow (see Figure 4 and legend).

In this model with the most ER α potency-selective ligand PPT (**4g**), we can see that the pyrazole core itself has the closest contact with Leu384 in the case of ER α and Met384 in the case of ER β . For the Met/Ile discriminating residues at ER α position 421, the C(4)-propyl group as well as the C(5)-aryl group appear to have close contacts, although this interaction seems less severe.

A model for PPT (4g) binding to ER β was generated by changing the two residues in the ER α ligand-binding pocket to those found in ER β , inserting the ligand according to mode C(3), and then conducting the ligand docking/minimization routine as was done before with ER α (see Experimental Section). The comparative fit of this pyrazole in the ligand-binding pocket of ER α and ER β can be appreciated by the composite receptor surface model shown in Figure 5 in crossed-stereoview. In this figure, we have displayed the pyrazoles in both of the receptors as skeletal structures, and the two pockets as colored surfaces. The ligand in the ER α model is shown with standard atom colors and the ER α surface in yellow; both the ligand and receptor in the ER β model are shown in red. In this figure, the view has been rotated, twisting the C(3)-phenol away from the viewer; this orientation is the best we have found to display differences between the two models.

The principal difference in the surface of the ligandbinding pocket is at the site where ERa has a smaller residue (Leu384) than $\text{ER}\beta$ (Met384). This is illustrated by the sharp red interpenetrating surface in the middle left region of the pocket. The increased steric bulk in $ER\beta$ at this site makes contact with the pyrazole core of this ligand, displacing it relative to the pyrazole position in the ER α model. Because the C(3)-phenol remains relatively fixed in the rather rigid A-ring binding pocket, this pyrazole core displacement causes a large shift in the position of the N(1)-phenol and a significant but somewhat smaller shift of the C(5)phenol and the propyl group. The large shift of the N(1)phenol is intriguing, because this group contributes significantly to the ER α affinity and potency selectivity of these pyrazoles. However, we have been unable to identify any interaction in the N(1)-phenol subpocket that would favor the presence of a hydroxyl in ER α and disfavor it in $ER\beta$.

The shift of the propyl group forces this substituent to adopt a gauche conformation in ER β (vs an anti orientation in ER α); this is a change that would most likely increase ligand internal energy and reduce binding affinity. The shift of the pyrazole position in the ER β model moves the C(5)-phenol toward the site where the smaller Ile residue in ER β replaces the bulkier Met421 in ER α . Thus, the different shape of the ligand-binding pocket in ER β , in particular the increased bulk of the ER β Met residue where ER α has a Leu, appears to interfere with the optimal binding of the pyrazole ligand and is therefore thought to account for its lowered binding affinity. It is interesting to note that even after extensive minimization, there is a substantial energy difference between the two complexes, with the ERa complex having greater stabilization by ca. 30 kcal/mol.

Conclusions

A series of C(4)-alkyl tetrasubstituted pyrazole analogues have been prepared and their RBA values determined for the ER. These compounds show an interesting binding affinity pattern, including highaffinity selectivity for ER α , which for the propylpyrazole triol (PPT, 4g) reaches 410-fold. This compound also has very high potency selectivity for ERa in reporter gene transcription assays in cells, and it is the first $ER\alpha$ specific agonist. Structure-affinity relationships in a series of mono-, di-, and triphenolic pyrazoles and molecular modeling suggests that these pyrazoles bind to the ER with the C(3)-phenol in the A-ring binding pocket. A model of PPT (4g) docked in the ER α LBD-DES crystal structure in the most likely binding orientation suggests that the very high $ER\alpha$ binding selectivity of this pyrazole derives from particular interactions between the pyrazole core and the C(4)-propyl group with a region on the ligand-binding pocket where $ER\alpha$ has a smaller residue (Leu384) than ER β (Met384). These interactions may serve as the basis for future structure-based design of ER α - and ER β -specific ligands.

Experimental Methods

General. All reagents and solvents were obtained from Aldrich, Fisher, or Mallinckrodt. Tetrahydrofuran was freshly distilled from sodium/benzophenone. Dimethylformamide was vacuum distilled prior to use and stored over 4 Å molecular sieves. Melting points were determined on a Thomas-Hoover UniMelt capillary apparatus and are uncorrected. All reactions were performed under a dry N₂ atmosphere unless otherwise specified. Reaction progress was monitored by analytical thinlayer chromatography using GF silica plates purchased from Analtech. Visualization was achieved by short-wave UV light (254 nm) or potassium permanganate stain. Radial preparative-layer chromatography was performed on a Chromatotron instrument (Harrison Research, Inc., Palo Alto, CA) using EM Science silica gel Kieselgel 60 PF₂₅₄ as adsorbent. Flash column chromatography was performed using Woelm 32-63 µm silica gel packing.²⁶ Synthetic procedures for 1a-f and spectral data for 1d,f are described below. Compounds 1a,27 1b,28 1c,29 and $1e^{30}$ were spectroscopically identical to the reported compounds. The synthesis of compounds 7b,c has been described elsewhere.15

¹H and ¹³C NMR spectra were recorded on either a Varian Unity 400 or 500 MHz spectrometers using CDCl₃ or MeOD d_4 as solvent. Chemical shifts are reported as parts per million downfield from an internal tetramethylsilane standard (δ 0.0 for ¹H) or from solvent references. NMR coupling constants are reported in hertz (Hz). ¹³C NMR were determined using either the attached proton test (APT) or standard ¹³C pulse sequence parameters. Low- and high-resolution electron impact mass spectra were obtained on Finnigan MAT CH-5 or 70-VSE spectrometers. Low- and high-resolution fast atom bombardment (FAB) were obtained on a VG ZAB-SE spectrometer. Elemental analyses were performed by the Microanalytical Service Laboratory of the University of Illinois. Those final components that did not give satisfactory combustion analysis (i.e., 4b-d, 6, 8a) gave satisfactory exact mass determinations and were found to be at least 98% pure by HPLC analysis under normal and reversed-phase conditions.

RBA Assays. Purified ER α and ER β binding affinities were determined using a competitive radiometric binding assay using 10 nM [³H]estradiol as tracer, commercially available ER α and ER β preparations (PanVera Inc., Madison, WI), and

hydroxylapatite (HAP) to adsorb bound receptor–ligand complex.^{16,17} HAP was prepared following the recommendations of Williams and Gorski.³¹ All incubations were done at 0 °C for 18–24 h. Binding affinities are expressed relative to estradiol (RBA = 100%) and are reproducible with a coefficient of variation of 0.3.

Transcription Activation Assays. Human endometrial cancer (HEC-1) cells were maintained in culture and transfected as described previously.³² Transfection of HEC-1 cells in 60-mm dishes used 0.4 mL of a calcium phosphate precipitate containing 2.5 μ g of pCMV β Gal as internal control, 2 μ g of the reporter gene plasmid, 100 ng of the ER expression vector, and carrier DNA to a total of 5 μ g DNA. CAT activity, normalized for the internal control β -galactosidase activity, was assayed as previously described.³²

Molecular Modeling for Pyrazole 4g. The starting conformation for **4g** used for receptor docking studies was generated from a random conformational search performed using the MMFF94 force field as implemented in Sybyl 6.6. The resulting lowest-energy conformer was then used for docking studies. Charge calculations were determined using the MMFF94 method and molecular surface properties displayed using MOLCAD module in Sybyl 6.6.

Pyrazole 4g, generated as noted above, was prepositioned in the DES-ERa LBD crystal structure (Protein Data Bank code $(3ERD)^6$ using a least-squares multifitting of select atoms within the DES ligand. Once prepositioned, DES was deleted and ligand 4g optimally docked in the ERa binding pocket using the Flexidock routine within Sybyl (Tripos). Both hydrogen-bond donors and acceptors within the pocket surrounding the ligand (Glu353, Arg394, and His524), the ligand itself, and select torsional bonds were defined. The best docked receptor-ligand complex from Flexidock then underwent a three-step minimization: first nonring torsional bonds of the ligand were minimized in the context of the receptor using the torsmin command, followed by minimization of the side chain residues within 8 Å of the ligand while holding the backbone and residues Glu353 and Arg394 fixed. A final third minimization of both the ligand and receptor was conducted using the Anneal function (hot radius 8 Å, interesting radius 16 Å from pyrazole 4g) to afford the final model.

The ER β molecular model was generated within Sybyl by first modifying residues Leu384 to Met384 and Met421 to Ile421 in the DES–ER α LBD crystal structure⁶ and conducting an initial minimization of these two residues while holding the remaining atoms fixed. This was followed by a final minimization using the Anneal function (hot radius 4 Å, interesting radius 8 Å from Met384 and Ile421) using conditions similar to above. The pyrazole 4g was introduced into this ER β model and the Flexidock routine implemented as for the ER α model (see above). All minimizations were done using the MMFF94 force field with the Powell gradient method (final rms < 0.02 kcal/mol·Å).

Chemical Syntheses. General Procedure for Preparation of Alkylphenones Using AlCl₃ (1a-c,f): Method A. To a stirred solution of AlCl₃ (46.8 mmol) in 1,2-dichloroethane (10 mL) at 0 °C was added the commercial acid chloride (39.0 mmol) dropwise over 10 min. The resulting solution was allowed to come to room temperature for 20 min until all of the AlCl₃ had dissolved. The reaction mixture was cooled to 0 °C and a solution of anisole (5.1 mL, 46.8 mmol) in 1,2dichloroethane (20 mL) was then added dropwise over 30 min. Upon completion, the reaction was allowed to reach room temperature and stir for 8-15 h. The mixture was then quenched by pouring over 100 g of ice and extracted with CH_2Cl_2 (3 × 25 mL). The combined organic layers were washed with water, NaHCO₃ (satd), brine, then dried over MgSO₄ and concentrated under reduced pressure. Excess anisole was removed in vacuo and then the ketone product distilled.

General Procedure for Preparation of Alkylphenones Using PPA (1d,e): Method B. A mechanically stirred mixture consisting of carboxylic acid (1.0 equiv), anisole (1.1 equiv) and polyphosphoric acid (PPA; 6.15 g/mL of anisole) was heated to 90–100 °C for 1.5 h. Upon cooling to near room temperature, the dark mixture was poured over ice (100 g/0.1 mol) and extracted repeatedly with EtOAc. The organic layers were then washed with satd NaHCO₃ followed by brine. After drying the extract over Na₂SO₄ and removal of solvent, a crude oil was obtained. The final products were purified by bulb-to-bulb distillation.

1-(4-Methoxyphenyl)-4-methylpentan-1-one (1d). Prepared according to method B outlined above to afford the title compound as a pale yellow oil (75%): ¹H NMR (CDCl₃, 500 MHz) δ 0.94 (d, 6H, *J* = 6.0), 1.60 (m, 3H, overlapping methine and β-CH₂), 2.90 (t, 2H, *J* = 6.8), 3.87 (s, 3H), 6.92 (d, 2H, *J* = 8.8), 7.90 (d, 2H, *J* = 8.8); ¹³C NMR (CDCl₃, 125 MHz) δ 22.6, 28.5, 33.9, 36.4, 55.2, 113.5, 130.2, 163.6, 198.7; MS (EI, 70 eV) *m/z* 206.1 (M⁴). Anal. (C₁₃H₁₈O₂) C, H.

1-(4-Methoxyphenyl)-3-methylbutan-1-one (1f). Prepared according to general method A outlined above to afford the title compound as a pale yellow oil (84%): ¹H NMR (CDCl₃, 500 MHz) ∂ 0.95 (d, 6H, J = 6.7), 2.24 (sept, 1H, J = 6.6), 2.73 (d, 2H, J = 6.7), 3.81 (s, 3H), 6.88 (d, 2H, J = 8.9), 7.90 (d, 2H, J = 9.2); ¹³C NMR (CDCl₃, 125 MHz) ∂ 22.6, 25.2, 46.2, 55.2, 113.5, 130.2, 130.3, 163.1, 198.7; HRMS (EI, M⁺) calcd for C₁₂H₁₆O₂ 192.1150, found 192.1153.

2-Methyl-1,3-bis(4-methoxyphenyl)propane-1,3-dione (2a). To a solution of 1a (160 mg, 0.97 mmol) and 4-nitrophenyl 4-methoxybenzoate (prepared from p-nitrophenol and 4-methoxybenzoic acid using diisopropylcarbodiimide and 4-(dimethyamino)pyridine in THF (35 mL) at 0 °C was added a 1.0 M solution of lithium hexamethyldisilamide (3.03 mL, 3.03 mmol) dropwise over 5 min. The reaction was warmed to room temperature and allowed to stir for 1.5 h. At this time the reaction was quenched by the addition of H₂O (25 mL). The mixture was then repeatedly extracted with diethyl ether. The organic layers were combined and washed with H₂O, then dried over Na₂SO₄ and concentrated under reduced pressure to afford a crude solid. Unreacted ester was removed by adding a solution of 40% ethyl acetate/hexanes and filtering off the insoluble ester. The remaining filtrate was concentrated and subjected to flash chromatography (40% ethyl acetate/hexanes) to afford the title compound as a light yellow oil (266 mg, 92%): ¹H NMR (CDCl₃, 500 MHz) δ 1.58 (d, 3H, J = 7.3), 3.85 (s, 3H), 5.13 (q, 1H, J = 7.2), 6.92 (d, 2H, 3H)J = 6.8), 7.93 (d, 2H, J = 7.0).

2-Ethyl-1,3-bis(4-methoxyphenyl)propane-1,3-dione (**2b**).¹¹ Prepared according to the procedure outlined above from **1b** and purified by flash chromatography (ethyl acetate/ hexanes) to afford a viscious oil (95%): ¹H NMR (CDCl₃, 500 MHz) δ 1.01 (t, 3H, J = 7.4), 2.12 (quint, 2H, J = 7.1), 3.80 (s, 6H), 4.98 (t, 1H, J = 6.6), 6.87 (AA'XX', 2H, J = 8.8, 2.1), 7.94 (AA'XX', 2H, J = 8.5, 2.3); APT ¹³C NMR (CDCl₃, 125 MHz) δ 12.9 (CH₃), 23.2 (CH₂), 55.6 (CH), 58.8 (CH₃O), 114.0 (ArCH), 129.3 (ArC), 131.0 (ArCH), 163.8 (ArC), 195.0 (C=O); MS (EI, 70 eV) m/z 312.3 (M⁺).

2-Propyl-1,3-bis(4-methoxyphenyl)propane-1,3-dione (2c). Prepared according to the procedure outlined above from 1c and purified by flash chromatography (ethyl acetate/ hexanes) to afford product as a clear viscous oil (76%): ¹H NMR (CDCl3, 500 MHz) δ 0.95 (t, 3H, J = 7.4), 1.36–1.47 (m, 2H), 2.06–2.11 (m, 2H), 3.83 (s, 6H), 5.05 (t, 1H, J = 6.7) 6.90 (AA'XX', 2H, J = 9.0, 2.5), 7.95 (AA'XX', 2H, J = 9.0, 2.5); ¹³C NMR (CDCl₃, 125 MHz) δ 14.0, 21.5, 31.6, 55.4, 57.1, 113.8, 129.0, 130.8, 163.5, 194.8; HRMS (EI, M⁺) calcd for C₂₀H₂₂O₄ 326.1518, found 326.1512.

2-Isobutyl-1,3-bis(4-methoxyphenyl)propane-1,3-dione (2d). Prepared according to the procedure outlined above from **1d** and purified by flash chromatography (ethyl acetate/ hexanes) to afford product as a clear viscious oil (77%): ¹H NMR (CDCl₃, 500 MHz) δ 0.96 (d, 3H, J = 6.5), 1.70 (m, 1H, J = 6.7), 2.00 (br t, 2H, J = 6.9), 3.82 (s, 6H), 5.18 (t, 1H, J =6.5), 6.89 (d, 2H, J = 8.6), 7.97 (d, 2H, J = 9.0); ¹³C NMR (CDCl₃, 125 MHz) δ 22.8 (CH₃), 27.2 (CH), 38.5 (CH₂), 55.7 (CH₃O), 55.8 (CH), 114.2 (ArCH), 129.4 (ArC), 131.2 (ArCH), 163.2 (C=O); HRMS (EI, M⁺) calcd for C₁₃H₁₈O₄ 340.1674, found 340.1679. Anal. (C₂₁H₂₄O₄) C, H.

2-n-Butyl-1,3-bis(4-methoxyphenyl)propane-1,3-di-

one (2e). Prepared according to the procedure outlined above from 1e and purified by flash chromatography (ethyl acetate/ hexanes) to afford product as a light yellow oil (69%): ¹H NMR (CDCl₃, 500 MHz) δ 0.89 (t, 3H, J = 7.0), 1.37 (m, 4H), 2.11 (q, 2H, J = 7.1), 3.85 (s, 6H), 6.91 (d, 4H, J = 9.0), 7.98 (d, 4H, J = 9.0); ¹³C NMR (CDCl₃, 125 MHz) δ 14.0 (CH₃), 22.9 (CH₂), 22.7 (CH₂), 30.7 (CH₂), 55.7 (CH₃O), 57.7 (CH), 114.2 (ArCH), 129.5 (ArC), 131.2 (ArCH), 163.2 (C=O); HRMS (EI, M⁺) calcd for C₁₃H₁₈O₄ 340.1674, found 340.1675. Anal. (C₁₃H₁₈O₄) C, H.

General Procedure for Pyrazole Synthesis. To a DMF (30 mL) and THF (10 mL) solution containing diketone (1.0 mmol) was added phenylhydrazine hydrochloride (3–5 equiv). The mixture was brought to reflux (oil bath temperature 120 °C) until disappearance of diketone as evident by TLC analysis (8–20 h). The reaction mixture was then allowed to cool to room temperature and diluted with H₂O (30 mL). The product was extracted repeatedly with ethyl acetate (3 × 25 mL) and the organic layers combined and washed sequentially with a satd LiCl solution (25 mL), satd NaHCO₃ (25 mL), and brine (25 mL). The oganic layer was dried over Na₂SO₄ and concentrated under reduced pressure to afford the crude product in the form of an oil, which was purified by flash chromatography or by passage through a short silica plug eluting with an ethyl acetate/hexane solvent system.

3,5-Bis(4-methoxyphenyl)-4-methyl-1-phenyl-1H-pyrazole (3a). The diketone **2a** (250 mg, 0.84 mmol) was reacted with phenylhydrazine hydrochloride (423 mg, 2.94 mmol) according to the general procedure above. Upon purification by flash chromatography (40% ethyl acetate/hexanes) the title compound was obtained as a tan solid (220 mg, 71%): mp 142–143 °C; ¹H NMR (CDCl₃, 500 MHz) δ 2.21 (s, 3H), 3.83 (s, 3H), 3.86 (s, 3H), 6.89 (AA'XX', 2H, J = 8.9, 2.5), 6.99 (AA'XX', 2H, J = 8.9, 2.5), 7.15 (AA'XX', 2H, J = 8.8, 2.5), 7.19–7.32 (m, 5H), 7.74 (AA'XX', 2H, J = 8.9, 2.5); ¹³C NMR (CDCl₃, 125 MHz) δ 10.2, 55.2, 55.3, 113.6, 113.9, 113.9, 123.1, 124.7, 126.5, 126.6, 128.7, 129.1, 131.3, 140.4, 141.2, 150.9, 159.2, 159.4; MS (EI, 70 eV) *m/z* 370 (M⁺). Anal. (C₂₄H₂₂N₂O₂) C, H, N.

4-Ethyl-3,5-bis(4-methoxyphenyl)-1-phenyl-1H-pyrazole (3b).¹¹ Diketone **2b** and phenylhydrazine hydrochloride were reacted as outlined above to afford **3b** as an orange solid after flash chromatography purification (87%): ¹H NMR (CDCl₃, 400 MHz) δ 1.04 (t, 3H, J = 7.6), 2.63 (q, 2H, J =7.6), 3.83 (s, 3H), 3.86 (s, 3H), 6.90 (AA'XX', 2H, J = 8.8, 2.4), 6.99 (AA'XX', 2H, J = 8.8, 2.6), 7.17 (AA'XX', 2H, J = 8.8, 2.4), 7.20 (m, 2H), 7.24 (m, 3H), 7.72 (AA'XX', 2H, J = 9.0, 2.4); ¹³C NMR (CDCl₃, 100 MHz) δ 15.8, 17.3, 55.4, 55.5, 114.1, 114.2, 120.7, 123.5, 124.8, 126.8, 127.0, 128.8, 129.3, 131.5, 140.5, 141.2, 150.8, 159.4, 159.6; HRMS (EI, M⁺) calcd for C₂₅H₂₄N₂O₂ 384.1835, found 384.1837.

3,5-Bis(4-methoxyphenyl)-1-phenyl-4-propyl-1H-pyrazole (3c). Diketone **2c** (200 mg, 0.61 mmol) was reacted with phenylhydrazine hydrochloride (444 mg, 3.07 mmol) according to the general procedure to afford **3c** as an orange oil (130 mg, 53%) after purification by flash chromatography (40% ethyl acetate/hexanes): ¹H NMR (CDCl₃, 500 MHz) δ 0.79 (t, 3H, J = 7.4), 1.42 (sext, 2H, J = 7.7), 2.57 (t, 2H, J = 8.0), 3.83 (s, 3H), 3.86 (s, 3H), 6.90 (AA'XX', 2H, J = 8.6, 2.5), 6.99 (AA'XX', 2H, J = 8.8, 2.5), 7.18–7.32 (m, 5H), 7.15 (AA'XX', 2H, J = 8.6, 2.5), 7.72 (AA'XX', 2H, J = 8.8, 2.5); ¹³C NMR (CDCl₃, 125 MHz) δ 14.05, 23.79, 25.86, 55.09, 55.15, 113.74, 113.85, 118.87, 118.86, 122.86, 124.60, 126.52, 129.49, 131.17, 139.66, 141.31, 150.36, 159.14, 159.33; HRMS (EI, M⁺) calcd for C₂₆H₂₆N₂O₂ 398.1994, found 398.2000.

4-Isobutyl-3,5-bis(4-methoxyphenyl)-1-phenyl-1*H*-pyrazole (3d). Diketone 2d was reacted with phenylhydrazine hydrochloride according to the general procedure to afford 3d as an orange oil (85%) after a short silica plug (20% ethyl acetate/hexanes). The purified material was then directly used in the subsequent deprotection step.

4-Butyl-3,5-bis(4-methoxyphenyl)-1-phenyl-1H-pyrazole (3e). Diketone **2e** was reacted with phenylhydrazine hydrochloride according to the general procedure to afford **3e** as a reddish oil (86%) after purification by flash chromatography (20% ethyl acetate/hexanes). This material was then directly used in the subsequent deprotection step.

4-Ethyl-1,3,5-tris(4-methoxyphenyl)-1*H*-pyrazole (3f). Diketone **2b** (400 mg, 1.28 mmol) was reacted with 4-methoxyphenylhydrazine hydrochloride (1.11 g, 6.40 mmol) according to the general procedure. The crude product was purified by flash chromatography (2% acetone/CH₂Cl₂) to afford **3f** as an oil (351 mg, 66%): ¹H NMR (CDCl₃, 500 MHz) δ 1.06 (t, 3H, J = 7.5), 2.65 (q, 2H, J = 7.5), 3.73 (s, 3H), 3.79 (s, 3H), 3.83 (s, 3H), 6.78 (AA'XX', 2H, J = 9.0, 2.8) 6.89 (AA'XX', 2H, J = 8.8, 2.4), 7.16 (AA'XX', 2H, J = 8.8, 2.4), 7.16 (AA'XX', 2H, J = 8.8, 2.4), 7.21 (AA'XX', 2H, J = 9.0, 2.7), 7.74 (AA'XX', 2H, J = 8.7, 2.4); ¹³C NMR (CDCl₃, 125 MHz) δ 15.52, 17.09, 55.05, 55.10, 55.22, 113.65, 113.80, 113.83, 119.78, 123.11, 125.96, 126.78, 128.95, 131.18, 133.45, 140.84, 149.94, 158.99, 157.97, 159.22; HRMS (EI, M⁺) calcd for C₂₆H₂₆N₂O₃ 414.1943, found 414.1942.

1,3,5-Tris(4-methoxyphenyl)-4-propyl-1H-pyrazole (3g). Diketone **2c** (500 mg, 1.52 mmol) was reacted with 4-methoxyphenylhydrazine hydrochloride (800 mg, 4.59 mmol) according to the general procedure to afford 362 mg of **3g** as a red oil (67%) after a short silica plug (20% ethyl acetate/ hexanes). This material was then directly used in the subsequent deprotection step.

4-Isobutyl-1,3,5-tris(4-methoxyphenyl)-1*H***-pyrazole (3h).** Diketone **2d** was reacted with 4-methoxyphenylhydrazine hydrochloride according to the general procedure to afford **3h** as a light orange oil (85%) after a short silica plug (20% ethyl acctate/hexanes). The purified material was then directly used in the subsequent deprotection step.

4-Butyl-1,3,5-tris(4-methoxyphenyl)-1*H***-pyrazole (3i).** Diketone **2e** was reacted with 4-methoxyphenylhydrazine hydrochloride according to the general procedure to afford **3i** as a light orange oil (87%) after a short silica plug (20% ethyl acetate/hexanes). The purified material was then directly used in the subsequent deprotection step.

General Demethylation Procedure. To a stirred solution of methyl-protected pyrazole (1 equiv) in CH_2Cl_2 at -78 °C was added dropwise a 1 M BBr₃ solution in CH_2Cl_2 (3–5 equiv). Upon complete addition of BBr₃, the reaction was maintained at -78 °C for 1 h and then allowed to reach room temperature and stir for an additional 16 h. The mixture was cooled to 0 °C and carefully quenched with H_2O (15–25 mL). The product was then repeatedly extracted with EtOAc and the organic layers dried over Na_2SO_4 . Upon solvent removal the crude phenolic products were purified by flash chromatography and/or recrystallization from MeOH/CH₂Cl₂ mixtures.

3,5-Bis(4-hydroxyphenyl)-4-methyl-1-phenyl-1H-pyrazole (4a). A stirred CH₂Cl₂ solution of **3a** (200 mg, 0.54 mmol) was deprotected using BBr₃ according to the general demethylation procedure. Purification by flash chromatography (5% CH₃OH/CH₂Cl₂) afforded the title compound as a tan solid (54 mg, 30%): mp 225-230 °C; ¹H NMR (MeOD- d_4 , 500 MHz) δ 2.13 (s, 3H), 6.76 (AA'XX', 2H, J = 8.4, 2.7), 6.93 (AA'XX', 2H, J = 8.7, 2.5), 7.01 (AA'XX', 2H, J = 8.8, 2.5), 7.54 (AA'XX', 2H, J = 8.8, 2.5), 7.54 (AA'XX', 2H, J = 8.6, 2.4), 7.35-7.22 (m, 5H); ¹³C NMR (MeOD- d_4 , 125 MHz) δ , 7, 112.8, 114.7, 114.8, 120.9, 124.4, 124.8, 126.7, 128.3, 128.9, 130.9, 139.8, 151.4, 157.1, 157.4; MS (EI, 70 eV) m/z 342. Anal. (C₂₂H₁₈N₂O₂·H₂O) C, H, N.

4-Ethyl-3,5-bis(4-hydroxyphenyl)-1-phenyl-1H-pyrazole (4b).¹¹ A stirred CH₂Cl₂ solution of **3b** (100 mg, 0.26 mmol) was deprotected using BBr₃ according to the general demethylation procedure. Purification by flash chromatography (5% CH₃OH/CH₂Cl₂) afforded the tile compound as a white solid (50 mg, 54%): mp 247–248 °C; ¹H NMR (MeOD- d_4 , 400 MHz) δ 0.98 (t, 3H, J = 7.4), 2.60 (q, 2H, J = 7.5), 6.78 (AA'XX', 2H, J = 8.8, 2.4), 6.88 (AA'XX', 2H, J = 8.7, 2.5), 7.05 (AA'XX', 2H, J = 8.7, 2.5); HRMS (EI, M⁺) calcd for C₂₃H₂₁N₂O₂ 357.1611, found 357.1603.

3,5-Bis(4-hydroxyphenyl)-1-phenyl-4-propyl-1*H***-pyrazole (4c).** A stirred CH₂Cl₂ solution of **3c** (107 mg, 0.27 mmol) was deprotected using BBr₃ according to the general demethylation procedure. The crude product was purified by flash chromatography (10% CH₃OH/CH₂Cl₂) to afford **4c** as a tan solid (85 mg, 86%): mp 240–245 °C; ¹H NMR (MeOD- d_4 , 500 MHz) \diamond 0.72 (t, 3H, J = 7.4), 1.35 (sext, 2H, J = 7.5), 2.55 (t, 2H, J = 7.7), 6.77 (AA'XX', 2H, J = 8.7, 2.5), 6.88 (AA'XX', 2H, J = 8.5, 2.4), 7.03 (AA'XX', 2H, J = 8.5, 2.4), 7.32–7.27 (m, 5H), 7.50 (AA'XX', 2H, J = 8.8, 2.5); ¹³C NMR (MeOD- d_4 , 400 MHz) \diamond 12.73, 13.28, 25.28, 114.75, 114.89, 118.26, 121.13, 124.78, 124.82, 126.69, 128.26, 128.99, 131.02, 139.71, 142.12, 151.30, 157.08, 157.49; HRMS (EI, M⁺) calcd for C₂₄H₂₂N₂O₂ 370.1681, found 370.1676.

4-Isobutyl-3,5-bis(4-hydroxyphenyl)-1-phenyl-1H-pyrazole (4d). A stirred CH₂Cl₂ solution of **3d** (100 mg, 0.24 mmol) was deprotected using BBr₃ according to the general demethylation procedure. The crude product was purified by flash chromatography (10% CH₃OH/CH₂Cl₂) to afford **4d** as a tan powder (70 mg, 76%): mp 225 °C dec; ¹H NMR (MeOD- d_4 , 500 MHz) δ 0.63 (d, 6H, J = 6,5), 1.51 (m, 1H, J = 7.0), 2.51 (d, 2H, J = 7.5), 6.76 (AA'XX', 2H, J = 9.0, 2.3), 6.87 (AA'XX', 2H, J = 8.9, 2.3), 7.02 (AA'XX', 2H, J = 8.5, 2.4), 7.26 (m, 5H), 7.49 (AA'XX', 2H, J = 8.0, 2.3); ¹³C NMR (MeOD- d_4 , 125 MHz) δ 22.8, 29.9, 33.8, 116.4, 116.6, 119.1, 123.0, 126.6, 126.8, 128.4, 129.9, 130.9, 132.9, 141.5, 144.2, 153.4, 158.8, 159.2; FAB-HRMS (M + 1) calcd for C₂₅H₂₅N₂O₂ 385.1916, found 385.1916.

4-Butyl-3,5-bis(4-hydroxyphenyl)-1-phenyl-1H-pyrazole (4e). A stirred CH₂Cl₂ solution of 3e (100 mg, 0.24 mmol) was deprotected using BBr3 according to the general demethylation procedure. The crude product was purified by flash chromatography (30% EtOAc/hexanes) to afford a tan solid. This material was subsequently recrystallized from 5-10% MeOH/CH₂Cl₂ to afford the title compound as small off-white crystals (59 mg, 64%): mp 205.5-207.5 °C; ¹H NMR (MeOD d_{J} , 400 MHz) δ 0.72 (t, 3H, J = 7.2), 1.16 (sext, 2H, J = 7.2), 1.34 (quint, 2H, J = 7.2), 2.60 (t, 2H, J = 7.2), 4.95 (br s, 2H exchange with D_2O), 6.78 (d, 2H, J = 9.0), 6.91 (d, 2H, J =8.0), 7.02 (d, 2H, J = 8.8), 7.26 (m, 5H), 7.53 (d, 2H, J = 8.4); ¹³C NMR (MeOD-*d*₄, 125 MHz) δ 14.6, 23.9, 24.8, 34.3, 116.8, 116.9, 120.5, 123.2, 126.8, 128.7, 130.3, 131.0, 131.1, 133.1, 141.7, 144.0, 153.3, 159.0, 159.4. Anal. $(C_{25}H_{24}N_2O_2 \cdot 0.1H_2O)$ C, H, N.

4-Ethyl-1,3,5-tris(4-hydroxyphenyl)-1H-pyrazole (4f). A stirred CH₂Cl₂ solution of **3f** (200 mg, 0.48 mmol) was deprotected using BBr₃ according to the general demethylation procedure. A crude oil was isolated which was triturated with a 10% CH₃OH/CH₂Cl₂ solution from which the desired product precipitated. The white powder was collected by filtration and recrystallized from CH₃OH/CH₂Cl₂ to afford the title compound **4f** (175 mg, 98%): mp 210–215 °C; ¹H NMR (MeOD-*d*₄, 400 MHz): δ 0.96 (t, 3H, J = 7.5) 2.58 (q, 2H, J = 7.5) 6.70 (AA'XX', 2H, J = 8.8, 2.6), 6.77 (AA'XX', 2H, J = 8.6, 2.3), 6.87 (AA'XX', 2H, J = 8.8, 2.3), 7.10 (m, 4H), 7.48 (AA'XX', 2H, J = 8.6, 2.4); ¹³C NMR (MeOD-*d*₄, 100 MHz) δ 14.5, 16.6, 114.8, 114.9, 114.9, 119.2, 121.4, 125.0, 126.8, 129.1, 131.2, 131.9, 142.1, 150.5, 156.7, 157.1, 157.5. Anal. (C₂₃H₂₀N₂O₃·0.7H₂O) C, H, N.

1,3,5-Tris(4-hydroxyphenyl)-4-propyl-1H-pyrazole (4g). A stirred CH_2Cl_2 solution of **3g** (200 mg, 0.48 mmol) was deprotected using BBr3 according to the general demethylation procedure. A crude oil was isolated which was triturated with a 10% CH₃OH/CH₂Cl₂ solution from which the desired product precipitated. The white powder was collected by filtration and recrystallized from CH_3OH/CH_2Cl_2 to afford the title compound 4g (125 mg, 68%): mp 230 °C dec; ¹H NMR (MeOD-d₄, 400 MHz) δ 0.72 (t, 3H, J = 7.2), 1.33 (sext, 2H, J = 7.6), 2.54 (t, 2H, J = 8), 6.70 (AA'XX', 2H, J = 8.8, 2.4), 6.76 (AA'XX', 2H, J = 6.8, 2.0, 6.87 (AA'XX', 2H, J = 8.8, 2.4), 7.02 (AA'XX', 2H, J = 8.8, 2.4), 7.05 (AA'XX', 2H, J = 9.2, 2.4), 7.47 (AA'XX', 2H, AA'XX', 2H, AA'2H, J = 8.8, 2.0; ¹³C NMR (MeOD- d_4 , 100 MHz) δ 25.8 (CH₃), 36.4 (CH₂), 38.4 (CH₂), 127.0 (C), 128.0 (C), 128.5 (C), 130.5 (CH), 134.3 (CH), 137.9 (CH), 138.9 (C), 140.3 (C), 141.3 (C), 142.8 (C), 143.3 (C), 144.1 (C),144.7 (CH), 155.3 (CH), 163.6 (CH), 169.5 (CH), 170.3 (CH). Anal. ($C_{24}H_{22}N_2O_3 \cdot 0.6H_2O$) C, H, N

4-Isobutyl-1,3,5-tris(4-hydroxyphenyl)-1*H*-pyrazole (4h). A stirred CH₂Cl₂ solution of **3h** (250 mg, 0.56 mmol) was deprotected using BBr₃ according to the general demethylation procedure. Recrystallization from CH₃OH/CH₂Cl₂ afforded the title compound **4h** as an off-white powder (80 mg, 37%): mp 226–228 °C; ¹H NMR (MeOD- d_4 , 500 MHz) δ 0.61 (d, 6H, J = 6.5), 1.50 (m, 1H, J = 7.0), 2.50 (d, 2H, J = 7.5), 4.87 (s, 3H, OH), 6.70 (AA'XX', 2H, J = 8.5, 2.0), 6.75 (AA'XX', 2H, J = 8.5, 2.3), 6.85 (AA'XX', 2H, J = 9.0, 2.3), 7.01 (AA'XX', 2H, J = 9.0, 2.3), 7.05 (AA'XX', 2H, J = 9.0, 2.3), 7.45 (AA'XX', 2H, J = 9.0, 2.3), 1³C NMR (MeOD- d_4 , 125 MHz) δ 22.9, 29.9, 33.9, 116.4, 116.5, 116.6, 118.3, 123.2, 126.9, 128.3, 130.9, 132.9, 133.5, 144.3, 152.7, 158.2, 158.6, 158.9. Anal. (C₂₅H₂₄N₂O₃· 0.7H₂O) C, H, N.

4-Butyl-1,3,5-tris(**4-hydroxyphenyl**)-**1***H*-**pyrazole** (**4**). A stirred CH₂Cl₂ solution of **3i** (200 mg, 0.45 mmol) was deprotected using BBr₃ according to the general demethylation procedure. Recrystallization from CH₃OH/CH₂Cl₂ afforded the title compound **4i** as an off-white powder (90 mg, 50%): mp **214**–230 °C dec; ¹H NMR (MeOD-*d*₄, 500 MHz) δ 0.71 (t, 3H, J = 7.5), 1.13 (sext, 2H, J = 7.0), 1.30 (quint, 2H, J = 8.5), 2.57 (t, 2H, J = 8.5), 2.67 (t, 2H, J = 8.5, 2.3), 6.86 (AA'XX', 2H, J = 9.0, 2.4), 6.76 (AA'XX', 2H, J = 8.5, 2.3), 7.05 (AA'XX', 2H, J = 8.5, 2.3), 7.05 (AA'XX', 2H, J = 8.5, 2.3), 7.46 (AA'BB', 2H, J = 8.5, 2.3); ¹³C NMR (MeOD-*d*₄, 400 MHz) δ 14.1, 23.5, 24.4, 33.9, 116.3, 116.4, 116.5, 119.3, 123.0, 126.7, 128.3, 130.7, 132.8, 133.5, 143.9, 152.4, 158.3, 158.7, 159.0. Anal. (C₂₅H₂₄N₂O₃·0.3H₂O) C, H, N.

4-Isopropyl-3,5-bis(4-methoxyphenyl)-1-phenyl-1*H***-pyrazole (6)**.¹³ Upon solid support cleavage and solvent removal, the crude solid was recrystallized from 25% ethyl acetate/hexane to afford **6** as small cubic crystals (30 mg, 11% over three steps): mp 225–230 °C; ¹H NMR (MeOD-*d*₄, 400 MHz) δ 1.09 (d, 6H, *J* = 7.0 Hz) 2.98 (septet, 1H, *J* = 7.11 Hz), 6.76 (AA'XX', 2H, *J* = 9.1, 2.6), 6.88 (AA'XX', 2H, *J* = 9.0, 2.6), 7.1 (AA'XX', 2H, *J* = 8.7, 2.5), 7.20–7.30 (m, 5H), 7.39 (AA'XX', 2H, *J* = 8.8, 2.40); ¹³C NMR (MeOD-*d*₄, 100 MHz) δ 22.5, 24.5, 114.5, 114.6, 121.6, 124.6, 125.0, 126.7, 128.2, 130.2, 131.8, 139.6, 141.3, 151.3, 157.2, 157.6; HRMS (EI, M⁺) calcd for C₂₄H₂₂N₂O₂ 370.1681, found 370.1674.

1,3-Bisphenylpropane-1,3-dione (10). To a stirred solution of diketone **9** (2g, 8.9 mmol) in freshly distilled THF (50 mL), was added 8.9 mL (8.9 mmol) of tetrabutylammonium fluoride (1 M in THF) and stirred for 30 min. The solution was concentrated under reduced pressure and dissolved in 50 mL of CHCl₃. Ethyl iodide (1.4 mL, 17.8 mmol) was added in one portion and stirred at room temperature overnight. The solution was concentrated and the crude was purified by flash chromatgraphy (petroleum ether/CH₂Cl₂, 1:1) to give 920 mg of **19** as a white solid in 41% yield: ¹H NMR (CDCl₃, 400 MHz) δ 1.05 (t, 3H, J = 7.4), 2.17 (quint, 2H, J = 7.4), 5.12 (t, 1H, J = 6.5) 7.42–7.58 (m, 6H), 7.94–7.98 (m, 4H); ¹³C NMR δ 12.8, 22.9, 58.7, 128.5, 128.8, 133.4, 136.1, 196.1; MS (EI, 70 eV) m/z 252.1 (M⁺).

4-Ethyl-1-(4-methoxyphenyl)-3,5-bisphenyl-1*H***-pyrazole (11). The diketone 10 (300 mg, 1.2 mmol) was reacted with 4-methoxyphenylhydrazine hydrochloride (830 mg, 4.7 mmol) according to the general procedure for pyrazole synthesis. The crude product was purified by flash chromatography (petroleum ether/CH₂Cl₂, 1:1) to afford 276 mg of 11 as a yellow oil (65%): ¹H NMR (CDCl₃, 400 MHz) \delta 1.04 (t, 3H, J = 7.5), 2.67 (q, 2H, J = 7.5), 3.77 (s, 1H), 6.77 (AA'XX', 2H, J = 1.0, 2.2), 7.19 (AA'XX', 2H, J = 9.1, 2.2), 7.23–7.48 (m, 9H), 7.78 (AA'XX', 2H, J = 8.2, 2.5); ¹³C NMR \delta 15.6, 17.1, 55.4, 113.8, 120.4, 126.1, 127.5, 127.9, 128.1, 128.41, 128.45, 130.1, 130.9, 133.4, 134.1, 141.2, 150.4, 158.2; MS (EI, 70 eV) m/z 354.2 (M⁺).**

4-Ethyl-1-(4-hydroxyphenyl)-3,5-bisphenyl-1H-pyrazole (7a). A stirred CH₂Cl₂ solution of **11** (274 mg, 0.77 mmol) was deprotected with BBr₃ according to the general demethylation procedure. The crude was purified by flash chromatography (CH₂Cl₂/acetone, 3:1) to give 84 mg of **7a** as a white solid (32%): mp 184–185 °C; ¹H NMR (CD₃OD, 400 MHz) δ 1.03 (t, 3H, J = 7.5), 2.66 (q, 2H, J = 7.5), 6.65 (AA'XX', 2H J= 8.8, 2.2), 7.09 (AA'XX', 2H, J = 9.0, 2,1), 7.2–7.5 (m, 8H), 7.77 (AA'XX', 2H, J = 8.4); ¹³C NMR δ 15.5, 17.0, 115.8, 120.2, 126.6, 127.7, 128.0, 128.2, 128.4, 128.5, 130.0, 130.6, 132.6, 133.8, 141.6, 150.5, 155.2; MS (EI, 70 eV) m/z 340.2 (M^+). Anal. $(\rm C_{23}H_{20}ON_2)$ C, H, N.

1,3-Bis(4-methoxyphenyl)-5-phenyl-4,5-dihydro-1Hpyrazole (13a). A mixture of commercially available 4'methoxychalcone (12a; 253 mg, 1.1 mmol) and 807 mg of 4-methoxyphenylhydrazine (4.6 mmol) in 10 mL of anhydrous DMF was heated to 85 °C overnight. The reaction solution was cooled to room temperature and partitioned with diethyl ether and water. The organic layer was washed with water, dried (MgSO₄), and concentrated. The crude was then recrystallized from a mixture of ethyl acetate and hexanes to give 198 mg of 13a as light yellow crystals (48%): ¹H NMR (CDCl₃, 400 MHz) δ 3.12 (dd, 1H, J = 16.7, 8.4), 3.74 (s, 3H), 3.86 (s, 3H), 3.74-3.88 (m, 1H), 5.14 (dd, 1H, J = 11.9, 8.6), 6.8 (AA'XX', 2H, J= 9.0), 6.94 (AA'XX', 2H, J = 9.0), 7.04 (AA'XX', 2H, J = 9.0), $7.26-7.42 \text{ (m, 5H)}, 7.68 \text{ (d, 2H, } J = 9.0\text{)}; {}^{13}\text{C} \text{ NMR} \delta 43.9, 55.3,$ 55.6, 65.8, 114.0, 114.4, 114.8, 126.1, 127.5, 127.8, 128.0, 129.1, 140.1, 142.9, 146.9, 153.2, 160.0; HRMS (EI, M⁺) calcd for C₂₃H₂₂N₂O₂ 358.1688, found 358.1681.

1,5-Bis(4-methoxyphenyl)-3-phenyl-4,5-dihydro-1*H***-pyrazole (13b).** A mixture of commercially available 4-methoxychalcone (**12b**; 2 g, 8.4 mmol) and 7.3 g of 4-methoxyphenylhydrazine (42 mmol) in 80 mL of anhydrous DMF was heated to 85 °C overnight. The reaction solution was cooled to room temperature and partitioned with diethyl ether and water. A yellow solid precipitated and was collected by filtration to give 2.6 g of **13b** (86%): ¹H NMR (CDCl₃, 400 MHz) δ 3.1 (dd, 1H, J = 15.6,7.8), 3.72 (s, 3H), 3.78 (s, 3H), 3.8 (d, 1H, J = 5.6), 5.13 (dd, 1H, J = 12.0, 9.0), 6.76 (AA'XX', 2H, J = 9.1, 2.3), 6.87 (AA'XX', 2H, J = 8.7, 2.1), 7.01 (d, 2H, J = 9.0), 7.2–7.4 (m, 5H), 7.7 (d, 2H, J = 7.3); ¹³C NMR δ 43.7, 55.2, 55.6, 65.2, 114.4, 114.4, 114.9, 125.6, 127.3, 128.3, 128.5, 132.9, 134.7, 139.7, 146.3, 153.3, 158.9; MS (EI, 70 eV) m/z 358.2 (M⁺).

4-Ethyl-1,3-bis(4-methoxyphenyl)-5-phenyl-4,5-dihydro-1H-pyrazole (14a). To a solution of lithium diisopropylamide in 20 mL of THF [prepared by dropwise addition of 0.88 mL of n-BuLi (1.41 mmol) to 0.21 mL (1.5 mmol) of diisopropylamine in 18 mL of THF at -78 °C] was added a solution of 317 mg (0.88 mmol) of pyrazoline 13a in 8 mL of THF dropwise via syringe at -78 °C and stirred for 1 h. To the dark red solution was added iodoethane (0.11 mL, 1.31 mmol) in one portion and the resulting yellow solution was warmed to room temperature overnight. The reaction was quenched with 5 mL of brine, the aqueous layer was separated and extracted with $\rm CH_2\rm Cl_2,$ the organic layer was dried (MgSO_4) and concentrated in vacuo. The residue was purified by flash chromatography (ether/hexanes, 3:2) to afford 262 mg of 14a as a yellow foam (77%): ¹H NMR (CDCl₃, 400 MHz) δ 0.8 (t, 3H, J = 7.3), 1.45 (tq, 1H, J = 4, 2), 1.55 (m, 1H), 3.2 (m, 1H), 3.25 (s, 3H), 3.3(s, 3H), 4.8 (d, 1H, 3.4), 6.77 (AA'XX', 2H, J = 8.9), 6.82 (AA'XX', 2H, J = 8.5), 7.0 (AA'XX', 2H, J = 8.7), 7.13 (AA'XX', 2H, J = 8.5), 7.25–7.40 (m, 3H), 7.70 (AA'XX', 2H, J = 7.3); ¹³C NMR & 10.5, 25.4, 54.8, 55.1, 57.8, 69.7, 114.4, 114.4, 115.0, 125.8, 126.0, 127.5, 127.7, 129.3, 139.3, 142.8, 148.4, 153.6, 160.2; HRMS (EI, M⁺) calcd for C₂₅H₂₆N₂O₂ 386.1999, found 386.1994.

4-Ethyl-1,5-bis(4-methoxyphenyl)-3-phenyl-4,5-dihydro-1H-pyrazole (14b). Prepared from lithium diisopropylamide (4.65 mmol), pyrazoline **13b** (1 g, 2.8 mmol), and ethyl iodide (0.45 mL, 5.6 mmol) by the procedures used to prepare pyrazoline **14a**. The crude product was purified by flash chromatography (hexanes/ether 3:2) to give 0.33 g (38%) of **14b** a yellow foam: ¹H NMR (CDCl₃, 400 MHz) δ 1.03 (t, 3H, J = 7.3), 1.69 (tq, 1H, J = 4, 2), 1.85 (m, 1H), 3.37 (m, 1H), 3.73 (s, 3H), 3.75 (s, 3H), 4.91 (d, 1H, 3.4), 6.77 (AA'XX', 2H, J = 8.7), 7.13 (AA'XX', 2H, J = 8.5), 7.25–7.40 (m, 3H), 7.70 (AA'XX', 2H, J = 7.3); ¹³C NMR δ 10.5, 25.1, 55.2, 55.6, 57.3, 68.9, 113.9, 114.4, 114.5, 125.8, 126.8, 127.9, 128.5, 132.6, 134.1, 138.5, 148.5, 152.8, 158.8; MS (EI, 70 eV) m/z 386.2 (M⁺).

4-Ethyl-1,3-bis(4-methoxyphenyl)-5-phenyl-1*H*-pyrazole (15a). To a stirred solution of 14a (27.8 mg, 0.07 mmol) in 3 mL of benzene was added 78 mg (0.9 mmol) of MnO₂. The solution was heated to 100 °C with a Dean–Stark trap for 2 h, cooled to room temperature, filtered through Celite and concentrated. The crude was chromatographed (EtOAc/hexanes, 1:10) to afford 28 mg of **15a** (100%): ¹H NMR (CDCl₃, 400 MHz) δ 1.12 (t, 3H, J = 7.5), 2.65 (q, 2H, J = 7.5), 3.78 (s, 3H), 3.85 (s, 3H), 6.77 (AA'XX', 2H, J = 9.0, 2.0), 7.0 (AA'XX', 2H, J = 9.0, 2.0), 7.22–7.40 (m, 5H), 7.71 (d, 2H, J = 8.6, 2.0); ¹³C NMR δ 15.5, 17.1, 55.2, 55.4, 113.8, 113.8, 120.1, 126.1, 126.8, 128.1, 128.4, 129.1, 130.1, 131.1, 133.5, 141.1, 150.2, 158.2, 159.1; HRMS (EI, M⁺) calcd for C₂₅H₂₄N₂O₂ 385.1912, found 385.1916.

4-Ethyl-1,5-bis(4-methoxyphenyl)-3-phenyl-1H-pyrazole (15b). A mixture of 147 mg (0.37 mmol) of pyrazoline 14b and 127 mg (0.56 mmol) of dichlorodicyanoquinone in 10 mL of benzene was heated to reflux for 5 h. The mixture was cooled to room temperature and filtered through a plug of Celite with diethyl ether. The filtrate was concentrated in vacuo and the residue was chromatographed (hexanes/ethyl acetate, 4:6) to give 130 mg (90%) of 15b as a white solid: ¹H NMR (CDCl₃, 400 MHz) δ 1.04 (t, 3H, J = 7.5), 2.65 (q, 2H, J= 7.5), 3.78 (s, 3H), 3.83 (s, 3H), 6.79 (AA'XX', 2H, J = 8.9, 2.0), 6.89 (AA'XX', 2H, J = 8.5, 2.0), 7.16 (AA'XX', 2H, J = 8.3, 1.8), 7.20 (AA'XX', 2H, J = 8.7, 2.1), 7.3–7.5 (m, 3H), 7.77 (AA'XX', 2H, J = 8.8, 1.8); ¹³C NMR δ 15.6, 17.12, 55.2, 55.4, 113.7, 113.9, 120.2, 123.2, 126.1, 127.4, 127.9, 128.4, 131.3, 133.5, 134.27, 141.07, 150.3, 158.1, 159.3; MS (EI, 70 eV) m/z 384.2 (M⁺).

4-Ethyl-1,3-bis(4-hydroxyphenyl)-5-phenyl-1H-pyrazole (8a). A stirred CH₂Cl₂ solution of **15a** (26 mg, 0.068 mmol) was deprotected with BBr₃ according to the general demethylation procedure. The crude was purified by flash chromatography (CH₂Cl₂/MeOH, 10:1) to give 17 mg of **8a** as a tan solid (70%): mp 225-227 °C; ¹H NMR (CD₃OD, 500 MHz) δ 0.8 (t, 3H, J = 7.5), 2.5 (q, 2H, J = 7.5), 6.58 (AA'XX', 2H, J = 8.5, 2.0), 6.94 (AA'XX', 2H, J = 8.9, 2.0), 7.1-7.3 (m, 5H), 7.39 (AA'XX', 2H, J = 8.9, 2.0), 7.1-7.3 (m, 5H), 7.39 (AA'XX', 2H, J = 8.9, 2.0); ¹³C NMR δ 15.8, 17.9, 116.3, 116.4, 121.0, 122.6, 128.2, 128.9, 129.3, 129.6, 132.6, 133.2, 135.3, 143.8, 151.7, 158.2, 159.0; HRMS (EI, M⁺) calcd for C₂₃H₂₀N₂O₂ 356.1598, found 356.1603.

4-Ethyl-1,5-bis(4-hydroxyphenyl)-3-phenyl-1H-pyrazole (8b). A stirred CH₂Cl₂ solution of **15b** (130 mg, 0.34 mmol) was deprotected with BBr₃ according to the general demethylation procedure. The crude was purified by flash chromatography (CH₂Cl₂/acetone, 3:1) to give 42 mg of **8b** as a white solid (35%): mp 225–226 °C; ¹H NMR (CD₃OD, 500 MHz) δ 0.96 (t, 3H, J = 7.5), 2.62 (q, 2H, J = 7.5), 6.71 (AA'XX', 2H, J = 9.1, 2.2), 6.77 (AA'XX', 2H, J = 9.1, 2.2), 7.06 (AA'XX', 2H, J = 9.1, 2.2), 7.07 (AA'XX', 2H, J = 9.1, 2.2), 7.35–7.47 (m, 3H), 7.67 (AA'XX', 2H, J = 8.9, 2.0); ¹³C NMR δ 15.8, 17.9, 116.3, 116.4, 121.0, 122.6, 128.2, 128.9, 129.3, 129.6, 132.6, 133.2, 135.3, 143.8, 151.7, 158.2, 159.0; HRMS (EI, M⁺) calcd for C₂₃H₂₀N₂O₂·0.75H₂O) C, H, N.

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Estrogen Pyrazoles: Defining the Pyrazole Core Structure and the Orientation of Substituents in the Ligand Binding Pocket of the Estrogen Receptor

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Abstract—Previously, we reported that certain tetrasubstituted 1.3,5-triaryl-4-alkyl-pyrazoles bind to the estrogen receptor (ER) with high affinity (Fink, B. E.; Mortenson, D. S.; Stauffer, S. R.; Aron, Z. D.; Katzenellenbogen, J. A. *Chem. Biol.* **1999**, *6*, 205–219; Stauffer, S. R.; Katzenellenbogen, J. A. *J. Comb. Chem.* **2000**, *2*, 318–329; Stauffer, S. R.; Coletta, C. J.; Sun, J.; Tedesco, R.; Katzenellenbogen, B. S.; Katzenellenbogen, J. A. *J. Med. Chem.* **2000**, submitted). To investigate how cyclic permutation of the two nitrogen atoms of a pyrazole might affect ER binding affinity, we prepared a new pyrazole core isomer, namely a 1,3,4-triaryl-5-alkyl-pyrazole (**2**), to compare it with our original pyrazole (**1**). We also prepared several peripherally matched core pyrazole isomer sets to investigate whether the two pyrazole series share a common binding orientation. Our efficient, regioselective synthetic route to these pyrazoles relies on the acylation of a hydrazone anion, followed by cyclization, halogenation, and Suzuki coupling. We found that the ER accommodates 1,3,4-triaryl-pyrazoles of the isomeric series only somewhat less well than the original 1,3,5-triaryl series, and it appears that both series share a common binding mode. This preferred orientation for the 1,3,5-triaryl-4-alkyl-pyrazoles is supported by binding affinity measurements of analogues in which the phenolic hydroxyl groups were systematically removed from each of the three aryl groups, and the orientation is consistent, as well, with molecular modeling studies. These studies provide additional insight into the design of heterocyclic core structures for the development of high affinity ER ligands by combinatorial methods. () 2000 Elsevier Science Ltd. All rights reserved.

Introduction

The estrogen receptor (ER) binds a remarkably wide range of non-steroidal ligands,¹ and the diverse core structures of these ligands span a wide range of synthetic accessibility.² We have been intrigued, in particular, by the design of non-steroidal ligand cores that might be easily prepared and thus would be well suited to combinatorial expansion. As part of this investigation, we identified pyrazoles as favorable heterocyclic core building blocks,³ and we found that by attaching a sufficient number of appropriate substituents onto this core system, we could obtain high affinity ligands for the ER.^{3–5}

An issue which had interested us initially about these systems was whether the heterocyclic core structure itself plays an active role in ligand–receptor interaction, or whether it acts merely as a inert scaffold, simply displaying groups in a geometry appropriate for filling the various subpockets that make up the ligand cavity of ER.³ Interestingly, we found that there were large differences in binding affinity (up to 50-fold) for ligands that had identical peripheral substitution patterns but different core structures (e.g. the two diazoles, imidazoles versus pyrazoles; Figure 1).³ Clearly, our initial thought that the ligand core structure might be acting as a passive entity—simply to position peripheral substituents—was not true. In the case shown below, the lower ER binding affinity for the imidazoles compared to the pyrazoles was attributed, at least in part, to the significantly greater dipole moment of the imidazoles.³

In further consideration, we wondered whether the distribution of heteroatoms within the same heterocyclic system would also have a significant effect on ER binding affinity. In the two *inidazoles* shown in Figure 1, the different position of the nitrogen atoms had little effect on their binding affinity. However, we were curious whether the related cyclic permutation of the two nitrogen atoms of a *pyrazole* might have a more significant effect on ER binding affinity.

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[Imidazoles]

Figure 1. Effect of different core structures of selected five-member heterocycles (diazoles) on their estrogen receptor relative binding affinity (RBA).

To investigate this issue, we have prepared a new pyrazole core isomer, namely a 1,3,4-triaryl-5-alkyl-pyrazole (2), to compare it with our original 1,3,5-triaryl-4-alkyl-pyrazole (1) (Fig. 2). This new compound (2) represents the only other possible pyrazole system capable of displaying the same relative substitution pattern of aryl, alkyl, aryl, and phenyl groups as does our original pyrazole 1. To extend this investigation further, we synthesized several other peripherally matched core pyrazole isomer





Figure 2. Original 1.3.5-triaryl-pyrazole 1 (the basis of Series 1) and the new 1.3.4-triaryl-pyrazole isomer 2 (the basis of Series II).

sets, in which some of the phenolic hydroxyl groups are systematically deleted, to establish more definitively which core isomer is most favored and to investigate whether they share a common binding orientation.

Results and Discussion

Synthesis of 1,3,4-triaryl-5-alkyl-pyrazoles

Our initial attempts to synthesize the pyrazole isomer 2 involved conditions similar to those used in previous studies (Scheme 1).³ We first prepared diketone 3 in high yield, starting from the deoxybenzoin and propionic anhydride. However, our attempts to generate the pyrazole by condensation of the dione with 4-methoxyphenylhydrazine were unsuccessful and resulted in a complex mixture that contained the two possible retro-Claisen condensation products from cleavage of the 1,3-dione.

Because of these difficulties and the fact that this route was not likely to be regioselective, we investigated an alternative, potentially regioselective approach that involved a one-pot acylation and cyclization procedure starting from an appropriate hydrazone, as illustrated in Figure 3. Unfortunately, we could not isolate the hydrazone (4), because it rapidly underwent Fischer indole cyclization to the 2,3-diaryl-indole product (5). Even under mild conditions, we obtained only starting material or indole product, but no hydrazone. Apparently, the 4-methoxyphenyl substituent in the deoxybenzoin starting material facilitates the [3,3]-sigmatropic cyclization of the ene-phenylhydrazole intermediate by stabilizing the transition state through a stilbene-like structure (Fig. 3).

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We wondered whether we could avoid this competing cyclization by omitting the 4-methoxyphenyl substituent during the synthesis of the pyrazole (Fig. 4). Of course, this approach would entail adding an aryl group later to the completed heterocycle, but this could presumably be done using an appropriate Pd(0)-mediated coupling reaction. This synthetic strategy was also attractive because it allows for the introduction of additional structural diversity into these systems at a late stage, just before phenol deprotection, an attractive feature for combinatorial library expansion.⁵

According to this approach, we were able to successfully synthesize pyrazole 2 as well as analogues 10a-d (Scheme 2). Initial formation of hydrazones 6a-c by reaction of



Scheme 1. Attempted synthesis of MeO-protected pyrazole isomer.



Figure 3. Attempted synthesis of pryazole isomer using an acylationcyclization.



Figure 4. Retro-synthetic strategy for Pd(0)-mediated coupling approach to pyrazoles.

the acetophenone compounds with either the hydrochloride salt of 4-methoxyphenylhydrazine or phenylhydrazine occurred in moderate yields, and as we had hoped, no competing cyclization to the indole occurred with this less stabilized system. Hydrazones 6a-c were then treated with butyllithium to form the corresponding dianion, which was acylated with an alkyl anhydride and then cyclized upon addition of HCl.⁶ The yields shown for pyrazole formation are based on the anhydride and are typical for this reaction. To introduce the last aromatic substituent, the tri-substituted pyrazoles 7a-d were iodinated by treatment with a solution of KI and I_2 ,⁷ and then subjected to Suzuki coupling conditions with either phenylboronic acid or *p*methoxyphenylboronic acid. Initial conditions involved using Pd(PPh₃)₄ as a Pd(0) catalyst and DME/H₂O as solvent (9c 52%, 72 h); however, by using Pd(OAc)₂ as a pre-catalyst and an *n*-PrOH/H₂O solvent mixture, we were able to obtain the tetra-substituted pyrazoles (9a,b,d,e) with slightly improved yields (60–71%) and shorter reaction times (1–14 h).⁸ The protected pyrazole isomers were subsequently demethylated using BBr₃ to afford the desired phenolic products 10a–d and 2.

Comparison of the estrogen receptor binding affinities of the isomeric 1,3,4-triaryl-pyrazoles (Series II) and the original 1,3,5-triaryl-pyrazoles (Series I)

The binding affinities of the pyrazoles for ER were assayed in a competitive radiometric binding assay; this assay has been described previously,⁹ and affinities are expressed as relative binding affinity (RBA) values, where estradiol has an affinity of 100% (Table 1). The 1,3,4-triaryl-pyrazole 2, which is the nitrogen ring isomer of our original 1,3,5-triaryl-pyrazole 1, has a very significant affinity of 5.8%, but is 2.6-fold less than the original pyrazole 1. To determine whether the 1,3,4-triaryl-pyrazole binds to ER in the same mode as pyrazole 1, we investigated the binding affinity of several other 1,3,4-triaryl-pyrazoles (10a–d) and compared their affinities with those of their corresponding 1,3,5-triaryl-pyrazole isomers (11a,b⁴ and 11c,d¹⁰).

When examining the first three sets of isomeric pyrazoles (the other two are discussed below), the pyrazoles in the new isomeric Series II (1,3,4-triaryl-pyrazoles) have somewhat lower affinity than those in the original Series I (1,3,5-triaryl-pyrazoles), but only by an average of 2-fold. Within the original pyrazole series (Series I), certain changes to the peripheral substituents resulted in an increase in RBA. For example, when a hydroxyl group is added to the N(1)-phenyl substituent (i.e. from pyrazole 1 to 11a), a slight increase in affinity occurs. A similar increase in RBA upon hydroxyl substitution is observed in the isomeric Series II, going from pyrazole 2 to 10a.

Previously, we also showed that modification of the alkyl chain at C(4)-position of the initial pyrazole Series I from an ethyl **1** to a propyl substituent **11b** results in a 1.6-fold increase in affinity, indicating a more favorable hydrophobic interaction. However, this favorable interaction ended at propyl, as the *n*-butyl isomer experienced a significant drop in binding affinity.⁴ In Series II, going from R_3 =Et (**2**) to *n*-propyl (**10b**) results in a similar but somewhat larger increase in affinity than it does in Series I (2.7-fold versus 1.6-fold).

Because the structure–binding affinity patterns observed for both pyrazole isomers (Series I and II) are quite similar, we postulated that these distinct core structure pyrazole isomers would be binding in the same orientation in the ER binding pocket.



Scheme 2. Synthesis of 1,3,4-triaryl-5-alkyl-pyrazoles.

 Table 1. Binding affinity data for pyrazole isomers and original pyrazole series

R ₁	R ₂	R ₃	R.4	Compound (I)	RBA (I) ^a	Compound (11)	RBA (II) ^a
11	ОН	Et	OH	1	15.3±3 、	2	5.8±0.3
ΟH	OH	Εt	OH	11a	20.3 ± 3	10a	7.8 ± 4
Н	OH	<i>n</i> -Pr	OH	11b	25 ± 0.01	10b	16.0 ± 1
Н	OH	Et	H	11e	0.46	10c	$0.43 {\pm} 0.07$
H	H	Εt	ОH	11d	0.007	10d	$0.008 {\pm} 0.005$

^aCompetitive radiometric binding assays were done using 10 nM $[^3H]E_2$ as tracer and lamb uterine cytosol as receptor source: for details, see Experimental. Compounds **11a d** were prepared as described elsewhere.^{4,10}

Elucidation of the aryl group in the pyrazole that mimics the A-ring of estradiol: Inferring pyrazole orientation in the ligand binding pocket from the binding affinity of isomeric monophenols

To further support the hypothesis that both pyrazole isomers have the same binding mode and to more confidently establish which phenolic group is playing the role of the A-ring phenol in estradiol, we prepared the individual monophenolic derivatives of pyrazole 1 (Table 1, Series I, 11c and 11d) and pyrazole 2 (Table 1, Series II, 10c and 10d). Systematic phenol deletion is a standard approach that has been used in the past to determine which phenol in an unsymmetrical bisphenolic non-steroidal ER ligand mimics the crucial A-ring phenol of estradiol (E_2).^{11,12} The monophenolic analogue having the highest affinity is then presumed to have preserved the phenol that is acting as the estradiol A-ring mimic, because the hydroxyl substituent at this position is known to be essential for high affinity binding to the ER.¹³ (At the outset, it is of note that removal of one of the phenols from the R₁ position (pyrazole 1 versus 11a in Series I, and pyrazole 2 versus 10a in Series II) has little effect

on binding affinity, suggesting that these rings are not the ones that correspond to the A-ring of estradiol.)

The route in Scheme 2 was used to prepare both the N(1) and the C(4) monophenolic pyrazole Series II analogues 10c and 10d, regioselectively and in good yield. The RBA of pyrazole isomer 10c is 0.43%, whereas that of pyrazole isomer 10d is only 0.008%. When these affinities are compared to that of the bisphenolic pyrazole 2 (5.8%) (Table 1), one can see that removal of the phenolic hydroxyl from the N(1)phenyl ring results in a 725-fold decrease in affinity, whereas removal of the hydroxyl from the C(4) phenyl group results in only a 13-fold decrease. Thus, the 56fold greater decrease in affinity that results from deleting the phenolic hydroxyl from the N(1) phenyl group versus from the C(4) phenyl group strongly suggests that the N(1)-phenolic substituent in the Series II pyrazoles is acting as the A-ring mimic (Fig. 5).

Initial attempts to prepare the C(3) and C(5) monophenol pyrazole analogues for Series I (11c and 11d) regioselectively using the acylation-cyclization strategy were unsuccessful. However, these compounds could successfully be prepared via the corresponding pyrazolines by a new regioselective route (not shown) that will be described elsewhere.¹⁰ The binding affinities for the Series I monophenolic isomers 11d and 11c are 0.007% and 0.46%, respectively. Compared to the corresponding bisphenolic pyrazole 1 (15.3%) (Table 1), a much greater decrease in binding affinity results when the hydroxyl group is removed from C(3) phenol (2185fold, 11d) than from the C(5) phenol (33-fold, 11c). The difference in affinity loss that results from the alternative hydroxy group deletion in this series (i.e. 2185/33- or 66fold) is comparable to that found in Series II (56-fold). This result indicates that in Series I the C(3) phenol is



Preferred Orientation of Phenol in Pyrazoles



Figure 5. Preferred orientation and proposed A-ring mimic for Series I and Series II pyrazoles.

mimicking the A-ring of E_2 . Thus, because the C(3) phenol in Series I and the N(1) phenol in Series II are positioned in a congruent fashion, we believe that the pyrazoles of both Series I and II share a single common binding mode, as illustrated in Figure 5. According to this mode, the other two aryl groups are displayed within a region of the binding pocket resembling the C/D region of E_2 , which is known to tolerate a large number of substituents.¹³

Modeling pyrazole orientation in the ligand binding pocket

In our original study of pyrazole 1, we used molecular modeling to show that this compound could bind in an orientation in which the C(5) phenol mimicked the A-ring of estradiol;³ this is different from that shown in Figure 5. However, in this earlier study, we had based our model on the ER α crystal structure with the antagonist ligand raloxifene.¹⁴ Upon further consideration, we now think that the ER α crystal structure we used previously was not an appropriate one for modeling ligands like pyrazole 1, because unlike raloxifene, these pyrazoles are ER α agonists, not antagonists.^{4,15}

We have recently completed a more extensive modeling of a pyrazole triphenol that corresponds to pyrazole *diphe*nol 11b,⁴ using as a starting point an X-ray structure of ER α complexed with the agonist diethylstilbestrol, a nonsteroidal estrogen agonist with an RBA of 98%.¹⁶ In this study, we considered six possible starting orientations (the three phenols placed in the A-ring pocket and in each case the two alternative orientations of the remaining two phenols that come from rotation about the bond linking the A-ring phenol and the pyrazole core). From this recent study,⁴ the orientation in which the C(3) phenol of the Series I pyrazole mimics the Aring of estradiol gave the lowest energy structure and appeared to be quite reasonable on steric grounds, although it is difficult to rigorously eliminate some of the other possible orientations. This is the orientation shown in Figure 5 for the Series I pyrazole.



Figure 6. A model for pyrazole 10a docked and minimized in the DES-ER α -LBD. The surface of the ligand is shown in yellow, and the solvent accessible surface of ER as purple dots.

In the present study, we have used this approach to examine the six possible binding orientations of the ethyl triphenol Series II pyrazole **10a**. Again, the orientation in which the N(1) phenol is the mimic of the Aring of estradiol (cf. Fig. 5) is well accommodated by the structure and has a reasonable binding energy, although an alternative orientation with the C(4) phenol in this position cannot be ruled out by binding energy considerations. Nevertheless, this latter orientation seems unlikely, given the much higher experimental binding affinity of monophenol **10c** versus **10d** (Table 1).

Thus, on the basis of the experimental determinations of monophenol binding affinities and molecular modeling, we suggest that the preferred binding modes of the pyrazoles in the estrogen receptor correspond to those illustrated in Figure 5. A pictorial representation of pyrazole **10a** in this orientation is shown in Figure 6.

Conclusions

To explore the effect of core structure on the ability of pyrazole ligands to bind to ER, we developed an efficient, regioselective, and flexible route to a new isomeric series of pyrazoles (1,3,4-triaryl-5-alkyl-pyrazoles), and we evaluated their binding affinity to ER. The synthesis relies on the acylation of a hydrazone anion, followed, after cyclization and halogenation, by a Pd(0) Suzuki coupling strategy. We found that the ER accommodates 1,3,4-triaryl-pyrazoles of the new isomeric series (Series II) only slightly less well than the original 1,3,5-triaryl series (Series I), and on the basis of the binding affinity of the corresponding monophenols and molecular modeling studies, both isomers appear to share a common binding mode. In this binding mode, the putative A-ring mimic is the C(3) phenol in Series I and the N(1)phenol in Series II. Thus, as in the case of the imidazoles discussed earlier (Fig. 1), it does appear possible to permute the position of heteroatoms in the azole ring without having a major effect on the ER binding

affinity, provided that the peripheral substituents remain disposed with the same geometry and provided that one remains in the same azole series (i.e. 1,2-azoles [pyrazoles] not 1,3-azoles [imidazoles]). so that compounds with equivalent dipole moments and polarities are being compared. These studies provide additional insight into the design of heterocyclic core structures for the development of high affinity ER ligands by combinatorial methods.

Experimental

General

Melting points were determined on a Thomas-Hoover UniMelt capillary apparatus and are uncorrected. All reagents and solvents were obtained from Aldrich, Fisher or Mallinekrodt. Tetrahydrofuran was freshly distilled from sodium/benzophenone. Dimethylformamide was vacuum distilled prior to use, and stored over 4 Å molecular sieves. *n*-Butyllithium and *t*-butyllithium were titrated with N-pivaloyl-o-toluidine. Et₃N was stirred with phenylisocyanate, filtered, distilled, and stored over 4 Å molecular sieves. All reactions were performed under a dry N₂ atmosphere unless otherwise specified. Reaction progress was monitored by analytical thin-layer chromatography using GF silica plates purchased from Analtech. Visualization was achieved by short wave UV light (254 nm) or potassium permanganate. Flash column chromatography was performed using Woelm 32/63 µm silica gel packing.¹⁷

¹H and ¹³C NMR spectra were recorded on either a Varian Unity 400 MHz or 500 MHz spectrometer using CDCl₃, MeOD or (CD₃)₂SO as solvent. Chemical shifts were reported as parts per million downfield from an internal tetramethylsilane standard (δ 0.0 for ¹H) or from solvent references. NMR coupling constants are reported in Hertz. ¹³C NMR spectra were determined using either the Attached Proton Test (APT) or standard ¹³C pulse sequence parameters. Low resolution and high resolution electron impact mass spectra were obtained on Finnigan MAT CH-5 or 70-VSE spectrometers. Elemental analyses were performed by the Microanalytical Service Laboratory of the University of Illinois. Hydrazones 6a-c are prone to decomposition and therefore could only be stored for 2– 3 days at 0 °C. Once the hydrazone products were confirmed by ¹H NMR they were typically used directly in the acylation cyclization step without further characterization.

Biological procedures: relative binding affinity assay

Ligand binding affinities (RBAs) using lamb uterine cytosol as a receptor source were determined by a competitive radiometric binding assay using 10 nM [³H]estradiol as tracer and dextran-coated charcoal as an adsorbent for free ligand.⁹ All incubations were done at 0 °C for 18 24 h. Binding affinities are expressed relative to estradiol (RBA = 100%) and are reproducible with a coefficient of variation of 0.3.

Molecular modeling of pyrazole 10a

The protocol for modeling followed that recently described for a related pyrazole triphenol.⁴ The starting conformation for pyrazole **10a** used for receptor docking studies was generated from a random conformational search performed using the MMFF94 force field as implemented in Sybyl 6.6. The resulting lowest energy conformer was then used for docking studies. Charge calculations were determined using the MMFF94 method and molecular surface properties displayed using MOLCAD module in Sybyl 6.6.

Pyrazole 10a, generated as noted above, was pre-positioned in the DES-ER α -LBD crystal structure¹⁶ using a least squares multifitting of select atoms within the DES ligand. Once pre-positioned, DES was deleted and ligand 10a was optimally docked in the ER α binding pocket in six orientations (as specified in the text) using the Flexidock routine within Sybyl (Tripos). Both hydrogen-bond donors and acceptors within the pocket surrounding the ligand (Glu₃₅₃, Arg₃₉₄, and His₅₂₄), the ligand itself and select torsional bonds were defined. The docked receptor ligand complexes from Flexidock then underwent a three step minimization. First, nonring torsional bonds of the ligand were minimized in the context of the receptor using the torsmin command. This was followed by minimization of the side chain residues within 8 Å of the ligand, while holding the backbone and residues Glu₃₅₃ and Arg₃₉₄ fixed. A final third minimization of both the ligand and receptor was conducted using the Anneal function (hot radius 8 Å, interesting radius 16 Å from pyrazole **10a**) to afford the final model.

Chemical syntheses

5-Ethyl-1,4-bis-(4-methoxyphenyl)-3-phenyl-1H-pyrazole (2). A stirred solution of **9e** (46 mg, 0.16 mmol) in CH₂Cl₂ (15 mL) was treated with BBr₃ (1.62 mL, 1.62 mmol) according to the general demethylation procedure. After work up and an SiO₂ plug (30% EtOAc/hexanes) a white solid was isolated (40 mg, 94%): mp 220 225 °C; ¹H NMR (MeOD- d_4 , 400 MHz) δ 0.90 (t, 3H, J = 7.5), 2.62 (q, 2H, J = 7.5), 6.79 (XX' of AA'XX', 2H, $J_{AX} = 8.3$, $J_{XX} = 2.3$), 6.94 (XX' of AA'XX', 2H, $J_{AX} = 8.6$, $J_{XX} = 2.7$), 7.04 (AA' of AA'XX', 2H, $J_{AX} = 8.2$, $J_{AA'} = 2.4$), 7.20–7.27 (m, 2H), 7.34 (AA' of AA'XX', 2H, $J_{AX} = 8.8$, $J_{AA'} = 2.6$), 7.37–7.44 (m, 3H); ¹³C NMR (MeOD- d_4 , 100 MHz) δ 12.4, 17.2, 114.8, 115.2, 118.5, 124.3, 126.9, 127.4, 127.5, 127.6, 131.0, 131.0, 132.9, 144.1, 148.9, 156.2, 157.8; MS (EI, 70 eV) *m/z* (relative intensity, %): 356 (M⁺, 100); Anal calcd for C₂₃H₂₀ N₂O₂·H₂O: C, 73.78; H, 5.92; N, 7.48. Found: C, 73.82; H, 5.95; N, 7.69.

4'-Methoxyacetophenone-4-methoxyphenylhydrazone (6a). An EtOH (40 mL) solution of 4-methoxyacetophenone (1g, 6.67 mmol), sodium acetate (1.09 g, 13.34 mmol), and 4-methoxyphenylhydrazine hydrochloride (1.74 g, 10.0 mmol) was heated to 80 °C for 3.0 h. The reaction mixture was cooled to room temperature and concentrated under reduced pressure. The residue was dissolved in ethyl acetate (30 mL) and washed with water (40 mL) and brine (2×40 mL). After drying over MgSO₄ and solvent concentration a red and white heterogeneous solid formed. The product was collected by filtration and rinsed with cold ethanol (20 mL) to afford **6a** as a white solid (1.18 g, 66%) which was used directly in the next step.

Acetophenone-4-methoxyphenylhydrazone (6b). A solution of acetophenone (250 mg, 2.00 mmol) and 4-methoxyphenylhydrazine hydrochloride (349 mg, 2.00 mmol) was reacted similarly to conditions used for **6a** to afford a red and white heterogeneous sold. Cold ethanol (20 mL) was added and the remaining solid collected via vacuum filtration to afford **6b** as a white solid (1.41 g, 59%) which was used directly in the next step.

Acetophenonephenylhydrazone (6c). A solution of acetophenone (1.8 g, 15 mmol), phenylhydrazine hydrochloride (2.17 g, 15 mmol), and sodium acetate (1.23 g, 15.0 mmol), in anhydrous ethanol (40 mL), was reacted similarly to conditions used for **6a** to afford a yellow and white heterogeneous solid. Cold ethanol (20 mL) was added to the solid and the product collected via vacuum filtration to afford **6c** as a white solid (1.67 g, 53%) which was used directly in the next step.

General acylation-cyclization procedure (7a-d). To a stirred solution of hydrazone (3.05 mmol) in THF (10 mL) at 0°C was added 1.29 M BuLi (4.73 mL, 6.10 mmol) dropwise. The resulting deep red solution was allowed to stir at 0°C for 0.25 h, then room temperature for 0.25 h, and re-cooled to 0 °C, whereupon a THF solution (2.0 mL) of the appropriate alkyl anhydride (1.53 mmol) was added dropwise. This mixture was stirred at 0 °C for 0.25 h, then treated with 3 M HCl (6 mL, 18 mmol) and refluxed for 1.5 h. The biphasic mixture was cooled to room temperature and the agueous layer separated and neutralized with saturated NaHCO₃. The neutralized solution was extracted with ethyl acetate $(3 \times 20 \text{ mL})$ and the organic layers combined with the THF layer from the reaction mixture. The final organic layers were washed with satd NaHCO₃ (3×20 mL), dried over MgSO₄, and concentrated to afford a red oil. The crude oil was purified by flash chromatography.

5-Ethyl-1,3-bis-(4-methoxyphenyl)-1H-pyrazole (7a). Following the general procedure above hydrazone **6a** (824 mg, 3.05 mmol) was acylated with propionic anhydride (198 mg, 0.20 mL, 1.53 mmol) and cyclized with HCl. Flash chromatography (25% EtOAc/hexanes) afforded an inseparable mixture of 4-methoxyacetophenone and 7a (474 mg). The phenone was reduced by the dropwise addition of a solution of NaBH₄ (27.3 mg, 0.72) in H₂O (5 mL) to a solution of the product mixture in ethanol (5 mL). The solution was stirred for 18 h at room temperature and poured over 1 M HCl (20 mL). Upon product isolation (EtOAc, satd NaHCO₃, brine) and solvent removal under reduced pressure an orange oil was isolated, which upon addition of 25% EtOAc/hexanes resulted in selective crystallization of pyrazole 7a as fine white crystals (200 mg, 44%): ¹H NMR (CDCl₃, 400 MHz) δ 1.24 (t, 3H, J=7.6), 2.64 (q, 2H, J=7.5), 3.83 (s, 3H) 6.45 (s, 1H), 3.85 (s, 3H), 6.98 (XX' of AA'XX', 2H, J_{AX} =8.9, J_{XX} =2.4), 7.09 (XX' of AA'XX', 2H, J_{AX} =9.0, J_{XX} =2.7), 7.39 (AA' of AA'XX', 2H, J_{AX} =8.9, $J_{AA'}$ =2.7), 7.79 (AA' of AA'XX', 2H, J_{AX} =8.9, $J_{AA'}$ =2.8); ¹³C NMR (CDCl₃, 100 MHz) δ 13.0, 19.6, 55.1, 55.4, 101.2, 113.7, 114.1, 126.2, 126.8, 126.8, 133.0, 146.5, 150.8, 158.9, 159.1.

1-(4-Methoxyphenyl)-3-phenyl-5-propyl-1H-pyrazole (7b). Following the general procedure above hydrazone **6a** (1.37 g, 5.71 mmol) was acylated with butyric anhydride (452 mg, 0.47 mL, 2.86 mmol) and cyclized with HCl. Flash chromatography (30% EtOAc/hexanes) afforded the title compound as a yellow oil (710 mg, 42%): ¹H NMR (CDCl₃, 400 MHz) & 0.96 (t, 3H, J=7.5) 1.66 (sext, 2H, J=7.5), 2.61 (t, 2H, J=7.7), 3.85 (s, 3H), 6.54 (s, 1H), 6.99 (XX' of AA'XX', 2H, J_{AX} =8.7, J_{XX} =2.7), 7.31 (app tt, 1H, J=7.3, 1.5), 7.38-7.43 (m, 4H), 7.89 (AA' of AA'XX', 2H, J_{AX} =7.2, $J_{AA'}$ =1.3); ¹³C NMR (CDCl₃, 100 MHz) & 13.7, 21.9, 27.1, 55.4, 102.2, 114.1, 125.5, 126.9, 127.5, 128.4, 132.9, 133.4, 145.2, 150.9, 159.1; HRMS (EI, M⁺) calcd for C₁₉H₂₀N₂O: 292.1576. Found: 292.1569.

5-Ethyl-1-(4-methoxyphenyl)-3-phenylpyrazole (7c). Following the general procedure above hydrazone 6b (4.50 g, 18.75 mmol) was acylated with propionic anhydride (1.22 g, 1.20 m L, 9.38 mmol) and cyclized with HCl. Upon solvent removal a crude orange solid was isolated. The crude solid was recrystallized from 20% ethyl acetate/hexanes to afford the title compound as pale yellow needles (1.82g, 35%): mp 70-75°C; ¹H NMR (CDCl₃, 400 MHz) δ 1.25 (t, 3H, J=7.5), 2.65 (q, 2H, J=7.5), 6.53 (s, 1H), 3.86 (s, 3H), 6.99 (AA'XX', 2H, J=8.6, 2.6), 7.28–7.33 (m, 1H), 7.37–7.43 (m, 4H), 7.85-7.90 (m, 2H); ¹³C NMR (CDCl₃, 125 MHz) δ 13.0, 19.6, 55.4, 101.6, 114.2, 125.5, 126.8, 127.5, 128.4, 132.9, 133.3, 146.6, 151.0, 159.0; MS (EI, 70 eV) m/z (relative intensity, %): 278 (M⁺, 100). Anal. calcd for $C_{18}H_{18}$ N₂O: C, 77.67; H, 6.52; N, 10.06. Found: C, 77.62; H, 6.49; N, 10.05.

1,3-Diphenyl-5-propyl-1H-pyrazole (7d). Following the general procedure above hydrazone **6c** (1.30 g, 6.19 mmol) was acylated with propionic anhydride (403 mg, 0.40 mL, 3.10 mmol) and cyclized with HCl. Flash chromatography (20% diethyl ether/hexanes) afforded the title compound as a yellow oil (600 mg, 40%): ¹H NMR (CDCl₃, 500 MHz) δ 1.28 (t, 3H, *J*=7.4), 2.72 (q, 2H, *J*=7.4), 6.58 (s, 1H), 7.32–7.53 (m, 8H), 7.91 (d, 2H, *J*=7.5); ¹³C NMR (CDCl₃, 100 MHz) δ 13.2, 19.9, 102.3, 125.7, 127.8, 127.9, 128.8, 129.1, 133.4, 140.0, 146.7, 151.5; HRMS (EI, M⁺) calcd for C₁₇H₁₆N₂: 248.1313. Found: 248.1312.

General iodination procedure (8a–d). To a refluxing solution of pyrazole (0.59 mmol) and sodium acetate (107 mg, 1.13 mmol) in H₂O (4.00 mL) was added a solution of KI (591 mg, 3.56 mmol) and I₂ (301 mg, 1.19 mmol) in H₂O (4 mL) dropwise. The resulting dark brown solution was allowed to reflux for 3 h and then was cooled to room temperature and the product extracted with diethyl ether (3×25 mL). The organic layers were combined, washed with NaS₂O₂ ($3 \times 20 \text{ mL}$), NaHCO₃ ($2 \times 20 \text{ mL}$), brine ($2 \times 20 \text{ mL}$) and then dried over MgSO₄ and concentrated under reduced pressure to afford the crude iodo-pyrazoles. The products were purified by recrystallization from hexanes or by flash chromatography on silica gel.

5-Ethyl-4-iodo-1,3-bis-(4-methoxyphenyl)-1H-pyrazole (8a). Pyrazole 7a (153 mg, 0.59 mmol) was iodinated according to the general procedure above. A crude tan solid was isolated and recrystallized from hexanes to afford the title compound as a white solid (224 mg, 87%): ¹H NMR (CDCl₃, 400 MHz) δ 1.13 (t, 3H, J=7.4), 2.73 (q, 2H, J=7.5), 3.84 (s, 3H), 3.85 (s, 3H), 6.94 7.02 (m, 4H), 7.36 (AA'XX', 2H, J=9.0, 2.7), 7.83 (AA'XX', 2H, J=8.8, 2.5); ¹³C NMR (CDCl₃, 125 MHz) δ 13.1, 20.1, 55.1, 55.4, 60.1, 113.4, 114.2, 125.4, 126.9, 129.4, 132.6, 147.2, 151.4, 159.4, 159.5.

4-Iodo-1-(4-methoxyphenyl)-3-phenyl-5-propylpyrazole (**8b**). Pyrazole **7b** (660 mg, 2.26 mmol) was iodinated according to the general procedure above. A crude oil was isolated and purified by flash chromatography (25% ethyl acetate/hexanes) to afford a mixture of unreacted pyrazole **7b** and product **8b** as a pale-yellow oil (yield as determined by NMR: 65%; resonances listed for product only): ¹H NMR (CDCl₃, 400 MHz) δ 0.90 (1, 3H, J=7.4) 1.56 (sext, 2H, J=7.6), 2.70 (1, 2H, J=7.8), 3.86 (s, 3H), 6.99 (XX' of AA'XX', 2H, J_{AX} = 8.7, $J_{XX'}$ =2.8), 7.90 (AA' of AA'XX', 2H, J_{AX} = 8.3, $J_{AA'}$ =1.6), 7.26–7.46 (m, 5H); ¹³C NMR (CDCl₃, 100 MHz) δ 13.8, 21.9, 28.4, 55.4, 60.9, 114.2, 127.1, 128.0, 128.0, 128.2, 132.8, 146.2, 151.5, 159.5; HRMS (EI, M⁺) calcd for C₁₉H₁₉N₂OI: 418.0542. Found: 418.0550.

5-Ethyl-4-iodo-1-(4-methoxyphenyl)-3-phenyl-1H-pyrazole (8c). Pyrazole **7c** (500 mg, 1.80 mmol) was iodinated according to the general procedure above to afford the title compound as a white solid (603 mg, 83%): mp 70-73 °C; ¹H NMR (CDCl₃, 500 MHz) δ 1.14 (t, 3H, J=7.5), 2.74 (q, 2H, J=7.5), 3.86 (s, 3H), 6.99 (AA' XX', 2H, J=8.6, 2.9), 7.34–7.47 (m, 5H), 7.86–7.92 (m, 2H); MS (EI, 70 eV) m/z (relative intensity, %): 404 (M⁺, 100). Anal. calcd for C₁₈H₁₇N₂OI: C, 53.48; H, 4.24; N, 6.93. Found: C, 53.66; H, 4.24; N, 6.89.

5-Ethyl-4-iodo-1,3-diphenyl-1H-pyrazole (8d). Pyrazole **7d** (248 mg, 1.13 mmol) was iodinated according to the general procedure above to afford a crude orange solid. Recrystallization from hexanes afforded **8d** as small white crystals (312 mg, 74%): mp 82–85 °C; ¹H NMR (CDCl₃, 500 MHz) δ 1.61 (t, 3H, *J*=7.5), 2.78 (q, 2H, *J*=7.5), 7.35 7.55 (m, 8H), 7.86–7.91 (m, 2H); ¹³C NMR (CDCl₃, 125 MHz) δ 13.3, 20.3, 61.1, 125.6, 128.2, 128.3, 128.4, 128.6, 129.3, 132.9, 139.9, 147.3, 152.1; HRMS (EI, M⁺) calcd for C₁₇H₁₅N₂I: 374.0280. Found: 374.0277.

General procedure for Suzuki coupling (9a,b,d,e). To a stirred solution of iodo-pyrazole (0.43 mmol) in *n*-propanol (2 mL) were added the appropriate boronic acid (0.45 mmol), Pd(OAc)₂ (3 mg, 0.013 mmol), PPh₃ (10 mg, 0.039 mmol), 2 M Na₂CO₃ (0.47 mL, 0.95 mmol), and

 H_2O (0.5 mL). The heterogeneous solution was heated to reflux for 5 h, cooled to room temperature, and filtered through Celite (EtOAc 4×5 mL). The filtrate was concentrated and partitioned in diethyl ether and water and the aqueous layer separated and extracted twice more with ether. The combined organic layers were washed with brine, dried over MgSO₄ and concentrated under reduced pressure. The tetrasubstituted pyrazoles were subsequently purified by flash chromatography.

5-Ethyl-1,3,4-tris-(4-methoxyphenyl)-1H-pyrazole (9a). Iodo-pyrazole 8a (187 mg, 0.43 mmol) was coupled with 4-methoxyphenylboronic acid (68 mg, 0.45 mmol) according to the general procedure above to afford a crude brown oil. Flash chromatography (40% ethyl acetate/hexanes) furnished the title compound as a clear oil (124 mg, 70%): ¹H NMR (CDCl₃, 400 MHz) δ 0.93 (t, 3H, J=7.5), 2.63 (q, 2H, J=7.5), 3.77 (s, 3H), 3.85 (s, 3H), 3.87 (s, 3H), 6.78 (AA'XX', 2H, J=8.9, 2.5), 6.93 (AA'XX', 2H, J=8.7, 2.5), 7.01 (AA'XX', 2H, J=8.9, 2.6), 7.20 (AA'XX', 2H, J=8.6, 2.5), 7.43(AA'XX', 2H, J=8.8, 2.4), 7.46 (AA'XX', 2H, J=8.9, 2.6); ¹³C NMR (CD₃OD, 100 MHz) δ 13.7, 17.8, 54.9, 55.0, 55.4, 113.3, 113.8, 114.1, 125.9, 126.4, 127.1, 127.4, 128.9, 131.3, 133.1, 143.5, 148.7, 158.3, 158.7, 159.1; HRMS (EI, M^+) calcd for $C_{26}H_{26}N_2O_3$: 414.1943. Found: 414.1937.

1,4-Bis-(4-methoxyphenyl)-3-phenyl-5-propyl-1H-pyrazole (9b). Iodo-pyrazole **8b** (300 mg, 0.72 mmol) was coupled with 4-methoxyphenylboronic acid (115 mg, 0.75 mmol) according to the general procedure above. Flash chromatography (25% EtOAc/hexanes) afforded the title compound as a glassy solid (203 mg, 71%): ¹H NMR (CDCl₃, 500 MHz) δ 0.70 (t, 3H, *J*=7.4), 1.31 (sext, 2H, *J*=7.6), 2.59 (t, 2H, *J*=7.9), 3.84 (s, 3H), 3.86 (s, 3H), 6.92 (AA'XX', 2H, *J*=8.4, 2.4), 7.00 (AA'XX', 2H, *J*=8.5, 2.5), 7.16-7.27 (m, 5H), 7.45 (AA'XX', 2H, *J*=8.7, 2.4), 7.48-7.54 (m, 2H); ¹³C NMR (MeOD-*d*₄, 100 MHz) δ 13.8, 22.2, 26.5, 55.2, 55.5, 113.9, 114.3, 114.6, 126.4, 127.2, 127.3, 127.9, 128.1, 131.5, 133.3, 133.4, 142.6, 149.0, 158.5, 159.3; HRMS (EI, M⁺) calcd for C₂₆H₂₇N₂O₂: 398.1994. Found: 398.1991.

5-Ethyl-1,4-bis-(4-methoxyphenyl)-3-phenyl-1H-pyrazole (9c). To a degassed solution of 2 M Na_2CO_3 (0.55 mL, 1.09 mmol; 0.5 mL H₂O), and DME (3.0 mL) was added 4-methoxyphenylboronic acid (83 mg, 0.55 mmol) and pyrazole 8c (200 mg, 0.45 mmol). To this solution Pd(Ph₃P)₄ (27 mg, 0.025 mmol) was added and the mixture heated to 80 °C for 72 h. The reaction mixture was then cooled to room temperature and filtered through Celite. The filtrate was transferred to a separatory funnel and the aqueous layers extracted with $Et_2O(3 \times 5 mL)$. The organic layers were combined and concentrated under reduced pressure. Flash chromatography (25% EtOAc/hexanes) afforded 9c as a white solid (100 mg, 53%): ¹H NMR (CDCl₃, 400 MHz) δ 0.93 (t, 3H, J = 7.5), 2.64 (q, 2H, J = 7.6), 3.85 (s, 3H), 3.86 (s, 3H), 6.93 (AA'XX', 2H, J=8.7, 2.4), 7.00 (AA'XX', 2H, J=9.2, 2.7), 7.18-7.27 (m, 5H), 7.46 (AA'XX', 2H, J = 9.0, 2.7), 7.49-7.53 (m, 2H); HRMS (EI, M⁺) calcd for C₂₅H₂₄N₂O₃: 384.1834. Found: 384.1838.

5-Ethyl-1-(4-methoxyphenyl)-3,4-diphenyl-1H-pyrazole (9d). Iodo-pyrazole 8c (400 mg, 0.99 mmol) was coupled with phenylboronic acid (144 mg, 1.04 mmol) according to the general procedure above to afford a crude brown oil. Flash chromatography (15% THF/ hexanes) furnished the product and a mixture of the product and unreacted starting material. The mixture was treated to a second column under the same conditions to afford additional 9d (50 mg combined, 14% isolated yield; 217 mg for remaining mixture, 60% total yield as determined by NMR): ¹H NMR (MeOD- d_4 , 500 MHz) δ 0.94 (t, 3H, J=7.4), 2.66 (q, 2H, J=7.5), 7.02 (AA'XX', 2H, J = 8.9, 2.7), 7.17–7.53 (m, 12H); ¹³C NMR (McOD-d₄, 100 MHz) δ 13.7, 17.8, 55.4, 114.1, 118.7, 126.6, 127.2, 127.9, 127.9, 128.3, 130.2, 132.8, 133.1, 133.9, 143.7, 148.8, 159.3.

5-Ethyl-4-(4-methoxyphenyl)-1,3-diphenyl-1H-pyrazole (9e). Iodo-pyrazole 8d (244 mg, 0.65 mmol) was coupled with 4-methoxyphenylboronic acid (105 mg, 0.69 mmol) according to the general procedure above to afford a yellow oil. Flash chromatography (15% EtOAc/hexanes) was performed to afford a pale yellow oil which was then recrystallized from hexanes to afford the title compound as white crystals (160 mg, 70%): ¹H NMR (McOD- d_4 , 500 MHz) δ 0.94 (t, 3H, J=7.5), 2.70 (q, 2H, J=7.5), 3.85 (s, 3H), 6.94 (AA'XX', 2H, J=8.8, 2.6), 7.19–7.28 (m, 5H), 7.43 (app tt, 1H, J = 7.4, 1.7), 7.48-7.55 (m, 4H), 7.55-7.60 (m, 2H); ¹³C NMR (McOD-d₄, 125 MHz) δ 13.7, 17.9, 55.1, 113.8, 118.8, 125.7, 126.1, 127.2, 127.8, 127.9, 129.0, 131.3, 133.2, 140.0, 143.6, 149.3, 158.4.

General procedure for demethylation using BBr₃ (2, 10a-d). To a stirred solution of pyrazole (9a-e, 0.28 mmol) in CH_2Cl_2 (10 mL) at -78 °C was added BBr₃ (2.82 mL. 2.82 mmol) dropwise as a 1 M solution in CH_2Cl_2 . The mixture was allowed to warm to room temperature and stirred for 3 h. The reaction was quenched at 0 °C by the addition of H_2O (10 mL). The resulting solid was solubilized by the addition of ethyl acetate and the resulting biphasic solution transferred to a separatory funnel and the aqueous layer isolated. The aqueous layer was acidified with 3 M HCl (3 mL) and extracted with ethyl acetate (2×15 mL). The combined organic layers were washed with brine and dried over MgSO₄. The solvent was removed under reduced pressure and the phenolic products purified by flash chromatography and/or crystallization.

5-Ethyl-1,3,4-tris-(4-hydroxyphenyl)-1H-pyrazole (10a). A stirred solution of 9a (46 mg, 0.11 mmol) in CH_2Cl_2 (5 mL) was treated with BBr₃ (1.10 mL, 1.10 mmol) according to the general demethylation procedure. After work up a reddish brown oil isolated. The oil was dissolved in methanol and concentrated under reduced pressure. No effort was made to remove trace amounts of methanol remaining in the sample. The residue was triturated by adding CH₂Cl₂ dropwise until a precipitate formed. The precipitate was isolated by vacuum filtration to afford **10a** as a white solid (40 mg, 93%): ¹H NMR (MeOD- d_4 , 400 MHz) δ 0.89 (t, 3H, J = 7.6) 2.59 (q, 2H, J=7.5), 6.65 (d, 2H, J=8.4), 6.79 (d, 2H,

J = 8.4), 6.93 (d, 2H, J = 8.8), 7.03 (d, 2H, J = 8.8), 7.21 (d, 2H, J=8.4), 7.32 (d, 2H, J=8.8); ¹³C NMR $(McOD-d_4, 100 MHz) \delta 12.4, 17.2, 114.3, 114.8, 115.1,$ 118.0, 124.2, 124.6, 127.4, 128.9, 131.0, 131.2, 143.9, 149.1, 156.0, 156.6, 157.7; HRMS (EI, M⁺) calcd for C₂₃H₂₀N₂O₃: 372.1474. Found: 372.1468.

1,4-Bis-(4-hydroxyphenyl)-3-phenyl-5-propyl-1H-pyrazole (10b). A stirred solution of 9b (100 mg, 0.25 mmol) in CH_2Cl_2 (10 mL) was treated with BBr₃ (2.50 mL, 2.50 mmol) according to the general demethylation procedure. After work up a red solid was isolated. The solid was recrystallized from 10% CH₃OH/CHCl₃ to afford 10b as small white crystals (75 mg, 81%): mp 233-236°C; ¹H NMR (MeOD-*d*₄, 400 MHz) δ 0.67 (t, 3H, J = 7.3), 1.29 (sext, 2H, J = 7.5), 2.59 (t, 2H, J = 7.9), 6.79 (AA'XX', 2H, J=8.4, 2.2), 6.94 (AA'XX', 2H, J=8.6, 2.5), 7.03 (AA'XX', 2H, J=8.3, 2.1), 7.20–7.26 (m, 3H), 7.31–7.43 (m, 4H); ^{13}C NMR (MeOD- d_4 , 100 MHz) δ 12.4, 21.5, 25.9, 114.9, 115.2, 119.1, 124.4, 127.0, 127.4, 127.5, 127.6, 131.1, 131.2, 132.9, 142.8, 148.9, 156.2, 157.8; HRMS (EI, M⁺) calcd for C₂₄H₂₂N₂O₂: 370.1681. Found: 370.1685.

5-Ethyl-1-(4-hydroxyphenyl)-3,4-diphenyl-1H-pyrazole (10c). A stirred solution of 9c (37 mg, 0.10 mmol) in CH_2Cl_2 (5 mL) was treated with BBr₃ (1.00 mL, 1.00 mmol) according to the general demethylation procedure. After work up a brown oil was afforded. The crude oil was purified by flash chromatography (5% CH₃OH/CH₂Cl₂) to afford **10c** as a white solid (30 mg, 86%): mp 210–220 °C; ¹H NMR (MeOD-*d*₄, 400 MHz) δ 0.90 (t, 3H, J=7.60 Hz), 2.65 (q, 2H, J=7.4), 6.95 (AA'XX', 2H, J=9.1, 2.7), 7.19-7.26 (m, 5H), 7.31-7.42 (m, 7H); 13 C NMR (MeOD- d_4 , 100 MHz) δ 13.8, 17.8, 116.5, 118.8, 126.8, 127.5, 127.6, 128.1, 128.1, 128.5, 130.3, 132.1, 132.9, 133.9, 144.2, 149.1, 156.5; HRMS (EI, M^+) calcd for $C_{23}H_{20}N_2O$: 340.1576. Found: 340.1576.

5-Ethyl-4-(4-hydroxyphenyl)-1,3-diphenyl-1H-pyrazole (10d). A stirred solution of 9e (100 mg, 0.28 mmol) in CH_2Cl_2 (10 mL) was treated with BBr_3 (2.82 mL, 2.82 mmol) according to the general demethylation procedure. After work up a crude brown solid was isolated. The solid was purified by flash chromatography (5% CH₃OH/CH₂Cl₂) to afford **10d** as a white solid (89 mg, 93%): ¹H NMR (MeOD- d_4 , 500 MHz) δ 0.93 (t, 3H, J=7.5), 2.68 (q, 2H, J=7.6), 6.78 (XX' of XX'AA', 2H, $J_{AX} = 8.4, J_{XX'} = 2.5$, 7.10 (AA' of AA'XX', 2H, $J_{AX} = 8.4, J_{AA'} = 2.5$), 7.21–7.28 (m, 3H), 7.40–7.52 (m, 6H), 7.53–7.58 (m, 2H); ¹³C NMR (MeOD- d_4 , 100 MHz) δ 13.7, 17.9, 115.6, 119.1, 125.8, 125.9, 127.4, 128.0, 128.1, 128.2, 129.2, 131.5, 133.0, 139.8, 143.9, 149.5, 154.9; HRMS (EI, M^+) calcd for $C_{23}H_{20}N_2O$: 340.1576. Found: 340.1571.

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Triarylpyrazoles with Basic Side Chains: Development of Pyrazole-Based Estrogen Receptor Antagonists

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Abstract—Recently, we developed a novel triaryl-substituted pyrazole ligand system that has high affinity for the estrogen receptor (ER) (Fink, B. E.; Mortenson, D. S.; Stauffer, S. R.: Aron, Z. D.; Katzenellenbogen, J. A. *Chem. Biol.* **1999**, *6*, 205). Subsequent work has shown that some analogues in this series are very selective for the ER α subtype in terms of binding affinity and agonist potency (Stauffer, S. R.; Coletta, C. J.; Tedesco, R.; Sun, J.; Katzenellenbogen, J. A. *J. Med. Chem.* **2000**, submitted). We now investigate how this pyrazole ER agonist system might be converted into an antagonist or a selective estrogen receptor modifier (SERM) by incorporating a basic or polar side chain like those typically found in antiestrogens and known to be essential determinants of their mixed agonist/antagonist character. We selected an *N*-piperidinyl-ethyl chain as a first attempt, and introduced it at the four possible sites of substitution on the pyrazole core structure to determine the orientation that the pyrazole might adopt in the ER ligand binding pocket. Of these four, the C(5) piperidinyl-ethoxy-substituted pyrazole **5** had by far the highest affinity. Also, it bound to the ER subtype alpha (ER α) with 20-fold higher affinity than to ER β . In cell-based transcription assays, pyrazole **5** was an antagonist on both ER α and ER β , and it was also more potent on ER α . Based on structure-binding affinity relationships and on molecular modeling studies of these pyrazoles in a crystal structure of the ER α -raloxifene complex, we propose that pyrazoles having a basic substituent on the C(5) phenyl group adopt a binding mode that is different from that of the pyrazole agonists that lack this group. The most favorable orientation appears to be one which places the N(1) phenol in the A-ring binding pocket so that the basic side chain can adopt an orientation similar to that of the basic side chain of raloxifene. \bigcirc 2000 Elsevier Science Ltd. All rights reserved.

Introduction

The term SERM, or selective estrogen receptor modifier, is used to describe ligands for the estrogen receptor (ER) that display an interesting balance of agonist and antagonist activity that can vary from tissue to tissue.¹ SERMs have particular utility in menopausal hormone replacement and in the treatment and prevention of breast cancer, although the pharmacological profiles of the currently available agents are not optimal.

SERMs typically consist of a non-steroidal core structure onto which is attached a side chain bearing a basic or polar function. X-ray crystal structures of the ER ligand binding domain (LBD) complexed with the SERMs raloxifene² or hydroxytamoxifen³ show that the large side chains of these ligands displace the C-terminal helix (helix-12) from the position it normally occupies in ER-LBD complexes with agonists. In this antagonist conformation, helix-12 moves into a position where it occludes a hydrophobic groove on the surface of the receptor that normally functions as a tethering site for the interaction of ER with coactivator proteins, an interaction that is important in mediating agonist activity.^{3,4}

It is clear, however, that more subtle changes in the conformation of ER must underlie some of the pharmacological differences among various SERMs.^{5,6} Recently, conformation-sensitive peptides have been used to discriminate among these different ligandinduced ER conformations.^{7–9} Nevertheless, it is not completely clear how the ligand structure of the various SERMs determines ER conformation, at least at this level of detail. In addition, differential interaction of SERMs with the two ER subtypes, ER α and ER β , might also underlie certain aspects of their tissue-selective activity.^{10–12}

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In recent studies of novel, heterocyclic non-steroidal estrogens amenable to combinatorial assembly and library synthesis, we identified tetrasubstituted pyrazoles as a system that afforded high affinity ligands for ER.¹³ Subsequently, we used parallel synthesis, as well as directed synthesis, to characterize structure affinity relationships in this series.^{14,15} In the process, we found several high affinity compounds, and in particular, some that show very high binding affinity and transcriptional efficacy for the ER alpha subtype $(ER\alpha)$.¹⁵ Although some of these compounds were also agonists on the ER beta subtype (ER β), their potency and binding affinity on this ER subtype was much lower.¹⁵ The high difference in ER-subtype binding affinities shown by some of these pyrazoles raised an interesting prospect that this system might be a favorable starting point for the development of ERa potency-selective antagonists or SERMs. This would depend, of course, on whether these ligands could be converted to antagonists by appropriate substitution of basic or polar side chains.

In this report, we describe the preparation of several analogues of an ER α high-affinity pyrazole which embody the type of basic or polar side chain that is typically found in SERMs. We evaluate how this substitution is tolerated in terms of ER binding affinity, and in some selected cases, we show it affects the transcriptional efficacy and potency of these ligands. We also develop a model for how these basic side chain-substituted pyrazoles are likely to be orientated in the ligand-binding pocket of ER.

Results and Discussion

An example of a favorable pattern of substitution for high affinity ER pyrazoles ligands consists of three phenols at the 1-, 3- and 5-positions and an alkyl group at the 4-position, and is shown in Figure 1 (pyrazole 1).¹³ ¹⁶ The piperidinyl-ethyl substituent, which is widely represented in many SERM classes (see, for example, Ralox-ifene, Figure 1),^{1,17} was chosen in this study as a typical basic side chain that might be expected to confer the characteristic mixed agonist/antagonist activity of these compounds. The question at hand, however, is at which position on a pyrazole such as 1 should this basic substituent be attached. Molecular modeling studies using the LBD crystal structures provided a starting point towards answering this question.

From published crystal structures, it is known that the ER-LBD adopts a different conformation when complexed with agonists (E₂, DES) versus antagonists (hydroxytamoxifen, raloxifene).^{2,3} In our initial molecular modeling studies, we examined various binding modes for the pyrazole core structure (i.e. without a basic side chain) based on the larger ligand binding pocket in the ER-LBD -raloxifene (antagonist) complex.¹³ However, in a subsequent publication,¹⁵ knowing now that these pyrazoles were agonists, we reexamined our initial predictions, this time using the ER-LBD DES (agonist) structure. As a result of this additional modeling work, as well as further structure activity relationship studies we have done in this series of compounds,¹⁵ we now believe that these core



Figure 1. Pyrazole triol core structure (1) and basic side chain-containing pyrazoles (2 5).

pyrazoles, which behave as agonists, bind in an orientation in which the C(3) phenol plays a role analogous to the critical A-ring of estradiol.¹⁵ We have illustrated this binding orientation with the relative positions we have chosen for pyrazole I and estradiol in Figure 1. The orientation of the SERM raloxifene relative to estradiol, known from X-ray crystallography,² is also displayed in Figure 1. In this orientation, the benzothiophene ring system of raloxifene mimics the AB ring system of estradiol, so that the basic side chain is directed roughly in the estradiol 11 β direction (Fig. 1), where it extends outward to displace helix-12, as noted above.²

On the basis of our current model, we might have considered it unlikely that the pyrazole ligand could even accommodate a basic side chain. When bound with the ER-LBD in the agonist conformation, there is no position on the pyrazole analogue where such a side chain could be disposed so as to occupy a region of the ligand binding pocket that is normally occupied by this group in other SERMs (i.e. the estradiol 11β region, Figure 1).^{2,3} However, because of the near symmetry of these triaryl pyrazoles and the known flexibility of the ligand-binding pocket of the ER,^{2.3} we were not convinced that the binding orientation preferences of the smaller core pyrazole agonist ligands would be retained. The additional bulk and polarity of the basic side chain that we were adding to engender antagonist activity might cause the ligand to adopt an entirely different orientation in the binding pocket.

Therefore, as an initial effort to develop pyrazole-based SERMs, we prepared four pyrazoles in which the basic piperidinyl-ethyl side chain is attached at four different positions (2–5, Fig. 1), and we determined the binding affinity of these derivatives for the ER. Each of these

isomers serves as a model to determine which position on the heterocycle scaffold can best accommodate the SERM-defining basic side chain in the ER-LBD. Once identified, such a site would obviously be a prime target for further study through a series of pyrazoles in which the structure of this basic substituent is varied.¹⁸

The four basic side chain-containing pyrazoles (2–5) shown in Figure 1 are designed to probe several potential ligand orientations within the binding pocket. For example, pyrazole 2 was proposed on the basis of our initial docking study of the N-aryl pyrazole 1 in the ER α -raloxifene X-ray crystal structure¹³ (now revised, as noted above¹⁵), which placed the C(5) phenol in the subpocket corresponding to the A-ring of E_2 . In this mode, the N(1) basic side chain group resides in what would be the estradiol 11β direction. Pyrazole 3 was envisioned to undergo a ligand "flipping" around the bond between the pyrazole core and the C(3) phenol, to place the C(4)-linked basic side chain in the estradiol 11ß direction. Pyrazoles 4 and 5 are regioisomers that have the basic side chain at the 3- and 5-positions, respectively. Pyrazole 5, a C(5)-substituted analogue, was predicted to have a potentially favorable interaction with ER by way of a ca. 120° counterclockwise rotation about an axis perpendicular to the pyrazole ring, resulting in a binding mode that would make the N(1)phenol the mimic of the A-ring in estradiol. Pyrazole 4, a regioisomer of pyrazole 5, was considered unlikely to have a favorable interaction with the ER, because the phenols and the basic side chain groups are displayed in a "long" orientation. This leaves the remaining two free phenols too close to one another for proper interaction with the A-ring and D-ring subpockets in ER. Nevertheless, it was prepared for the sake of completeness.

Chemical syntheses

The synthesis of pyrazole **2** is shown in Scheme 1. For the preparation of the required piperidinyl-ethoxy-substituted

phenylhydrazine, we initially investigated the classical diazotization–reduction route, starting from the corresponding substituted aniline **6**. However, we had difficulty isolating the desired hydrazine precursor. Fortunately, we found the approach of Demers and Klaubert,¹⁹ involving metallation of the corresponding aromatic bromide (7) followed by reaction with an azodicarboxylate ester, conveniently afforded the di-Boc protected hydrazine **8**. Because isolation of the free hydrazine proved to be problematic, we performed the Boc deprotection and pyrazole cyclization in one-pot, which gave pyrazole **9** in 65% yield. Selective demethylation using either AlCl₃– EtSH or BBr₃ afforded pyrazole **2** in 55–75% yield.

The synthesis of the C(4) basic side chain-containing pyrazole 3 is shown in Scheme 2. Introduction of a basic side chain into the pyrazole structure at the 4-position was done early in the synthesis, starting with Freidel-Crafts acylation of anisole with 6-bromohexanoyl chloride. For this transformation, the temperature had to be carefully maintained at -10 °C to avoid reaction of the product ketone 10 with a second equivalent of anisole, a side reaction that results in the formation of a 1,1-diaryl-alkene by-product upon dehydration. The bromoketone 10 was then aminated with excess piperidine in DMF to afford amino-ketone 11 in high yield. Condensation with the *p*-nitrophenyl benzoate furnished the β-diketone 12, which was cyclized with phenyl hydrazine hydrochloride to give the desired protected pyrazole 13. Deprotection using AlCl₃-EtSH yielded the final pyrazole 3.

The final pyrazoles 4 and 5 were initially prepared as an approx. 1:1 mixture of regioisomers by the route shown in Scheme 3. Starting from 4-hydroxybutryophenone, we used a Mitsunobu reaction to prepare the piperidinyl-ethoxy-substituted ketone 14. Claisen condensation afforded the requisite β -diketone 15. Treatment with phenylhydrazine hydrochloride followed by BBr₃ deprotection afforded the desired pyrazoles 4 and 5 as a



Scheme 1. Synthesis of N(1) basic side chain-containing pyrazole 2.



Scheme 2. Synthesis of C(4) basic side chain-containing pyrazole 3.



Scheme 3. Synthesis of C(3) and C(5) basic side chain-containing pyrazoles 4 and 5.

1:1 mixture that could be separated by chromatography. Pyrazole **4** was further purified by recrystallization from MeOH, and definitive regiochemical assignment of this isomer was made on the basis of a single crystal X-ray analysis. An ORTEP diagram of this compound is shown in Figure 2.

Isolation of the last pyrazole (5) proved to be more problematical because we found that the basic side chain of these pyrazoles undergoes partial cleavage when BBr₃ is used as a deprotection reagent. This is a significant problem, because the triphenolic by-product resulting from this cleavage was difficult to separate from the desired C(5) basic side chain isomer by chromato-graphy and it has distinctly different biological activity.¹⁵ Later, we found that we could do this deprotection selectively, without this side reaction, using the milder AlCl₃ EtSH reagent. Ultimately, however, we used a regiose-lective route (described elsewhere in detail¹⁸) to produce the desired pyrazole **5**.

Estrogen receptor binding affinity of pyrazoles 2-5

The binding affinity of pyrazoles 2–5 for ER was assayed in a competitive radiometric binding assay as described previously,^{20,21} and affinities are expressed as relative binding affinity (RBA) values, where the affinity of estradiol is considered to be 100% (Table 1). The affinity of all compounds was tested both in a natural ER preparation from lamb uterine cytosol,²¹ which is predominantly ER α ,^{10,11} as well as with purified recombinant human ER α and ER β (see Experimental).²⁰

When uterine cytosol was used as a source of estrogen receptor, pyrazole **2** had a relatively low binding affinity, just 1.5% that of estradiol, a value which is 10-fold lower than that of the parent N(1) aryl pyrazole (1).^{13,15} Pyrazole **3** with the basic side chain on C(4) also displayed very low affinity for the receptor. Apparently, even with the reorientation proposed above, this analogue, in which the basic side chain is attached to the pyrazole
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Figure 2. X-ray crystal structure for 4 (ORTEP: ellipsoids drawn at the 35% probability level).

core solely through aliphatic linkages, does not fit well in the ligand binding pocket of ER. More intriguing are the two regioisomeric pyrazoles 4 and 5; they display very distinctive preferences for binding to the ER, pyrazole 5 having a 65- to 90-fold higher affinity than pyrazole 4 in all three test systems. As noted previously, the higher affinity of isomer 5 versus 4 was expected.

Pyrazole **5** has the highest affinity of all basic side chain pyrazoles investigated thus far, and, in fact, is among the highest affinity of all pyrazoles we have reported as ER ligands.^{13–15} When we evaluated the binding affinity of these pyrazoles with pure human ER α and ER β (Table 1),²⁰ pyrazole **5** showed a distinctive preference for binding to ER α (ca. 20-fold), a characteristic that it shares with the parent pyrazole triol **1** (ca. 230-fold).¹⁵ Pyrazole **2** shows only a 5-fold affinity preference for ER α . Pyrazoles **3** and **4** had very low affinity for both ER α and ER β , and they were not investigated further either for binding affinity or transcriptional activity.

Transcription activity of pyrazole 5

Based on the promising ER α binding selectivity of pyrazole **5** and because of its piperidine side chain, we were interested in whether this compound had antagonistic properties, and in particular, whether it might function as an antagonist on one or both of the ER subtypes in a potency selective manner. Figure 3 shows the transcriptional activity profiles of pyrazole **5** assayed in HEC-1 cells with either ER α or ER β . Cells were transfected with an expression plasmid for ER α or ER β , together with an estrogen-responsive reporter gene construct ((ERE)₃pS2-CAT), and reporter gene expression was determined at different concentrations of ligand. Antagonism was

Table 1. Estrogen receptor binding affinity of pyrazoles 1-5

Compound	Site of basic side chain	RBA lamb uterine cytosol ^a	RBA hERα ^a	RBA hERβª
1	[none]	20.3±3	35.7±6	0.15±0.01
2	N(1)	1.5 ± 0.4	$0.65 {\pm} 0.13$	0.13 ± 0.03
3	C(4)	$0.017 {\pm} 0.006$	0.05 ^b	0.019 ⁶
4	C(3)	$0.29 {\pm} 0.26$	0.13 ^b	0.01 ^b
5	C(5)	24.5 ± 1	11.5 ± 1	$0.65 {\pm} 0.02$

"Relative binding affinity (RBA) where estradiol = 100%. Values are the average of repeat determinations \pm SD ($n \ge 3$) or \pm range (n = 2). For details of the assay procedure, see Experimental. "Single determinations.



Figure 3. Transcription activation profile for pyrazole 5 with ER α and ER β . Human endometrial cancer (HEC-1) cells were transfected with expression vectors for ER α or ER β and an (ERE)₃-pS2-CAT reporter gene and were treated with indicated concentrations of pyrazole 5 alone (solid lines) or pyrazole 5 in the presence of 10⁻⁹ M estradiol (E₂) (dashed lines) for 24 h. CAT activity was normalized for β -galactosidase activity from an internal control plasmid. Values are the mean ±SD for three separate experiments, and are expressed as a percent of the ER α or ER β response with 10⁻⁹ M E₂, which is set at 100%. For some values, error bars are too small to be visible.

assayed at different concentrations of ligand in the presence of 10^{-9} M E₂.

Pyrazole **5** showed no stimulation of transcriptional activity of ER α or ER β , and it was a full antagonist on ER α and ER β , completely antagonizing activity stimulated by estradiol (Fig. 3). Thus, introduction of the type of basic side chain typically found in the mixed agonist/antagonist SERMs at the proper site on the parent pyrazole **1**, which is an agonist, converts this pyrazole system into an antagonist (pyrazole **5**). The relatively good ER α binding affinity selectivity of pyrazole **5** (ca. 20-fold; cf. Table 1) is preserved to a considerable degree in its potency as a transcriptional antagonist. Its IC₅₀ value is ca. 20 nM on ER α and ca. 160 nM on ER β , which gives an ER α antagonist potency selectivity of ca. 10-fold (Fig. 3).

In work to be presented elsewhere, we have prepared additional analogues of pyrazole **5** having other basic and polar side chains as substituents on the C(5) phenyl group.¹⁸ Some of these have very high affinity selectivity for ER α over ER β , and they act as ER α potency selective antagonists (J. Sun et al., in preparation).

Binding models for pyrazoles 2 and 5 in the ER α -raloxifene crystal structure

As previously mentioned, the low binding affinity of pyrazole 4 was not surprising. The low affinity of pyrazole 3, however, was unexpected, because, as mentioned earlier, when the pyrazole core is flipped, the basic side chain attached to C(4) appears to be orientated properly for this substituent to project through the 11 β channel.

Pyrazole 2, in which the basic side chain is substituted on the N(1) phenyl group, was initially believed to be the most favorable candidate for interacting with ER. However, this turned out not to be the case; pyrazole 5 has the



Figure 4. Pyrazoles 2 and 5 oriented with corresponding aryl rings oriented in a congruent fashion.

highest affinity. These two pyrazoles have somewhat related structures (cf. Fig. 1); the three aryl groups (two unsubstituted and one side chain-substituted) can be superimposed in a congruent fashion, as illustrated in Figure 4. When oriented in this fashion, the only difference between these two isomers is the position of the two ring nitrogen atoms and the C(4) ethyl substituent.

In an attempt to understand the 16-fold higher affinity of pyrazole 5 than pyrazole 2, we performed some docking studies of these two ligands with the ER α -raloxifene

crystal structure.² Both ligands were docked in the $ER\alpha$ -raloxifene crystal structure in a similar manner, and they were subjected to a three-step minimization protocol to generate the final receptor-ligand model complexes shown (see Experimental, Molecular Modeling for details). Residue side chains that were close to each ligand in its initial position were allowed to undergo torsional movement during the docking operation, in order to obtain an optimal fit, prior to minimization. A crossed-stereo view of pyrazoles 2 and 5 is shown in Figure 5(A) and (B). The final models have similar energies, which at least at this level of refinement, simply indicates that they have no bad contacts or unfavorable conformations.

In both of the views shown in Figure 5, the three aryl groups displayed peripherally from the pyrazole core are accommodated comfortably and in a very similar manner within the ligand binding pocket. This was achieved with only minimal and energetically favorable movement of a few residues that were initially in close contact with the ligand.



Figure 5. Crossed-stereo views of pyrazole 2 (top) and 5 (bottom) docked and minimized in ER α -raloxifene structure ligand binding pocket showing residues within 2.5 Å of ligand. The pyrazoles were initially positioned with respect to certain reference atoms of the ligand raloxifene, and then minimized together with the same contact residues in the protein using the Flexidock routine within SYBYL. For details, see Experimental. The yellow surface is a cutaway of the Connolly surface of the protein ligand-binding pocket.

The most significant difference between the complexes with pyrazoles 2 and 5 is the position of the ethyl group (cf. Fig. 4). In Figure 5(B), the C(4) ethyl group of pyrazole 5 projects into a region of ER that accommodates the 18-methyl of estradiol in the ER- E_2 complex.² In this subpocket, the favorable van der Waals interaction that normally occurs between Leu525 and the 18-methyl group of E_2 is conveniently replaced by a similar interaction with the ethyl substituent of pyrazole 5 (Fig. 5(B)).

By contrast, in the complex with pyrazole 2, the ethyl substituent is oriented in a different position (downward), where it no longer benefits as much from a favorable hydrophobic-hydrophobic interaction, as was the case in the complex with pyrazole 5. Leu525 is somewhat closer to the pyrazole core, where it interacts with the N(1) phenyl ring. The closest residue for interaction with the C(4) ethyl of pyrazole 2 (Fig. 5(A)) is Met388. This residue is a somewhat "softer" partner for interaction than Leu525, and may be less stabilizing. Other differences between the ER complexes with pyrazoles 2 and 5 which could account for their differing binding affinity may have to do with electronic effects and are not considered in these models. As noted earlier (Fig. 4), the positions of the nitrogen atoms are different between the two binding modes, so the electronic molecular surfaces that are presented to the receptor are predicted to be quite different.

Conclusions

We have prepared a series of basic side chain-substituted pyrazoles as potential selective antagonists for ER subtypes. The initial four analogues were designed to explore the four possible orientations that the pyrazole core structure might adopt in the ER ligand binding pocket, and thereby to identify which site on the pyrazole could best accommodate the rather large, basic side chain substituent. Of the four analogues, the C(5)piperidinyl-ethoxy-substituted pyrazole 5 was found to have the highest affinity for $ER\alpha$, with an affinity among the highest observed for all ER ligands of the pyrazole type. It also shows considerable affinity selectivity for ER α versus $ER\beta$. Through cell-based transfection assays, we found that pyrazole 5 was an antagonist on both ER α and ER β , and that its potency on ER α was somewhat higher, reflecting its $ER\alpha$ affinity selectivity. Modeling studies using the crystal structure of the ER α -raloxifene complex were used to compare potential binding modes for pyrazoles 2 and 5. The binding modes appear to be quite similar, and the origin of the higher affinity for pyrazole 5 versus 2 may result from a more favorable positioning of the ethyl group in the binding pocket in the complex with pyrazole 5. In work to be presented elsewhere, we have prepared additional analogues of pyrazole 5 having other basic and polar side chains as substituents on the C(5) phenyl group.¹⁸ Some of these have very high affinity selectivity for $ER\alpha$ over $ER\beta$, and they act as $ER\alpha$ potency selective antagonists (J. Sun et al., in preparation). This work is the first effort towards developing a novel SERM based on a pyrazole core structure.

Experimental

General

Melting points were determined on a Thomas-Hoover UniMelt capillary apparatus and are uncorrected. All reagents and solvents were obtained from Aldrich, Fisher or Mallinckrodt. Tetrahydrofuran was freshly distilled from sodium/benzophenone. Dimethylformamide was vacuum distilled prior to use, and stored over 4 A molecular sieves. *n*-Butyllithium was titrated with *N*-pivaloyl-*o*-toluidine. Et₃N was stirred with phenylisocyanate, filtered, distilled, and stored over 4 A molecular sieves. All reactions were performed under a dry N₂ atmosphere unless otherwise specified. Reaction progress was monitored by analytical thin-layer chromatography using GF silica plates purchased from Analtech. Visualization was achieved under illumination with short wave UV light (254 nm) or with a potassium permanganate indicator spray. Radial preparative-layer chromatography was performed on a Chromatotron instrument (Harrison Research, Inc., Palo Alto, CA) using EM Science silica gel Kieselgel 60 PF₂₅₄ as adsorbent. Flash column chromatography was performed using Woelm 32-63 µm silica gel packing. The preparation of pyrazole 5 by a regioselective route is described elsewhere.¹⁸

¹H and ¹³C NMR spectra were recorded on either a Varian Unity 400 or 500 MHz spectrometers using CDCl₃ or MeOD- d_4 as solvent. Chemical shifts were reported as parts per million downfield from an internal tetramethylsilane standard ($\delta 0.0$ for ¹H) or relative to solvent peaks. NMR coupling constants are reported in Hertz. ¹³C NMR spectra were determined using either the Attached Proton Test (APT) or standard ¹³C pulse sequence parameters. Low-resolution and high-resolution electron impact mass spectra were obtained on Finnigan MAT CH-5 or 70-VSE spectrometers. Elemental analyses were performed by the Microanalytical Service Laboratory of the University of Illinois. Single crystal Xray analysis on pyrazole 4 was performed to determine connectivity. A rigorous analysis was not performed because of the presence of a disordered solvent molecule in the unit cell. However, for the purposes of regiochemical assignment, the analysis was satisfactory.

Biological procedures

Relative binding affinity assay. Ligand binding affinities (RBAs) using lamb uterine cytosol as a receptor source were determined by a competitive radiometric binding assay using 10 nM [3H]estradiol as tracer and dextrancoated charcoal as an adsorbent for free ligand, as previously described.²¹ Binding affinities with purified recombinant human ER α and ER β were determined by a competitive radiometric binding assay using 10 nM $[^{3}H]$ estradiol as tracer, commercially available ER α and $ER\beta$ preparations (PanVera Inc. Madison, WI), and hydroxylapatite (HAP) to adsorb bound receptor-ligand complex.²⁰ HAP was prepared following the recommendations of Williams and Gorski.²² All incubations were done at 0°C for 18-24 h. Binding affinities are expressed relative to estradiol (RBA = 100%) and are reproducible with a coefficient of variation of 0.3.

Transcriptional activation assay. Human endometrial cancer (HEC-1) cells were maintained in culture and transfected as described previously.^{23,24} Transfection of HEC-1 cells in 60 nm dishes used 0.4 mL of a calcium phosphate precipitate containing 0.5 µg of pCMVβGal as internal control, 2 µg of the reporter gene plasmid, 100 ng of the ER expression vector, and carrier DNA to a total of 5 µg DNA. Compounds were added to the cell culture media as ethanol solutions so as to yield a final ethanol concentration of 0.1%. CAT activity, normalized for the internal control β-galactosidase activity, was assayed as previously described.^{23,24}

Molecular modeling

Small molecule modeling. The starting structures of pyrazoles 2 and 5 used for the ER-LBD docking studies were generated from random conformational searches using the TRIPOS force field as implemented in Sybyl 6.5.2. Low energy conformers then underwent full minimization until a convergence criterion of 0.001 kcal/ mol was met using the AM1 semi-empirical force field. Charge calculations were done using Gasteiger–Huckel method and molecular surface properties displayed using MOLCAD module in Sybyl 6.5.2.

Receptor docking studies. The lowest energy conformers for pyrazoles 2 and 5, generated as indicated above, were used for docking studies. Prior to docking, a + 1formal charge and a proton were added to each piperidinyl nitrogen. Ligands were then pre-positioned in the crystal structure of the ER α raloxifene complex,² using a least squares multifitting of the most congruent atoms of the pyrazole and raloxifene. In raloxifene this included the A-ring carbon atoms and the 1.4-carbon atoms in both the 2- and 3-substituted phenyl groups. The same fitting strategy was used in the pyrazoles, all six carbon atoms in the ring corresponding to the A-ring and the 1,4-carbon atoms of the other two aryl rings. Once pre-positioned, the raloxifene structure was deleted and then the pyrazole ligand was optimally docked in the ER α binding pocket using the TRIPOS Flexidock routine. Hydrogen bond donors (Arg394, His524) and hydrogen bond acceptors (Glu353, His524, and Asp351) in ER were indicated in Flexidock, as well as the hydrogen bond donors and acceptors in the ligand. Side chains that were allowed to rotate during docking of the pyrazoles 2 and 5 included Met543, Leu539, Leu536, Leu354, Asp351, Trp383, all of which are near the basic side chain group; Phe404, Leu387, and Met388, which are near the A-ring mimic: Leu384, Leu346, Leu428, and Trp383, which are near the B/C ring region; and Ile424, His524, and Leu525, which are near the D-ring subpocket. Arg394 and Glu353, the hydrogen bonding partners of the critical A-ring phenol mimic, were kept fixed during the entire docking and minimization.

The best docked ligand ER-LBD complex from the Flexidock routine then underwent a three-step minimization: First, non-ring torsional bonds of the ligand were minimized in the context of the receptor, using the torsmin command. This was followed by minimization of the side chain residues within 8 Å of the ligand while holding the backbone and residues Glu353 and Arg394 fixed. A final, third minimization of both the ligand and receptor was conducted using the Anneal function (hot radius 8 Å, interesting radius 16 Å), again holding residues Glu353 and Arg394 fixed. The result of this was considered to be the final model. Minimizations were done using the TRIPOS Forcefield (MAXIMIN) with the Powell gradient method and default settings (final RMS $< 0.05 \text{ kcal/mol} \cdot A$). The total energy for the ER α complex with pyrazole 2 was -806 kcal/mol and for the complex with pyrazole 5 was -868 kcal/mol. At this level of refinement, these numbers cannot be used to predict relative binding affinities. However, they indicate that there are no serious errors or unfavorable contacts in the ER-LBD ligand models.

Chemical synthesis

1-[4-(2-N-Piperidinyl-ethoxy)-phenyl]-3,5-di-(4-hydroxyphenyl)-4-ethyl-pyrazole (2). To a stirred solution of the protected pyrazole 9 (100 mg, 0.19 mmol) in CH₂Cl₂ (1.0 mL) at 0 °C was added BBr₃ (1.0 M CH₂Cl₂, 5 equiv, 0.95 mL) dropwise. The reaction was kept at 0 °C for 4 h and then quenched with water. The solution was neutralized with satd NaHCO₃ and then repeatedly extracted with EtOAc. The combined organic layers were washed with brine, dried over Na2SO4 and concentrated to afford a tan residue. Purification by flash chromatography (5% TEA and 5% MeOH in CH₂Cl₂) afforded 2 as an off-white powder (75%): mp 125-130 °C (decomposed); ¹H NMR (CDCl₃, 400 MHz) δ 0.98 (t, 3H, J = 7.3, 1.51 (br s, 2H), 1.62 (quint, 4H, J = 5.5), 2.58 (q, 2H, 7.5), 2.67 (br s, 4H), 2.89 (t, 2H, J = 5.5), 4.13 (t, 2H, J = 5.5), 6.76 (d, 2H, J = 8.5), 6.87 (d, overlapping, 2H, J=8.5), 6.88 (d, overlapping, 2H, J=9), 7.03 (d, 2H, J = 8.5), 7.16 (d, 2H, J = 9.0), 7.49 (d, 2H, J = 8.5); APT ¹³C NMR (CDCl₃, 100 MHz) δ 9.9 (CH₃), 18.5 (CH₂), 24.8 (CH₂), 26.2 (CH₂), 54.1 (CH₂), 56.1 (CH₂), 58.8 (CH₂), 116.1 (CH), 116.8 (CH), 116.9 (CH), 121.4 (C), 123.1 (C), 126.8 (C), 128.5 (CH), 130.9 (CH), 133.0 (CH), 135.2 (C), 144.0 (C), 152.6 (C), 159.1 (C), 159.4 (C), 159.5 (C); MS (EI, 70 eV) m/z 483.3 (M); HRMS (EI) calcd for $C_{30}H_{33}N_3O_2$ (M⁺-HBr): 483.2522. Found: 481.2365 (M⁺-HBr-2).

4-(4-N-Piperdinyl-butyl)-3,5-di-(4-hydroxyphenyl)-1-phenylpyrazole hydrochloride (3). To a stirred CH_2Cl_2 (1.0 mL) solution of the protected pyrazole 13 (72 mg, 0.15 mmol) was added AlCl₃ (116 mg, 0.87 mmol). The mixture was cooled to 0 °C and EtSH (55 µL, 0.73 mmol) added dropwise. The mixture was then allowed to reach rt and stirred for an additional 0.5 h. At this time the reaction was concentrated under a stream of nitrogen in a fume hood, brought up in CH₂Cl₂ and concentrated once again. The crude mixture was re-dissolved in CH_2Cl_2 and a mixture of THF (1 mL), water (1 mL), and 6 M HCl (0.25 mL) added. The resulting precipitate was isolated by vacuum filtration and was washed with four 5mL portions of water followed by four 5mL portions of ether to afford pyrazole 3 as a white solid (37 mg, 54%): ¹H NMR (McOD-d₄, 500 MHz) δ 1.30-1.60 (m, 6H), 1.75 (br s, 6H), 2.73 (t, 2H, J = 7.2), 2.78

(t, 4H, J=8.1), 6.90 (XX' of AA'XX', 2H, $J_{AX}=8.8$, $J_{XX''}=2.5$), 7.07 (AA' of AA'XX', 2H, $J_{AX}=8.8$, $J_{AA'}=2.5$), 7.23 7.36 (m, 5H), 7.32 (XX' of AA'XX', 2H, $J_{AX}=8.6$, $J_{XX'}=2.4$), 7.52 (AA' of AA'XX', 2H, $J_{AX}=8.6$, $J_{AA'}=2.2$); ¹³C NMR (MeOD- d_4 , 125 MHz) δ 21.5, 22.5, 22.3, 23.0, 27.0, 52.8, 56.6, 115.1, 115.2, 117.4, 121.1, 124.8, 124.9, 127.0, 127.3, 128.5, 131.2, 139.8, 142.4, 151.5, 157.5, 157.9; MS (EI, 70 cV) m/z (relative intensity, %): 467 (M⁺-HCl, 21); HRMS (EI, M⁺) calcd for C₃₀H₃₃N₃O₂: 467.2573. Found: 467.2579.

1 - (4 - Hydroxyphenyl) - 3 - |4 - (2 - N - piperidinyl - ethoxy) phenyl]-4-ethyl-5-(4-hydroxy-phenyl)-1H-pyrazole (4). A CH₂Cl₂ solution (10 mL) of the crude protected pyrazole isomers (300 mg, 0.60 mmol) prepared from the 1,3dione 15 was cooled to 0°C and treated dropwise with BBr₃ (1 M in hexane, 5 equiv, 3 mmol). The reaction was allowed to reach temperature and stir overnight. The mixture was re-cooled to 0° C and guenched with 5 mL of MeOH and the solvent removed under reduced pressure. To the crude mixture 8 mL of H₂O was added and the aqueous layer neutralized with a satd aqueous NaHCO₃ solution and extracted repeatedly with EtOAc $(5 \times 20 \text{ mL})$. The organic layers were washed with brine, dried over Na₂SO₄ and concentrated in vacuo to afford a brown residue. Pyrazole 4 was cleanly separated from the regioisomer 5 and other byproducts by flash chromatography (7% Et₃N and 7% MeOH in CH₂Cl₂). It was further purified by recrystallization from MeOH to afford the C(3) isomer (4) as small white crystals whose structure was verified by X-ray analysis (23%). Pyrazole 4: mp 151–153 °C; ¹H NMR ((CD₃)₂SO, 400 MHz) δ 1.01 (t, 3H, J = 7.6), 1.43 (m, 2H), 1.55 (m, 4H), 2.49 (br s, 4H), 2.59 (q, 2H, J=7.6), 2.72 (t, 2H, J=5.6), 4.15 (t, 2H, J = 6.0), 6.74 (d, 2H, J = 8.4), 6.83 (d, 2H, J = 8.4), 7.08 (apparent triplet due to three overlapping doublets. 6H, J=7.6), 7.67 (d, 2H, J=8.4), 9.71 (br s, 2H, OH exchangeable with D₂O); APT ¹³C NMR ((CD₃)₂SO, 100 MHz) δ 15.4 (CH₃), 16.8 (CH₂), 23.9 (CH₂), 25.6 (CH₂), 54.5 (CH₂), 57.5 (CH₂), 65.6 (CH₂), 114.5 (CH), 115.1 (CH), 115.4 (CH), 118.7 (C), 120.9 (C), 126.1 (CH), 126.5 (C), 128.5 (CH), 131.2 (CH), 131.8 (C), 141.2 (C), 148.5 (C), 156.2 (C), 157.4 (C), 157.9 (C); HRMS (EI, M^+) calcd for $C_{30}H_{33}N_3O_3$: 483.2521. Found: 483.2516.

As was noted in the text, isolation of the other regioisomer (pyrazole 5) from this reaction mixture was complicated by the fact that it could not be separated effectively from the triphenolic byproduct that resulted from partial cleavage of the basic side chain during the BBr₃ deprotection. This compound was prepared in a regioselective fashion by another route that is described elsewhere.¹⁸

1-Bromo-4-(2-*N***-piperidinyl-ethoxy)benzene (7).** In a 250 mL flask charged with Ph_3P (35 mmol) in 60 mL of THF at 0 °C was added di-*iso*-propyl-azodicarboxylate (DIAD, 35 mmol) reagent dropwise. The solution was kept at 0 °C and stirred for 50 min. At this time *N*-(2-hydroxyethyl)-piperidine was added dropwise via syringe. After 10 min, *p*-bromophenol (35 mmol) and TEA (86.7 mmol) in 20 mL THF were added dropwise via

an addition funnel. After 1 h at 0 °C the reaction was warmed to rt and stirred for 6 h. The crude mixture was then concentrated in vacuo and the Ph₃PO byproduct removed by vacuum filtration. The filtrate was again concentrated and the product purified by gradient flash chromatography using 5% TEA and 30% EtOAc in hexanes up to 5% TEA and 50% EtOAc in hexanes to afford 7 as a clear oil (85%):^{25 1}H NMR (CDCl₃, 500 MHz) δ 1.41 (m, 2H), 1.58 (m, 4H), 2.47 (br s, 4H), 2.72 (d, 2H, *J*=7.2), 4.05 (d, 2H, *J*=7.6), 6.79 (d, 2H, 8.3), 7.82 (d, 2H, *J*=8.1); ¹³C NMR (CDCl₃, 125 MHz) δ 22.1, 38.0, 67.5, 113.0, 116.5, 121.8, 123.8, 132.3, 136.6, 149.6, 158.0, 158.3; MS (EI, 70 eV) *m*/*z* 284 (M⁺).

N,*N*-di-*tert*-Butoxycarbonyl-*N*-[4-(2-*N*-piperidinyl-ethoxy) phenyllhydrazine (8). To a THF solution at $-70 \,^{\circ}\text{C}$ containing the substituted bromobenzene 7 (10.56 mmol) was added *n*-BuLi (1.3 equiv) dropwise over 10 min. After 30 min, *tert*-butoxycarbonyl-azodicarboxylate (15.84 mmol) was added as a solid in one portion. The mixture was then slowly warmed to rt and 1 equiv of dilute AcOH added. After stirring the guenched mixture for 30 min, the reaction was partitioned between water (30 mL) and Et₂O (50 mL). The layers were separated and the aqueous layer extracted twice more with Et₂O $(2 \times 50 \text{ mL})$. The combined organic layers were washed with brine and dried over Na₂SO₄. Subsequent solvent removal afforded 6.8 g of a crude red oil. Flash chromatography (5% TEA and 25% EtOAc in hexanes) produced 8 as a white solid (75%): mp 50-55°C; (note: spectrum complicated by Boc rotamers); ¹H NMR (CDCl₃, 400 MHz) δ 1.38 (s, 9H), 1.40 (overlapping s, 11H), 1.61 (m, 4H), 2.45 (m, 4H), 2.73 (t, 2H, J = 5.6), 4.05 (t, 2H, J = 5.3), 6.82 (d, 2H, J = 7.8), 7.25 (br s, 2H); ¹³C NMR (CDCl₃, 100 MHz) δ 24.2 (CH₂), 24.8 (CH₂), 28.2 (CH₃), 28.3 (CH₃), 36.8 (CH₂), 46.0 (CH₂), 46.7 (CH₂), 55.1 (CH₂), 57.8 (CH₂), 60.4 (CH₂), 65.7 (CH₂), 66.0 (CH₂), 81.0 (CH₂), 81.8 (CH₂), 113.9 (CH), 114.6 (CH), 120.7 (CH), 129.4 (CH), 156.7 (C), 158.7 (C), 171.2 (C); MS (EI, 70 eV) m/z 435.4 (M, 35%). Anal. (C₂₃H₃₇N₃O₅): C, 63.42; H, 8.56; N, 9.65. Found: C, 63.20; H, 8.72; N, 9.47.

1-[4-(2-N-Piperidinyl-ethoxy)-phenyl]-3,5-bis-(4-methoxyphenyl)-4-ethyl-1H-pyrazole (9). To a stirred solution of DMF (14mL), THF (6mL), and 1,3-bis-(4-methoxyphenyl)-2-ethyl-propane-1,3-dione (500 mg, 1.52 mmol)¹⁵ was added hydrazine 8 in 3-fold excess (800 mg, 4.59 mmol). The mixture was brought to reflux for 5h and then recooled to rt. The THF was removed under reduced pressure and the remaining residue diluted with H₂O (30 mL) and extracted repeatedly with EtOAc. The organic layers were washed with brine, dried over Na₂SO₄ and concentrated to produce a crude redorange oil. Flash chromatography using 10% TEA and 20% EtOAc in hexanes afforded the pure product 9 as an orange oil (65%): ¹H NMR (CDCl₃, 400 MHz) δ 1.03 (t, 3H, J=8.1), 1.42 (m, 2H), 1.58 (quin, 4H, J=6); 2.47 (br s, 2H), 2.61 (q, 2H, J = 6.2), 2.72 (t, 2H, J = 6), 3.79 (s, 3H), 3.82 (s, 3H), 4.04 (t, 2H, J = 6.2), 6.76 (d, 2H, J = 9.0), 6.86 (d, 2H, J = 9.0), 6.96 (d, 2H, J = 9.1), 7.13 (d, 2H, J=9.1), 7.16 (d, 2H, J=9.1), 7.69 (d, 2H, J = 9.0); ¹³C NMR (CDCl₃, 100 MHz) δ 15.6 (CH₃),

17.2 (CH₂), 24.3 (CH₂), 26.0 (CH₂), 55.1 (CH₃), 55.2 (CH₃), 57.9 (CH₂), 66.2 (CH₂), 113.8 (CH), 113.9 (CH), 114.5 (CH), 119.9 (C), 123.3 (C), 126.0 (CH), 126.9 (C), 129.0 (CH), 131.3 (CH), 133.6 (C), 141.0 (C), 150.1 (C), 157.4 (C), 159.1 (C), 159.4 (C); HRMS (EI, M⁺) calcd for $C_{32}H_{37}N_3O_2$: 511.2834. Found: 511.2825.

6-Bromo-1-(4-methoxy-phenyl)-hexan-1-one (10). To a stirred solution of AICI₃ (12.40 g, 92.75 mmol) in 1,2dichloroethane (30 mL) at 0 °C was added 6-bromohexanoyl chloride (23.70 g, 17 mL, 111.30 mmol) dropwise over 10 min. The resulting solution was stirred at rt for 0.5 h, cooled to $-15 \circ \tilde{C}$ and a solution of anisole (10.00 g, 10.05 mL, 92.75 mmol) in 1,2-dichloroethane (10 mL) was added dropwise over 20 min. The reaction mixture was allowed to stir at -15 °C for 1 h, then quenched with $H_{2}O$ (50 mL). The aqueous layer was isolated and extracted with CH_2Cl_2 (2×50 mL), and the organic layers were combined and washed with H₂O $(2 \times 50 \text{ mL})$, saturated NaHCO₃ $(2 \times 50 \text{ mL})$, and brine (2×50 mL). Subsequent drying over Na₂SO₄ and solvent removal in vacuo afforded a crude orange oil. Upon standing at rt, large crystals formed after 48 h. The crystals were isolated and rinsed with cold hexane to afford 10 (23 g, 87%): mp 50-52 °C; ¹H NMR (CDCl₃, 400 MHz) δ 1.52 (quint, 2H, J=7.6), 1.76 (quint, 2H, J = 7.5), 1.91 (quint, 2H, J = 7.1), 2.94 (t. 2H, J = 7.3), 3.42 (t, 2H, J = 6.7), 3.87 (s, 3H), 6.93 (d, 2H, J = 8.8), 7.94 (d, 2H, J=9.0); ¹³C NMR (CDCl₃, 100 MHz) δ 23.39, 27.79, 32.51, 33.59, 37.80, 55.34, 113.56, 129.88, 130.13, 163.24, 198.48; HRMS (EI, M⁺) calcd for C13H17O2Br: 286.0391. Found: 286.0394.

1-(4-Methoxy-phenyl)-6-(N-piperidinyl)-hexan-1-one (11). To a stirred solution of ketone 10 (248 mg, 1.00 mmol) in DMF (30 mL) at 0 °C was added piperidine (170 mg, 0.20 mL, 2.00 mmol) dropwise over 5 min. The solution was placed in a 70 °C oil bath for 4 h and then cooled to rt. The crude mixture was concentrated under reduced pressure to remove a majority of the DMF and excess piperidine to afford an orange residue. The residue was taken up in CH₂Cl₂ (30 mL), washed with brine (2×20 mL), dried over MgSO₄ and concentrated under reduced pressure to afford 11 as a white solid (227 mg, 79%): mp 163-165°C; ¹H NMR (CDCl₃, 500 MHz) δ 1.35 1.45 (m, 2H), 1.65 2.00 (m, 8H), 2.26 (br g, 2H, J=8), 2.59 (br q, 2H, J=7), 2.88 2.96 (m, 4H), 3.51 (br d, 2H), 3.88 (s, 3H), 6.93 (d, 2H, J_8.9), 7.93 (d, 2H, J = 8.9; ¹³C NMR (CDCl₃, 100 MHz) δ 22.1, 22.3, 23.3, 23.4, 26.4, 37.5, 53.1, 55.5, 57.2, 113.8, 129.9, 130.3, 163.5, 198.5; MS (EI, 70 eV) m/z (relative intensity, %): 289 (M⁺, 3); HRMS (EI, M⁺) calcd for $C_{19}H_{27}NO_2$: 289.2042. Found: 289.2060.

1,3-Bis-(4-methoxy-phenyl)-2-(4-*N***-piperidinyl-butyl)-propane-1,3-dione (12).** To a solution of ketone **11** (350 mg, 1.21 mmol) and TMEDA (140.6 mg, 0.18 mL, 1.21 mmol) in THF (15 mL) at 0 °C was added a 1.0 M solution of LiHMDS (3.02 mL, 3.02 mmol) dropwise. The solution was allowed to stir at rt for 0.5 h, then was cooled back to 0 °C. At this time a solution of 4-nitrophenyl 4-methoxybenzoate (990 mg, 3.63 mmol; prepared from *p*-nitrophenol and 4-methoxybenzoic acid using DIC and DMAP coupling conditions) in THF (5mL) was added dropwise. The resulting solution was allowed to stir for 15h at rt and 4h at reflux (oil bath temperature of 70 °C). At this time the mixture was allowed to come to rt and quenched by the addition of $H_2O(10 \text{ mL})$. The organic layer was isolated and washed with H₂O $(3 \times 15 \text{ mL})$, dried over Na₂SO₄, and concentrated under reduced pressure to afford a yellow solid. The remaining ester was removed by warming the crude mixture in 15% ethyl acetate/hexanes and filtering off the excess crystalline ester. The filtrate was concentrated under reduced pressure and purified by flash chromatography (10% TEA and 55% EtOAc in hexanes) to afford 12 as a yellow oil (227 mg, 51%): ¹H NMR (CDCl₃, 400 MHz) δ 1.22-1.48 (m, 4H), 1.50 -1.75 (m, 6H), 2.10 (q, 2H, J=7.5), 2.34 (t, 2H, J=9.8), 2.42 (br s, 4H), 3.83(s, 6H), 5.05 (t, 1H, J = 6.6), 6.89 (d, 4H, J = 8.9), 7.94 (d, 4H, J = 8.9); ¹³C NMR (CDCl₃, 100 MHz) δ 24.0, 25.3, 26.1, 26.3, 29.3, 54.3, 55.4, 57.0, 58.7, 113.9, 128.9, 128.9, 130.8, 130.8, 163.6, 194.6.

3.5-Bis-(4-methoxyphenyl)-4-(4-N-piperidinyl-butyl)-1phenyl-1*H*-pyrazole (13). To a stirred solution of DMF (45 mL), THF (20 mL), and β -diketone 12 (170 mg, 0.40 mmol) was added phenylhydrazine hydrochloride (289 mg, 2.00 mmol). The solution was brought to reflux (oil bath temperature between 110 and 120 °C) for 22 h. The reaction mixture was allowed to cool to rt, and the THF was evaporated under reduced pressure. The remaining mixture was diluted with H₂O (40 mL) and extracted with EtOAc (3×40 mL). The organic layers were combined and washed with saturated LiCl $(3 \times 40 \text{ mL})$, saturated NaHCO₃ $(2 \times 40 \text{ mL})$, H₂O $(2 \times 40 \text{ mL})$, and brine $(2 \times 40 \text{ mL})$. The organic layer was dried over MgSO₄ and concentrated under reduced pressure to afford a crude brown oil. Purification by flash chromatography (10% TEA and 55% EtOAc in hexanes) afforded 13 as a reddish oil (79 mg, 40%). Compound 13 was verified by ¹H NMR and used in the next step of the reaction scheme without further characterization.

1-[4-(2-N-Piperidinyl-ethoxy)-phenyl]-butan-1-one (14). To a solution of PPh₃ (10.05 g, 38.31 mmol) in THF (150 mL) at 0 °C was added diisopropyl azodicarboxylate (7.5 mL, 38.31 mmol) dropwise. After stirring at 0°C for 50 min, N-piperidinylethanol (5.09 mL, 38.31 mmol) was added followed 10 min later by addition of a solution of 4-hydroxy-butyrophenone (5.24 g, 31.9 mmol) in Et₃N (13.4 mL, 95.8 mmol) and THF (30 mL). The mixture was allowed to warm to rt and stirred overnight and then concentrated in vacuo. To the crude residue was added Et₂O (150 mL) and the insoluble Ph₃PO by-product removed by filtration. The filtrate was concentrated once again and the residue purified by flash chromatography (10% TEA and 25% EtOAc in hexanes) to afford an oil. This material was dissolved in Et₂O (60 mL) and acidified with HCl (4 M in dioxane, 6.3 mL). The precipitate that formed was collected by vacuum filtration and partitioned between saturated $NaHCO_3$ (35 mL) and ether (50 mL). The organic layer was washed with brine, dried over Na2SO4 and concentrated under reduced pressure to give 14 as pale-yellow

crystals (3.05 g, 35%). An analytically pure sample was obtained by recrystallization from hexane: mp 48–50 °C; ¹H NMR (CDCl₃, 400 MHz) δ 0.99 (t, 3H, *J*=7.5), 1.43–1.49 (m, 2H), 1.64 (quint, 4H, *J*=5.9), 1.75 (sext, 2H, *J*=7.6), 2.55 (br s, 4H), 2.87 (t, 2H, *J*=5.9), 2.90 (t, 2H, *J*=7.2), 4.19 (t, 2H, *J*=6.0), 6.93 (XX' of AA'XX', 2H, *J*_{AX}=8.9, *J*_{XX'}=2.5), 7.93 (AA' of AA'XX', 2H, *J*_{AX}=9.0, *J*_{AA'}=2.5); ¹³C NMR δ 14.1, 18.2, 24.2, 26.0, 40.4, 55.3, 57.9, 66.3, 114.4, 130.5, 162.7, 199.2; HRMS calcd for C₁₇H₂₆NO₅: 276.1963. Found: 276.1964.

2 - Ethyl - 1 - (4 - methoxyphenyl) - 3 - [4 - (2 - N - piperidinyl - 1)]ethoxy)-phenyl|-propane-1,3-dione (15). To a THF (25 mL) solution of ketone 14 (500 mg, 1.82 mmol) and 4-nitrophenyl 4-methoxybenzoate (745.3 mg, 2.73 mmol) at 0°C was added LiHMDS (1 M solution in hexane, 4.55 mL, 4.55 mmol) dropwise. Upon complete addition of LiHMDS the solution was allowed to warm to rt and stir overnight. The reaction mixture was concentrated to near dryness and EtOAc (35 mL) added. The crude mixture was washed sequentially with saturated NaHCO3 and brine. The solvent was dried over Na₂SO₄, removed in vacuo and the resulting oil purified via flash chromatography (10% TEA and 50% EtOAc in hexanes) to give 15 as a viscous red oil (465 mg, 63%): ¹H NMR (CDCl₃, 400 MHz) δ 1.03 (t, J=7.31), 1.49 (br s, 2H), 1.70 (br s, 4H), 2.16 (quint, J = 7.29 Hz, 2H), 2.63 (br s, 4H), 2.90 (br s, 2H), 3.85 (s, 3H), 4.25 (br s, 2H), 4.96 (t, J = 6.76 Hz, 1H), 6.91 (d, J = 9.05 Hz, 4H), 7.95 (d, J = 8.96 Hz, 2H), 7.96 (d, J = 8.85 Hz, 2H); ¹³C NMR (CDCl₃, 125 MHz) δ 12.8, 23.1, 24.0, 25.8, 55.0, 55.4, 57.6, 59.0, 66.2, 113.9, 114.5, 116.5, 129.2, 129.3, 130.9, 131.0, 163.0, 163.7, 194.9; HRMS (EI, M⁺) calcd for C₂₅H₃₁NO₄: 409.2253. Found: 407.2096 $(M^+ - 2).$

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Regioselective Synthesis of 1,3,5-Triaryl-4-alkylpyrazoles: Novel Ligands for the Estrogen Receptor

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ABSTRACT



A regioselective synthesis of 4-alkyl-1,3,5-triarylpyrazoles has been developed for the preparation of unsymmetrically substituted systems of interest as ligands for the estrogen receptor.

In our efforts to discover novel ligands for the estrogen receptor (ER) that might act as selective estrogen receptor modifiers (SERMs),¹ we found that 1,3,5-triaryl-4-alkylpyrazoles such as 1 and 2 (Scheme 1) were good ligands for ER, demonstrating high binding affinities and transcriptional efficacy that in some cases were very selective for the ER α subtype (ER α).^{2,3} Initially, we synthesized these pyrazoles by condensation of 2-alkyl-1,3-diketones with arylhydrazines.⁴⁻⁶ Of course, when the 1,3-diketones were unsymmetrical, this approach did not afford any significant regioselectivity. This lack of regioselectivity became of concern when we needed the corresponding monophenols 3 and 4 for structure-activity studies to determine which phenol in pyrazole 2 mimics the A-ring of estradiol. According to a classical approach, the monophenol with the higher affinity can be presumed to be the one that corresponds to the A-ring

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of estradiol.^{7.8} However, when the original 1,3-dione hydrazine condensation pyrazole synthesis was used to prepare these monophenols, only an inseparable mixture of the two regioisomers **3** and **4** was afforded (Scheme 1). Thus, a regioselective approach to these and related compounds was needed.

In a related effort, we wanted to develop these novel 1,3,5triaryl-4-alkylpyrazole ligands into the sort of mixed agonist/ antagonists that typically have SERM activity.^{9,10} This generally involves incorporating a basic or polar side chain (such as a piperidinylethoxy group) onto either the C(3) or C(5) phenyl groups. However, when pyrazoles **6** and **7** were prepared by condensation of 4-methoxyphenylhydrazine with unsymmetrical 1,3-diketone **5**, we obtained the regiosiomeric pyrazoles **6** and **7** in pure form only after exhaustive chromatography, and we had to obtain an X-ray structure of the more crystalline isomer **6** to establish the identity of these regioisomers (Scheme 1).

The results from cell-based transcriptional assays showed that pyrazole 7 has the desired antagonistic character typical for a SERM. However, to examine this series further we

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pyrazole 7 and its analogues, we prepared α , β -unsaturated ketone 21 by an aldol condensation of 4-methoxyacetophenone and *p*-hydroxybenzaldehyde, according to a literature procedure¹⁹ with modifications (Scheme 4). Despite numerous attempts, we were unable to obtain good yields in this simple reaction. However, we were able to isolate the highly crystalline enone 21 easily.

Enone 21, protected as its silvl ether (22), reacted with 4-methoxylphenylhydrazine to give pyrazoline 23. This material was alkylated, as before, with various iodides to



 MnO_2 or DDQ to afford the desired pyrazoles 27-29. Fluoride ion cleavage of the silvl group the gave the C(5)phenolic pyrazoles 30-32. An X-ray structure of one of these pyrazoles (31, Scheme 4) secured the structure of this compound, in the process confirming the regioselectivity of this route to pyrazoles. Installation of the piperidinylethoxy side chain was accomplished by a Mitsunobu reaction. Although BBr3 cleaved all three ether groups, AlCl3-EtSH selectively cleaved only the methyl ethers, leaving the basic side chain unaffected and giving pyrazoles 33-35 in very high yield. A number of other side chain derivatives (36-41) were prepared in the C(4) ethyl series. Pyrazole 41 was prepared by a closely related route (not shown). We are currently investigating the biological activities of all of the new pyrazole compounds bearing the various polar/basic side chains.

Basis of Regioselectivity. The regioselectivity of this pyrazole synthesis derives from the initial enone-arylhydrazine condensation, which results in the attachment of the aryl-substituted hydrazine nitrogen to the enone β -carbon and the unsubstituted hydrazine nitrogen to the enone carbonyl carbon. Two mechanisms seem plausible for this transformation (Scheme 5), and they differ in the timing of bond formation. In path a, the aryl-substituted nitrogen reacts first, undergoing a Michael-type addition to the β -carbon of the enone which is followed by an intramolecular imine formation between the carbonyl group and the free amine. In path b, imine formation between the unsubstituted nitrogen and the carbonyl group occurs first, this being followed by a cyclization process to a zwitterionic species that undergoes proton tautomerization to furnish the pyrazoline. Whereas pathway b might first appear to be an ionic 5-endo-trig process, it can more reasonably be considered to be a concerted, symmetry-allowed closure of a 1,2-diaza analogue





of a pentadienyl anion to a 1-azaallyl anion. No intermediates are observed in this transformation. Thus, at this point, there is no definitive basis for favoring one mechanism over the other. However, the fact that α -substituted unsaturated ketones (e.g., 8) fail to react under conditions where the unsubstituted congeners (e.g., 10) react well (see Scheme 2) would be more consistent with mechanism a.

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Supporting Information Available: Procedures for the preparation of all of the compounds mentioned in this paper and their spectroscopic characterization. This material is available free of charge via the Internet at http://pubs.acs.org.

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Synthesis and Biological Evaluation of a Novel Series of Furans: Ligands Selective for Estrogen Receptor Alpha

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Deborah S. Mortensen, Alice L. Rodriguez, Kathryn E. Carlson, Jun Sun, Benita S. Katzenellenbogen and John A. Katzenellenbogen

3-Alkyl-2,4,5-triarylfurans are potent agonists for the estrogen receptor (ER), having highselectivity for ER α



Abstract

A variety of non-steroidal systems can function as ligands for the estrogen receptor (ER), in some cases showing selectivity for one of the two ER subtypes, ER α or ER β . We have prepared a series of heterocycle-based ligands for the estrogen receptor, furans, thiophenes, and pyrroles, and assessed their behavior as ER ligands. An aldehyde enone conjugate addition approach and an enolate alkylation approach were developed to prepare the 1,4-dione systems that were precursors to the tri-substituted and tetra-substituted systems, respectively. All of the diones were easily converted into the corresponding furans, but formation of the thiophenes and pyrroles from the more highly substituted 1,4-diones was problematical. Of the systems investigated, the tetra-substituted furans proved to be most interesting. They were ER α bindingand potency-selective agents, with the triphenolic 3-alkyl-2,4,5-tris(4-hydroxyphenyl)furans (15a-d) displaying generally higher subtype binding selectivity than the bis-phenolic analogs (15f-i). Binding selectivity for ER α was a high as 50-70 fold, and transcriptional activation studies showed that several members of this series were ER α selective agonists, with the best compound (3-ethyl-2,4,5-tris(4-hydroxyphenyl)furan, 15b) having full transcriptional activity on ER α while being inactive on ER β . Comparative binding affinity analysis and molecular modeling were used to investigate the preferred binding mode adopted by the furan ligands, which appears to have the C(2) phenol mimicking the important role of the A-ring of estradiol. These ligands should be useful in studying the biological roles of both ER α and ER β , and they might form the basis for the development of novel estrogen pharmaceuticals.

The estrogen receptor (ER) is a ligand-regulated transcription factor whose activity as an inducer or repressor of gene transcription depends upon the nature of the ligand with which it is bound, as well as the nature of the co-regulator proteins with which it associates.¹ Estrogen action is important in many tissues, and ER is involved in the development and function of the reproductive system and plays a role in bone density maintenance,²⁻⁴ regulation of blood-lipid profiles,³⁻⁷ and brain function.^{8,9} Not surprisingly, ER is a target for the discovery of new drugs for treating or regulating a variety of hormone-related conditions.

There are two ER genes, the well known gene that produces subtype ER α and the more recently discovered gene that produces the subtype ER β .^{10,11} The tissue distribution of ER α and ER β differ, though the biological significance of this difference is not yet well understood.¹² A number of previously known ER ligands,¹²⁻¹⁵ as well as some recently developed novel ligand systems,¹⁶ have different potencies¹⁷ or are able to effect different response levels through ER α and ER β .¹⁸ Ligands having very high ER sub-type selectivity would be effective probes of the respective biological roles of ER α and ER β , and might also function as tissue-selective agents having improved endocrine profiles.

The ER sub-types α and β have only fifty-five percent amino acid identity in the ligand binding domain, yet all but two of the residues that define the ligand binding pocket are the same.¹⁹ The two differences are conservative substitutions, with Met421 in ER α corresponding to an isoleucine in ER β , and similarly Leu384 in ER α being replaced with a methionine in ER β . These subtle changes in the ligand-binding pocket of the two ER sub-types do not provide a definitive basis for understanding the selectivity in either binding or potency that is seen with some ligands; so, further structural analyses will be required.

In previous investigations, we found that certain pyrazoles, particularly those displaying a 1,3,5-triaryl-4-alkyl substitution pattern, were very selective for ER α , in terms of affinity, potency and efficacy.²⁰⁻²⁵ Pyrazole **1** was found to have the highest ER α binding affinity, but pyrazole **2** had the greatest ER α subtype selectivity (Figure 1). While certain azoles, such as oxazoles, thiazoles, isoxazoles, and imidazoles, did not produce good ER ligands in our hands,²⁰

we wondered whether other five-membered ring heterocycles bearing only one heteroatom, namely, furans, thiophenes and pyrroles, might provide ER ligands with interesting biological character. Several furans (**3-5**) and pyrroles (**6-8**) have been described as potential anti-fertility agents.²⁶⁻²⁹ Few, however, showed significant activity, the most active being the furans (**3-5**), and binding and transcriptional activity were reported only in one instance.²⁶

[Figure 1]

[Figure 2]

In this report we describe an investigation of single heteroatom-containing fivemembered ring heterocyclic analogs of the high affinity pyrazole ligands. We find that certain furans are ER ligands that display ER α -selective biological character equivalent to that of the pyrazoles. We have also developed a structure-activity relationship within the furan series to optimize ligand affinity and selectivity, and to investigate the binding mode adopted by this class of ligands.

Results

Synthesis of Ligands

While the literature provides a plethora of approaches to furans, thiophenes, and pyrroles, we wished to prepare all three of these heterocycles from a common precursor, namely, a 1,4-dione. The trisubstituted 1,4-diketones **10a-c** were prepared by the conjugate addition of an aldehyde to an α,β -unsaturated ketone **9a-c** (Scheme 1). Treatment of commercially available acetophenones and aromatic aldehydes with ethanolic KOH afforded chalcones **9a-c**. The α,β -unsaturated ketones **9a-c** then underwent conjugate addition reactions with commercially available aldehydes, utilizing Stetter's thiazolium salt catalysts, either 3-ethyl-5-(2-hydroxyethyl)-4-methyl-thiazolium bromide for aromatic aldehydes.³⁰ The Stetter reactions produced the desired 1,2,4-*tri-substituted* butane-1,4-diones **10a-c** in good yields. However, all

of our attempts to use this approach with an α -substituted enone to produce the desired *tetra-substituted* diones were unsuccessful.

[Scheme 1]

The 1,2,3,4-tetrasubstituted butanediones were instead prepared by the alkylation of an enolate with an α -bromoketone (Scheme 2). Treatment of the 1,2-diaryl-ethanones **11a-d** with potassium hexamethyldisilyl amide (KHMDS), followed by the addition of an α -bromoketone **12a-f**, provided the desired 1,3,4-triaryl-2-alkyl-butane-1,4-diones **13a-l**. These reactions afforded the product diones as mixtures of diastereomers, which in some cases were separable by crystallization. Separation was not required, however, because these centers become trigonal in the final products. The α -bromoketones **12a-f** were produced in nearly quantitative yields from the parent ketones upon treatment with bromine and a catalytic amount of aluminum chloride.

[Scheme 2]

Furans: The 1,4-diketones were converted into furans upon treatment with catalytic *p*-toluenesulfonic acid in refluxing toluene. Table 1 lists the yields for conversion of diones **13a-1** and **10a-b** to furans **14a-n**. The methyl ether protecting groups of furans **14a-n** were removed using boron trifluoride-dimethyl sulfide complex, to afford furans **15a-n** as free phenols. The demethylation reactions and yields for conversion of **14a-n** to **15a-n** are listed in Table 2.

[Table 1 and Table 2]

Thiophenes: Upon treatment with Lawesson's reagent, diones **10a-b** were converted into thiophenes **16a-b**.^{31,32} Furan formation was found to be a competing process in these reactions, sometimes giving as high as a 1:1 furan to thiophene ratio when the 1,4-diones were treated with Lawesson's reagent alone. Competing furan formation not only reduced yields, but the furan byproducts could not be separated from the corresponding thiophenes by flash column chromatography or recrystallization. Fortunately, the addition of solid potassium carbonate as an acid scavenger minimized furan formation, so the thiophenes could be isolated in pure form. Demethylation of **16a-b** with boron trifluoride-dimethyl sulfide complex afforded the phenolic thiophene analogs **17a-b** in good yields (Scheme 3).

[Scheme 3]

Unfortunately, all attempts to convert the 1,2,3,4-*tetra-substituted* butanediones into thiophenes afforded solely furan products. Even with the addition of potassium carbonate, Lawesson's reagent gave only quantitative yields of the furans. An alternate approach using hydrogen sulfide-hydrogen chloride also failed to give the desired thiophenes, instead producing furan in low yield and leaving unreacted starting material. It is known that these latter reagents can be used to produce *di-substituted* thiophenes either from 1,4-diones^{33,34} or directly from the corresponding furans.³³ However, these conditions were reported not to work for the conversion of *tetra-substituted* systems, such 1,4-diaryl-2,3-dibromo-butane-1,4-diones, into the corresponding thiophenes.³⁴ While our failure to convert tetra-substituted 1,4-diones into thiophenes was disappointing, it is not surprising considering the ample precedent for preferential furan formation from hindered 1,4-diones.³³⁻³⁶ In fact, to our knowledge, there has been only one report of the formation of thiophenes from 1,2,3,4-tetra-substituted-1,4-dicarbonyl precursors.³⁷

Pyrroles: Treatment of diones **10c** with *p*-anisidine and *p*-toluenesulfonic acid produced pyrrole **18** in modest yield; however, unreacted dione may be recycled to increase production of the desired heterocycle. Demethylation of **18** provided the target pyrrole **19** (Scheme 4). As was the case with the thiophenes, all attempts to convert 1,2,3,4-tetrasubstituted-butane-1,4-diones into pyrroles resulted in quantitative furan formation.

[Scheme 4]

Biological Studies

Relative Binding Affinities: The relative binding affinity of the heterocyclic ligands were determined by a competitive radiometric receptor binding assay first using lamb uterine cytosol as a source of receptor.³⁸ (The uterus is thought to contain almost exclusively ER α .)¹² The higher affinity ligands were then studied further with purified human ER α and ER β .^{18,39} The

results of these studies, given as relative binding affinity (RBA) values where the affinity of estradiol is considered to be 100%, are summarized in Table 3.

While the members of these series of heterocycles displayed somewhat lower affinities for uterine cytosol ER than did those of the pyrazole class,²⁰ they showed good affinity nevertheless, with the highest RBA of 6.5% being observed for the 2,4,5-tris(4-hydroxyphenyl)-3-ethyl furan **15b**. As was the case with the pyrazoles,²⁰ the tri-substituted ligands, furans **15m**,**n** and thiophenes **17a**,**b**, showed very low affinities and thus were not assayed further with ER α and ER β . The one pyrrole we prepared (**19**) also had low affinity; this was unexpected, considering that it is a tetra-substituted ligand that is a close structural analog to a high affinity pyrazole (cf., Fig 1).²⁰

[Table 3]

All of the tetra-substituted furans (15a-l) were assayed for binding to ER α and ER β , and a number of structure-affinity trends are of note. All of these furans had much higher affinity for purified ERa than for ER from lamb uterine cytosol. We believe that this is due to reduced nonspecific binding of these relatively lipophilic ligands in the assays with the purified receptor, $ER\alpha$, which are performed at much lower total protein concentrations than are the assays using uterine cytosol. While all of the furans proved to be ER α binding selective ligands, the data indicates that the presence of a third phenolic hydroxyl is required to achieve the highest binding selectivity for ER α versus ER β (compare tris-phenols vs the corresponding bis-phenols: 15c vs 15g, and 15b vs 15f,h,i, respectively). As the size of the 3-alkyl substituent is changed in the triphenols, the highest ER α binding affinity is found with the ethyl and propyl analogs (15b and **15c** with of RBAs 100-140), but the highest ER α binding selectivity was found with the methyl furan 15a (RBA ER α /ER β = 65). The symmetrical tetra-(4-hydroxyphenyl)furan (15e) has respectable ER α affinity and selectivity, but is not as good as the best of the 3-alkyl analogs. The homologous series of bis-phenols (15f,h,i) and mono-phenols (15j-l) were prepared to investigate the orientation of these furans in the ligand binding pocket of ER; their affinities are lower, and the structure-affinity correlations are considered further in the Discussion section.

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Transcriptional Activation Assays: Four high affinity tetra-substituted furan ligands (**15a,b,c,g**) were assessed for transcriptional activation activity through both ER α and ER β . These cotransfection assays are conducted in human endometrial carcinoma (HEC-1) cells, using expression plasmids for full-length human ER α or ER β and an estrogen-responsive luciferase reporter gene system.¹⁶ The results of these initial screening assays are summarized in Figure 3.

All four ligands tested were more potent on ER α than ER β , and they were fully efficacious on ER α . On ER β , however, the three tri-phenols (**15a-c**) were inactive, but the bisphenol (**15g**) showed partial agonist activity. Thus, while having a third phenol on the C(5) phenyl group is not essential for high ER α binding affinity (Table 3), it does appear to be important to ensure that the furans are highly selective for ER α and not efficacious on ER β .

A full dose-response curve for transcriptional activation through ER α and ER β for one of the most selective of the ligands assayed, 3-ethyl-2,4,5-tris(4-hydroxyphenyl)furan (**15b**), is shown in Figure 4. This furan is highly selective in its activation of ER α , having an EC₅₀ = 0.33 nM for ER α and having essentially no agonist or antagonist activity on ER β at concentrations up to 1 μ M.

[Figure 3]

[Figure 4]

Discussion

We have prepared a series of aryl-substituted 5-membered ring heterocycles containing a single heteroatom, namely, furans, thiophenes, and pyrroles, and we have assessed them as ligands for the estrogen receptor (ER). Certain tetra-substituted furans proved to be very selective for ER α , in terms of binding affinity and potency as agonists, with the triphenolic 3-alkyl-2,4,5-tris(4-hydroxyphenyl)furans (**15a-d**) displaying generally higher subtype binding selectivity than the bis-phenolic analogs (**15f-i**)(Table 3). The highest ER α binding selectivity (65 fold) was obtained with the 2,4,5-tris(4-hydroxyphenyl)-3-methyl furan **15a**, but the best overall combination of high affinity and ER α selectivity was given by the corresponding 3-ethyl

and 3-propyl analogs, furans **15b** and **15c** (Table 3). The ethyl furan **15b** is sufficiently ER α selective, that it can be used to activate ER α fully at concentrations (10-100 nM) where it has no agonist or antagonist activity on ER β (Figure 4).

Structure-Activity Relationships in the Triaryl-Furan Series: The heterocycles most closely related to the furans, thiophenes, and pyrroles studied here are the pyrazoles that we have investigated extensively (Fig. 1),^{20,22-25} and certain features that we noted in the behavior of the pyrazoles are also found with the new heterocycles. The triaryl-substituted systems have low binding affinity, as was the case with the pyrazoles (and other related heterocycles), and high affinity and ER α selectivity was engendered by having a fourth group that was aliphatic and moderate in size.²⁰ With the furans, the most favorable substituents at this position were ethyl and propyl, whereas with the pyrazoles it was a propyl substituent. The best furan (RBA 140 for furan 15b) has considerably higher ER α binding affinity increasing with substituent size up to a point, beyond which further increase proves detrimental, was noted. Such trends are well precedented in both steroidal and non-steroidal systems, ^{40,41} and they have been interpreted to indicate the filling of a preformed pocket of limited volume within the receptor.⁴⁰

Where it is possible to make direct comparisons between furans and thiophenes (15m vs. 17a and 15n vs. 17b), it appears that the furans might be higher affinity ligands. However, this comparison remains limited by difficulties we encountered in the preparation of tetra-substituted thiophenes. We prepared only one pyrrole (19), and though it was tetrasubstituted, its affinity was disappointing. Because it has strong structural analogy to other heterocycles that have high affinity,²⁰ we have no explanation for this fact.

Investigation of the Orientation of the Triaryl Furans in the ER Ligand Binding Pocket

We have been interested in the manner in which non-steroidal ER ligands having more than one phenol are oriented within the ligand binding pocket of ER.⁴² Most of the furans we studied here that show high binding selectivity for ER α (15a-c) have three phenols, any one of which could be serving as the analog of the phenolic A-ring of estradiol, a ring that is known to be important for the high affinity of this natural estrogen (see Figure 5).⁴⁰ While multiple binding modes could be operative, we used comparative binding affinity considerations and molecular modeling to see whether we could determine which of these three phenols might most likely be functioning as the A-ring mimic. One should note that for each of the three potential A-ring mimics, there are two possible binding modes for the rest of the furan (i.e., the ligands could be bound in the orientations shown in Fig. 4, or in alternative binding modes, flipped 180° around the bond connecting the A-ring mimic phenol to the furan core, not shown, giving a total of six basic binding orientations). ⁴²

[Figure 5]

Comparative Binding Analysis: In the comparative binding affinity approach, we prepared furan bis-phenols (**15f,h,i**) and monophenols (**15j-I**) to investigate the effect that removing phenolic hydroxyl groups had on binding affinity. We imagined that deletion of the hydroxy group from the furan phenol substituent that was playing the role of the A-ring of estradiol would have the greatest effect on binding affinity.⁴⁰ Also, if two phenols were deleted, we imagined that the monophenol having the highest affinity would be the one that retained the hydroxyl on the phenyl substituent in the furan that mimics the estradiol A-ring. The ER α binding affinities of the mono and diphenols were given in Table 3, but are schematized here in Figure 6 for ready analysis.

[Figure 6]

In the monophenols (15j-l) (See RBA values shown in boxes in Figure 6) the C(4) monophenol (15k) has a very low affinity; so, it is unlikely that the C(4) aryl group is the mimic of the estradiol A-ring. However, it is difficult to choose between the C(2) and C(5) phenols (15j,l), because their affinities are quite comparable. So, from the monophenols, it appears that either the C(2) or C(5) phenol rings could be the A-ring mimic. The highest affinity bis-phenol (15f) (See RBA values shown in parentheses in Italics, Figure 6) has hydroxyl groups on both the C(2) and C(4) phenyl groups, consistent with the importance of the C(2) phenol, noted above.

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However, the other two bis-phenols (15h,i) again have very similar affinities; so, a definitive distinction cannot be made between the importance of the other two phenols, at C(4) and C(5).

Based on these data, we make the following tentative conclusions: (a) the C(4) phenol is unlikely to be the A-ring mimic (based on the affinity pattern of the monophenols), (b) the C(2) phenol is most likely to be the A-ring mimic (based on the higher affinity of the 2,4-bis-phenol (**15f**) than the 2,5- and 4,5-bis-phenols (**15h,i**), but (c) the C(5) phenol might also function as the A-ring mimic. This tentative conclusion is the same as the one we reached in our analysis of the binding orientations of the pyrazoles,²⁴ and it is supported by further work we have done, both in the pyrazole and the furan series, to develop ER α -selective antagonists by attaching basic side chains to these heterocyclic systems.^{25,43} Thus, we believe that the furan triphenols bind with the C(2) phenol in the A-ring binding pocket, but that the C(5) phenol can play this role in certain analogs (notably furan antagonists).⁴³

Molecular Modeling Studies: We have also used molecular modeling to investigate the binding orientation of a furan ligand in ER α . Using the FlexiDock routine in the modeling program SYBYL, according to a modification of the protocol we described previously for the pyrazoles (See Experimental),²⁴ we docked the ER α -selective furan **15b** into the ligand-binding pocket of ER α , positioning the furan initially in each of six possible orientations (Cf., Fig 5, and earlier discussion). In each case, the FlexiDock routine oriented the molecule so that the C(2) phenol remained or returned to the position so as to be the A-ring mimic. With the C(2) phenol in the A-ring orientation, we obtained the best fit when the C(3) ethyl group projected downward in roughly the direction of a 6 α or 7 α -substituent in estradiol. The result of this docking-minimization study is shown in Figure 7. The purple dots represent the solvent-accessible surface of the ligand binding pocket, and the green surface is the molecular volume for the furan ligand. At the right of the figure is an overlay of furan **15b** (gray), onto estradiol (purple), illustrating their relative position within the ligand binding pocket.

This final minimized model shows that the hydrogen bond contacts found between the phenol and the Glu353 and Arg394 residues in the ER α /estradiol and ER α /diethylstilbestrol

crystal structures^{44,45} persist with the C(2) phenol of the furan ligand. However, because of the overall greater length of the furan, we found that the phenol appears to be driven more deeply into the A-ring binding pocket in the furan structure than in the estradiol structure. The second phenol on C(4) of the furan makes a hydrogen bond with His524, as is also found with the distal hydroxyl groups of ligands in the crystal structures.^{44,45} The third phenol on C(5) appears to be making a hydrogen bond with the hydroxy group of Thr347, which is the only other polar residue in the ligand binding pocket. As was the case with the pyrazoles, it is not clear what interactions are responsible for the high ER α binding selectivity of this ligand, although this is obviously a very interesting issue.²⁴

[Figure 7]

The novel heterocyclic systems we have investigated here, the 3-alkyl-2,4,5-triarylfurans, in particular, are ligands for the estrogen receptor that have very high selectivity for ER α in terms of affinity and potency in transcription assays. These ligands should be useful in studying the biological role of both of the ER subtypes, and they might form the basis for the development of novel estrogen pharmaceuticals.

Experimental Section

General Methods

All reagents and solvents were purchased from Aldrich or Fischer. Solvents were distilled prior to use. THF was distilled over sodium/benzophenone. Methylene chloride was distilled over calcium hydride. Hexane was distilled over calcium sulfate. Triethylamine was distilled over calcium hydride. Reactions were all monitored by TLC, performed on 0.25 mm silica gel glass plates containing F-254 indicator (Merck). Visualization on TLC was achieved by UV light or phosphomolybdic acid indicator. Column chromatography was performed with Woelm 32-63 µm silica gel packing. Melting points were measured using a Thomas Hoover

capillary melting point apparatus and are uncorrected. NMR spectra chemical shifts are reported in parts per million downfield from TMS and referenced with either TMS internal standard for d_1 -chloroform or d_6 -acetone solvent peak. NMR coupling constants are reported in Hertz. All compounds used in structure-activity relationship considerations gave either satisfactory microanalyses or satisfactory exact mass determinations by high resolution mass spectrometry and were shown to be pure by HPLC under two distinct reversed phase conditions.

Chemical Synthesis

General Procedure for the Preparation of Trisubstituted Butane-1,4-diones by the Stetter Reaction. Aldehyde, α , β unsaturated ketone, triethlyamine, and either 3-benzyl-5-(2-hydroxyethyl)-4-methyl-thiazolium chloride (with aliphatic aldehydes) or 3-ethyl-5-(2-hydroxyethyl)-4-methyl-thiazolium bromide (with aromatic aldehydes) were combined and heated to reflux in ethanol for 60-96 h. Solvent was removed under reduced pressure, the resulting residue was taken up into CH₂Cl₂ and extracted with 3M HCl (aq.), water, sat. NaCl, and dried over sodium sulfate. Solvent was removed under reduced pressure and crude product was purified by flash column chromatography and recrystallization to afford 1,4 diones.

General Procedure for the Preparation of Tetrasubstituted Butane-1,4-diones A solution of KHMDS (1.1 equiv, 0.5 M in toluene) was added to a stirring solution of diarylethanone (1 equiv) in THF (2-5 mL), at -78 °C. The mixture was stirred 1 h followed by the dropwise addition of α -bromoketone (1.1 equiv). The reaction mixture was allowed to stir at -78 °C for 1 h and then allowed to gradually warm to rt, stirring 5 h to overnight (monitored for disappearance of starting material) and then quenched with the addition of H₂O. The reaction was further diluted with EtOAc and washed with H₂O and sat. NaCl. Organic extract was dried over Na₂SO₄, filtered and solvent removed under reduced pressure. Purification by flash column chromatography afforded diones as mixtures of diastereomers (by NMR).

General Procedure for Furans In toluene, the 1,4-dione and p-toluenesulfonic acid monohydrate (cat. amount) were heated to reflux for 3-12 h. The reaction was the cooled,

filtered and solvent removed under reduced pressure and the crude product was purified by flash column chromatography.

General Demethylation Procedure Using $BF_3 \cdot SMe_2$ To a stirring solution of the methyl ether precursor (1 equiv) in CH_2Cl_2 (~8 mL) at rt was added $BF_3 \cdot SMe_2$ complex (75 equiv). After stirring for 12-18 h, solvent and excess reagent were evaporated under nitrogen stream in hood. Residue was taken up in EtOAc and washed with H_2O and sat NaCl. Organic extract was dried over Na₂SO₄, filtered and solvent removed under reduced pressure. The resulting residue was purified by silica gel flash column chromatography.

General Demethylation Procedure Using BBr₃ The methyl ether protected compound was dissolved in CH_2Cl_2 and stirred at rt. A solution of 1M boron tribromide in CH_2Cl_2 was added via syringe, and the reaction was left to stir over-night or until all starting material had been consumed. The reaction poured into separatory funnel and extracted with H_2O (3x10 ml), sat NaCl, and dried over sodium sulfate. Solvent was removed under reduced pressure and the resulting crude product was purified by flash column chromatography and recrystallization to afford deprotected phenols.

2,3,5-Tris-(4-hydroxy-phenyl)-4-methyl-furan (15a) Furan **14a** (58.0 mg, 0.14 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15a**. The crude material was purified by flash column chromatography (1:2 EtOAc:hexanes) and recrystallized from EtOAc:hexanes to give **15a** (40.2 mg, 77% yield). mp 231-233 °C (decomp); ¹H NMR (500 MHz, acetone- d_6) δ 2.05 (3H, s, CH₃), 6.74 (2H, AA'XX', J_{AX} = 8.77, $J_{AA'}$ = 2.48, ArH ortho to OH), 6.94 (2H, AA'XX', J_{AX} = 8.46, $J_{AA'}$ = 2.42, ArH ortho to OH), 6.95 (2H, AA'XX', J_{AX} = 8.83, $J_{AA'}$ = 2.49, ArH ortho to OH), 7.16 (2H, AA'XX', J_{AX} = 8.81, $J_{XX'}$ = 2.39, ArH meta to OH), 7.34 (2H, AA'XX', J_{AX} = 8.99, $J_{XX'}$ = 2.50, ArH meta to OH), 7.61 (2H, AA'XX', J_{AX} = 8.77, $J_{XX'}$ = 2.46, ArH meta to OH), 8.44 (1H, bs, OH), 8.48 (1H, bs, OH), 8.51 (1H, bs, OH); ¹³C NMR (125 MHz, acetone- d_6) δ 10.5, 116.1(2), 116.4(2), 116.6(2), 117.5, 124.1, 124.6, 124.7, 125.8, 127.5(2), 127.7(2), 132.2(2), 147.4, 147.9, 157.4, 157.5, 157.7; MS (EI, 70 eV) *m*/*z* 358.1 (M⁺); HRMS calcd for C₂₃H₁₈O₄: 358.12051, found: 358.12040.

3-Ethyl-2,4,5-tris(4-hydoxyphenyl)furan (15b) Furan **14b** (28.0 mg, 0.07 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15b**. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) and recrystallized from EtOAc:hexanes to give **15b** (16.3 mg, 93% yield). mp 219-220 °C; ¹H NMR (500 MHz, acetone- d_6) δ 1.01 (3H, t, J=7.52, CH_3CH_2), 2.51 (2H, q, J= 7.52, CH_3CH_2), 6.72 (2H, AA'XX', J_{AX} = 8.99, $J_{AA'}$ = 2.52, Ar*H* ortho to OH), 6.95 (2H, AA'XX', J_{AX} = 8.85, $J_{AA'}$ = 2.54, Ar*H* ortho to OH), 6.96 (2H, AA'XX', J_{AX} = 8.67, $J_{AA'}$ = 2.55, Ar*H* ortho to OH), 7.17 (2H, AA'XX', J_{AX} = 8.89, $J_{XX'}$ = 2.41, Ar*H* meta to OH), 7.31 (2H, AA'XX', J_{AX} = 9.01, $J_{XX'}$ = 2.47, Ar*H* meta to OH), 7.60 (2H, AA'XX', J_{AX} = 8.95, $J_{XX'}$ = 2.49, Ar*H* meta to OH), 8.39 (1H, bs, O*H*), 8.46 (1H, bs, O*H*), 8.49 (1H, bs, O*H*); ¹³C NMR (125 MHz, acetone- d_6) δ 14.1, 17.1, 115.2(2), 115.6(2), 115.7(2), 123.2, 123.4, 123.5, 123.6, 125.1, 126.5(2), 126.8(2), 131.3(2), 146.5, 146.7, 156.5, 156.7, 156.9; MS (EI, 70 eV) *m/z* 372.2 (M⁺); HRMS calcd for C₂₄H₂₀O₄ : 372.136159, found: 370.136761; Anal. calcd for C₂₄H₂₀O₄ (+0.5 H₂O), C: 74.43, H: 6.12. Found, C: 74.82, H: 5.86.

2,3,5-Tris(4-hydroxyphenyl)-4-propylfuran (15c) Furan **14c** (40.0 mg, 0.09 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15c**. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) to provide **15c** (32.0 mg, 93% yield) as a white powder. ¹H NMR (500 MHz, acetone- d_6) δ 0.79 (3H, t, *J*= 7.24, *CH*₃CH₂CH₂), 1.42 (2H, m, CH₃CH₂CH₂), 2.48 (2H, m, CH₃CH₂CH₂), 6.73 (2H, AA'XX', *J*_{AX}= 8.89, *J*_{AA'}= 2.53, Ar*H* ortho to OH), 6.95 (4H, AA'XX', *J*_{AX}= 8.55, *J*_{AA'}= 2.53, Ar*H* ortho to OH), 7.16 (2H, AA'XX', *J*_{AX}= 8.82, *J*_{XX'} = 2.41, Ar*H* meta to OH), 7.31 (2H, AA'XX', *J*_{AX}= 8.86, *J*_{XX'} = 2.47, Ar*H* meta to OH), 7.60 (2H, AA'XX', *J*_{AX}= 9.02, *J*_{XX'} = 2.42, Ar*H* meta to OH), 8.42 (1H, bs, O*H*), 8.48 (1H, bs, O*H*), 8.51 (1H, bs, O*H*); ¹³C NMR (125 MHz, acetone- d_6) δ 14.4, 23.8, 26.8, 116.0(2), 116.4(2), 116.6(2), 122.9, 124.1, 124.4, 124.5, 126.1, 127.3(2), 127.7(2), 132.2(2), 147.5, 147.7, 157.4, 157.5, 157.7; MS (EI, 70 eV) *m*/*z* 386.2 (M⁺); HRMS calcd for C₂₅H₂₂O₄: 386.15181, found: 386.15274.

3-Butyl-2,4,5-tris(4-hydroxyphenyl)furan (15d) Furan **14d** (50.0 mg, 0.11 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15d**. The crude material was purified by flash column chromatography (1:1 EtOAc:hexanes) to provide **15d** (39.5 mg, 88% yield) as a solid. A small amount of **15d** for biological testing was further purified by reverse phase HPLC; ¹H NMR (500 MHz, acetone-*d*₆) δ 0.76 (3H, t, *J*= 7.41, CH₃CH₂CH₂CH₂), 1.22 (2H, sextet, *J*= 7.39, CH₃CH₂CH₂CH₂), 1.39 (2H, m, CH₃CH₂CH₂CH₂CH₂), 6.72 (2H, AA'XX', *J*_{AX}= 8.72, *J*_{AA'} = 2.53, Ar*H* ortho to OH), 6.949 (2H, AA'XX', *J*_{AX}= 8.83, *J*_{AA'} = 2.49, Ar*H* ortho to OH), 6.952 (2H, AA'XX', *J*_{AX}= 8.63, *J*_{AA'}= 2.53, Ar*H* ortho to OH), 7.16 (2H, AA'XX', *J*_{AX}= 8.67, *J*_{XX'} = 2.41, Ar*H* meta to OR), 7.31 (2H, AA'XX', *J*_{AX}= 9.11, *J*_{XX'} = 2.45, Ar*H* meta to OR), 7.60 (2H, AA'XX', *J*_{AX}= 9.01, *J*_{XX'} = 2.53, Ar*H* meta to OR), 8.47 (1H, bs, OH), 8.52 (1H, bs, OH), 8.56 (1H, bs, OH); ¹³C NMR (125 MHz, acetone-*d*₆) δ 13.9, 23.2, 24.3, 32.7, 116.0(2), 116.4(2), 116.6(2), 123.0, 124.1, 124.4, 124.5, 126.1, 127.3(2), 127.7(2), 132.2(2), 147.5, 147.6, 157.4, 157.5, 157.7; MS (EI, 70 eV) *m*/z 400.2 (M⁺); HRMS calcd for C₂₆H₂₄O₄: 400.16746, found: 400.16743.

2,3,4,5-Tetrakis-(4-hydroxy-phenyl)-furan (15e) Furan **14e** (40.0 mg, 0.10 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15e**. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) and recrystallized from EtOAc:hexanes to give **15e** (49.0 mg, 85% yield). mp 247-250 °C (decomp); ¹H NMR (500 MHz, acetone- d_6) δ 6.76 (4H, AA'XX', J_{AX} = 8.72, $J_{AA'}$ = 2.45, Ar*H* ortho to OH), 6.77 (4H, AA'XX', J_{AX} = 8.65, $J_{AA'}$ = 2.45, Ar*H* ortho to OH), 7.37 (4H, AA'XX', J_{AX} = 8.62, $J_{XX'}$ = 2.32, Ar*H* meta to OH), 8.38 (4H, AA'XX', J_{AX} = 8.73, $J_{XX'}$ = 2.44, Ar*H* meta to OH), 8.38 (2H, bs, O*H*), 8.53 (2H, bs, O*H*); ¹³C NMR (125 MHz, acetone- d_6) δ 116.08(4), 113.12(4), 123.9(2), 124.2(2), 125.5(2), 127.7(4), 132.4(4), 147.7(2), 157.4(2), 157.6(2); MS (EI, 70 eV) *m*/z 436.2 (M⁺); HRMS calcd for C₂₈H₂₀O₅: 436.13107, found: 436.13124.

3-Ethyl-2,4-bis(4-hydroxyphenyl)-5-phenylfuran (15f) Furan **14f** (20.0 mg, 0.05 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15f**. The crude material was purified by flash column chromatography (1:1

hexane:EtOAc) followed by recrystallization from CH₂Cl₂:hexanes to give **15f** as a solid (17.4 mg, 94 % yield). mp 226-230 °C; ¹H NMR (500 MHz, acetone- d_6) δ 1.02 (3H, t, J= 7.49, CH₃CH₂), 2.53 (2H, q, J= 7.49, CH₃CH₂), 6.97 (2H, AA'XX', J_{AX} = 8.74, $J_{AA'}$ = 2.60, Ar*H* ortho to OH), 6.98 (2H, AA'XX', J_{AX} = 8.70, $J_{AA'}$ = 2.45, Ar*H* ortho to OH), 7.16 (1H, m, Ar*H* para to furan), 7.19 (2H, AA'XX', J_{AX} = 8.75, $J_{XX'}$ = 2.48, Ar*H* meta to OH), 7.24 (2H, m, Ar*H* meta to furan), 7.46 (2H, m, Ar*H* ortho to furan), 7.64 (2H, AA'XX', J_{AX} = 8.66, $J_{XX'}$ = 2.53, Ar*H* meta to OH), 8.55 (2H, bs, O*H*); ¹³C NMR (125 MHz, acetone- d_6) δ 14.9, 17.9, 116.5(2), 116.7(2), 124.2, 124.6, 125.6(2), 125.7, 126.5, 127.5, 127.9(2), 129.2(2), 132.1(2), 132.3, 147.0, 148.4, 157.8, 157.9; MS (EI, 70 eV) *m*/*z* 356.1 (M⁺); HRMS calcd for C₂₄H₂₀O₃ : 356.14125, found: 328.14129.

2,4-Bis(4-hydroxyphenyl)-5-phenyl-3-propylfuran (15g) Furan **14g** (30.0 mg, 0.075 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15g**. The crude material was purified by flash column chromatography (2:1 hexane:EtOAc) and recrystallized from EtOAc:hexanes to give **15g** (23.2 mg, 86% yield). ¹H NMR (500 MHz, acetone- d_6) δ 0.80 (3H, t, *J*=7.39, CH₃CH₂), 1.44 (2H, m, CH₃CH₂CH₂), 2.49 (2H, m, CH₂-furan), 6.969 (2H, AA'XX', *J*_{AX}= 8.87, *J*_{AA'}= 2.55, Ar*H* ortho to OH), 6.972 (2H, AA'XX', *J*_{AX}= 8.55, *J*_{AA'} = 2.41, Ar*H* ortho to OH), 7.16 (1H, m, Ar*H* para to furan), 7.18 (2H, AA'XX', *J*_{AX}= 8.78, *J*_{XX'} = 2.44, Ar*H* meta to OH), 7.24 (2H, m, Ar*H* meta to furan), 7.45 (2H, m, Ar*H* ortho to furan), 7.64 (2H, AA'XX', *J*_{AX}= 8.68, *J*_{XX'} = 2.62, Ar*H* meta to OH), 8.52 (1H, bs, O*H*), 8.57 (1H, bs, O*H*); ¹³C NMR (125 MHz, acetone-*d*₆) δ 14.1, 23.6, 26.6, 116.3(2), 116.5(2), 123.0, 124.0, 125.4(2), 125.5, 126.5, 127.3, 127.7(2), 128.9(2), 131.9(2), 132.1, 146.7, 148.5, 157.6, 157.7; MS (EI, 70 eV) *m*/*z* 370.2 (M⁺); HRMS calcd for C₂₅H₂₂O₃ : 370.156890, found: 370.15661.

3-Ethyl-2,5-bis(4-hydroxyphenyl)-4-phenylfuran (15h) Furan 14h (30.0 mg, 0.08 mmol) was reacted according to the general demethylation procedure using $BF_3 \cdot SMe_2$ to afford crude 15h. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) to provide 15h (25.8 mg, 93% yield) as a white powder. mp 127-132 °C

(decomp); ¹H NMR (500 MHz, acetone- d_6) δ 1.00 (3H, t, J= 7.47, CH_3CH_2), 2.52 (2H, q, J= 7.50, CH_3CH_2), 6.72 (2H, AA'XX', J_{AX} = 8.86, $J_{AA'}$ = 2.48, ArH ortho to OH), 6.96 (2H, AA'XX', J_{AX} = 8.75, $J_{AA'}$ = 2.56, ArH ortho to OH), 7.27 (2H, AA'XX', J_{AX} = 8.91, $J_{XX'}$ = 2.47, ArH meta to OH), 7.36 (2H, m, ArH ortho to furan), 7.41 (1H, tt, J= 7.51, 1.34, ArH para to furan), 7.48 (2H, m, ArH meta to furan), 7.62 (2H, AA'XX', J_{AX} = 8.84, $J_{XX'}$ = 2.46, ArH meta to OH), 8.44 (1H, bs, OH), 8.54 (1H, bs, OH); ¹³C NMR (125 MHz, acetone- d_6) δ 14.9, 17.9, 116.1(2), 116.5(2), 123.8, 123.9, 124.0, 124.3, 127.5(2), 127.8(2), 128.3, 129.7(2), 131.1(2), 135.4, 147.6, 147.7, 157.5, 157.6; MS (EI, 70 eV) m/z 356.2 (M⁺); HRMS calcd for C₂₄H₂₀O₃ : 356.14125, found: 356.14168.

3-Ethyl-4,5-bis(4-hydroxyphenyl)-2-phenylfuran (15i) Furan **14i** (40.0 mg, 0.10 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15i**. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) and recrystallized from EtOAc:hexanes to give **15i** (31.3 mg, 84% yield). mp 204-206 °C; ¹H NMR (500 MHz, acetone-*d*₆) δ 1.04 (3H, t, *J*=7.53, C*H*₃CH₂), 2.57 (2H, q, *J*= 7.50, CH₃CH₂), 6.75 (2H, AA'XX', *J*_{AX}= 9.14, *J*_{AA'} = 2.46, Ar*H* ortho to OH), 6.97 (2H, AA'XX', *J*_{AX}= 8.59, *J*_{AA'} = 2.40, Ar*H* ortho to OH), 7.18 (2H, AA'XX', *J*_{AX}= 8.79, *J*_{XX'} = 2.32, Ar*H* meta to OH), 7.30 (1H, m, Ar*H* para to furan), 7.34 (2H, AA'XX', *J*_{AX}= 8.86, *J*_{XX'} = 2.47, Ar*H* meta to OH), 7.47 (2H, m, Ar*H* meta to furan), 7.77 (2H, m, Ar*H* ortho to furan), 8.52 (1H, bs, O*H*); ¹³C NMR (125 MHz, acetone-*d*₆) δ 14.8, 18.0, 116.1(2), 116.6(2), 123.8, 124.4, 125.7, 125.9(2), 126.6, 127.5(2), 127.6, 129.6(2), 132.2(2), 132.7, 146.8, 148.5, 157.7, 157.8; MS (EI, 70 eV) *m*/*z* 356.2 (M⁺); MS (CI, 130 eV) *m*/*z* 357.2 (M⁺+H); HRMS calcd for C₂₄H₂₀O₃: 356.14125, found: 356.14045.

3-Ethyl-2-(4-hydroxyphenyl)-(4,5)-bisphenylfuran (15j) Furan 14j (40 mg, 0.11 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude 15j. The crude material was purified by flash column chromatography (75:25 hexane:EtOAc) and recrystallized from hexane:EtOAc to give 15j (21 mg, 55% yield) as a white crystalline solid. mp 159-160 °C; ¹H NMR (400 MHz, CDCl₃) δ 1.04 (3H, t, *J* = 7.57, CH₃),

2.53 (2H, q, J = 7.51, CH_2), 4.76 (1H, s, OH), 6.93 (2H, d, J = 8.34, ArH ortho to OH), 7.13-7.47 (10H, m, ArH), 7.65 (2H, d, J = 8.75, ArH meta to OH); ¹³C NMR (500 MHz, CDCl₃) δ 14.6, 17.3, 115.6(2), 124.1, 124.8, 125.1(2), 126.7, 127.2(2), 127.4, 128.2(2), 128.8(2), 130.2(2), 131.1, 134.2, 146.6, 147.2, 154.6, 164.5; MS (EI, 70 EV) *m*/z 340.2 (M⁺); HRMS calcd for C₂₄H₂₀O₂: 340.14633, found: 340.14611.

3-Ethyl-4-(4-hydroxyphenyl)-(2,5)-bisphenylfuran (15k) Furan **14k** (42 mg, 0.12 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15k**. The crude material was purified by flash column chromatography (75:25 hexane:EtOAc) and recrystallized from hexane:EtOAc to give **15k** (28 mg, 70% yield) as a white powder. mp 144-147 °C ¹H NMR (400 MHz, CDCl₃) δ 1.11 (3H, t, *J* = 7.45, CH₃), 2.62 (2H, q, *J* = 7.36, CH₂), 4.92 (1H, s OH), 6.97 (2H, d, *J* = 8.21, ArH ortho to OH), 7.28-7.49 (10H, m, ArH), 7.8 (2H, d, *J* = 7.87, ArH meta to OH); ¹³C NMR (500 MHz, CDCl₃) δ 14.6, 17.4, 115.8, 125.1, 125.2(2), 125.4(2), 125.8, 126.4, 126.8, 126.9, 128.3(2), 128.6 (2), 131.1, 131.5(2), 131.6, 147.1, 147.2, 154.9; MS (EI, 70 EV) *m/z* 340.2 (M⁺); HRMS calc. for C₂₄H₂₀O₂: 340.14633, found: 340.14649.

3-Ethyl-5-(4-hydroxyphenyl)-(2,4)-bisphenylfuran (15I) Furan **141** (13.4 mg, 0.04 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15I**. The crude product was purified by flash column chromatography (95:5 hexane:EtOAc) and recrystallized from hexane:EtOAc to give furan **15I** (10 mg, 78% yield) as a light yellow powder. mp 133-136 °C ¹H NMR (400 MHz, CDCl₃) δ 1.06 (3H, t, *J* = 7.35, CH₃), 2.57 (2H, q, *J* = 7.48, CH₂), 4.65 (1H, s, OH), 6.69 (2H, d, *J* = 8.47, ArH ortho to OH), 7.26-7.46 (10H, m, ArH), 7.74 (2H, d, *J* = 7.26, ArH meta to OH); ¹³C NMR (500 MHz, CDCl₃) δ 14.6, 17.4, 115.2(2), 124.7, 124.3, 125.4(2), 125.5, 126.8, 127.0(2), 127.4, 128.5, 128.6(2), 128.8(2), 130.3(2), 131.7, 134.2, 146.6, 154.6; MS (EI, 70 eV) *m/z* 340.2 (M⁺); HRMS calcd for C₂₄H₂₀O₂: 340.14633, found: 340.14659.

2,5-Bis(4-hydroxyphenyl)-3-phenylfuran (15m) Furan **14m** (22.0mg, 0.06 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15m**.

The crude material was purified by flash column chromatography (1:1 EtOAc:hexane) followed by recrystallization from CH₂Cl₂:hexanes to give **15m** as a solid (18.7 mg, 92% yield). mp 167-170 °C; ¹H NMR (500 MHz, acetone- d_6) δ 6.82 (2H, AA'XX', J_{AX} = 8.77, $J_{AA'}$ = 2.32, Ar*H* ortho to OH), 6.83 (1H, s, furan*H*), 6.92 (2H, AA'XX', J_{AX} = 8.90, $J_{AA'}$ = 2.52, Ar*H* ortho to OH), 7.31 (1H, m, Ar*H* para to furan), 7.39 (2H, m, Ar*H* meta to furan), 7.43 (2H, AA'XX', J_{AX} = 8.817, $J_{XX'}$ = 2.45, Ar*H* meta to OH), 7.46 (2H, m, Ar*H* ortho to furan), 7.67 (2H, AA'XX', J_{AX} = 8.93, $J_{XX'}$ = 2.44, Ar*H* meta to OH), 8.56 (2H, bs, OH); ¹³C NMR (100 MHz, acetone- d_6) δ 107.3, 115.6(2), 115.9(2), 122.78, 122.81, 123.1, 125.4(2), 127.2, 127.8(2), 128.6(2), 128.8(2), 134.9, 147.5, 152.5, 157.36, 158.38; MS (FAB) m/z 328.2 (M⁺), HRMS calcd for C₂₂H₁₆O₃ : 328.10995, found: 328.10980.

3,5-Bis(4-hydroxyphenyl)-2-phenylfuran (15n) Furan **14n** (65.0 mg, 0.18 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **15n**. The crude material was purified by flash column chromatography (1:1 EtOAc:hexane) followed by recrystallization from CH₂Cl₂:hexanes to give **15n** as a solid (46.0 mg, 77% yield). mp 160-163 °C; ¹H NMR (400 MHz, acetone-*d*₆) δ 6.79 (1H, s, furan*H*), 6.89 (2H, AA'XX', *J*_{AX}= **8.65**, *J*_{AA'} = 2.47, Ar*H* ortho to OH), 6.93 (2H, AA'XX', *J*_{AX}= **8.82**, *J*_{AA'} = 2.45, Ar*H* ortho to OH), 7.23 (1H, m, Ar*H* para to furan), 7.30 (2H, AA'XX', *J*_{AX}= **8.67**, *J*_{XX'} = 2.53, Ar*H* meta to OH), 7.32 (2H, m, Ar*H* meta to furan), 7.66 (2H, m, Ar*H* ortho to furan), 7.69 (2H, AA'XX', *J*_{AX}= **8.81**, *J*_{XX'} = 2.45, Ar*H* meta to OH), 8.58 (2H, bs, OH); ¹³C NMR (100 MHz, acetone-*d*₆) δ 108.7, 116.5(2), 116.6(2), 123.4, 125.63, 126.3(2), 126.5(2), 128.0, 129.3(2), 130.7(2), 132.4, 147.0, 153.7, 157.9, 158.36 (2); MS (EI, 70 eV) *m*/*z* 328.2 (M⁺); HRMS calcd for C₂₂H₁₆O₃ : 328.10995, found: 328.10955.

General Procedure for Thiophenes The 1,4-dione and Lawesson's Reagent were stirred in CH_2Cl_2 at 40 °C until all 1,4-dione had been consumed by TLC (2-3 h). The reaction mixture was then poured into separatory funnel and washed with H_2O , 10% sodium bicarbonate, sat. NaCl, and dried over sodium sulfate. Solvent was removed under reduced pressure, and the

crude product was purified by flash column chromatography and recrystallization to afford cyclized thiophenes.

2,5-Bis(4-hydroxyphenyl)-3-phenylthiophene (17a) Thiophene **16a** (22.6 mg, 0.06 mmol) was reacted according to the general demethylation procedure using BBr₃ to afford crude **17a**. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) to give **17a** as a solid (18.3 mg, 88% yield). mp 125-130 °C; ¹H NMR (500 MHz, acetone- d_6) δ 6.78 (2H, AA'XX', J_{AX} = 8.73, $J_{AA'}$ = 2.47, Ar*H* ortho to OH), 6.90 (2H, AA'XX', J_{AX} = 8.78, $J_{AA'}$ = 2.60, Ar*H* ortho to OH), 7.14 (2H, AA'XX', J_{AX} = 8.52, $J_{XX'}$ = 2.47, Ar*H* meta to OH), 7.26 (1H, m, Ar*H* para to thiophene), 7.32 (4H, m, Ar*H* meta and ortho to thiophene), 7.34 (1H, s, thiophene*H*), 7.56 (2H, AA'XX', J_{AX} = 8.79, $J_{XX'}$ = 2.52, Ar*H* meta to OH), 8.54 (1H, bs, O*H*), 8.57 (1H, bs, O*H*); ¹³C NMR (125 MHz, acetone- d_6) δ 115.5(2), 115.8(2), 124.9, 125.5, 125.8, 126.7(2), 126.8, 128.3(2), 128.9(2), 130.3(2), 136.6, 136.9, 138.0, 141.9, 157.2, 157.4; MS (EI, 70 eV) *m/z* 344.1 (M⁺); HRMS calcd for C₂₂H₁₆SO₂ : 344.08710, found: 344.08621.

3,5-Bis(4-hydroxyphenyl)-2-phenylthiophene (17b) Thiophene **16b** (108.0 mg, 0.29 mmol) was reacted according to the general demethylation procedure using BF₃•SMe₂ to afford crude **17b**. The crude material was purified by flash column chromatography (1:1 hexane:EtOAc) followed by recrystallization from CH₂Cl₂:hexanes to give **17b** as a solid (85.1 mg, 85 % yield). mp 198-200 °C; ¹H NMR (500 MHz, acetone-*d*₆) δ 6.79 (2H, AA'XX', *J*_{AX}= 8.66, *J*_{AA'} = 2.47, Ar*H* ortho to OH), 6.90 (2H, AA'XX', *J*_{AX}= 8.80, *J*_{AA'} = 2.60, Ar*H* ortho to OH), 7.17 (2H, AA'XX', *J*_{AX}= 8.69, *J*_{XX'} = 2.47, Ar*H* meta to OH), 7.24-7.34 (5H, m, Ar*H* phenyl), 7.32 (1H, s, thiophene*H*), 7.57 (2H, AA'XX', *J*_{AX}= 8.72, *J*_{XX'} = 2.52, Ar*H* meta to OH), 8.50 (2H, bs, O*H*); ¹³C NMR (125 MHz, acetone-*d*₆) δ 115.3(2), 115.8(2), 125.3, 125.7, 126.8(2), 127.2, 127.9, 128.5(2), 128.8(2), 130.1(2), 134.7, 134.9, 139.1, 142.6, 156.7, 157.5; MS (EI, 70 eV) *m*/z 344.1 (M⁺); HRMS calcd for C₂₂H₁₆SO₂: 344.08710, found: 344.08620.

General Procedure for N-Substituted Pyrroles In toluene, the 1,4-dione, amine and *p*-toluene sulfonic acid monohydrate were heated to reflux for 24 h, using a Dean-Stark trap. The

reaction was cooled, filtered, and solvent removed in-vacuo. Crude product was purified by flash column chromatography and recrystallization to afford N-substituted pyrroles.

2-Ethyl-1,3-bis-(4-hydroxyphenyl)-5-phenylpyrrole (19) Methyl-ether protected pyrrole **18** (38.8 mg, 0.10 mmol) and boron tribromide (0.6 mL, 0.6 mmol) were reacted as outlined in general procedure for demethylation to give 29.0 mg brownish solid. Trituration with hexane gave product as off-white solid, (21.8 mg), 60.8% yield. ¹H NMR (500 MHz, acetoned6) δ 0.90 (3H, t, *J*=7.42, *CH*₃), 2.62 (2H, q, *J*=7.42, *CH*₂), 6.43 (1, s, pyrrole*H*), 6.88 (2H, AA'XX', *J*_{AX}= 8.84, *J*_{AA'} = 2.63, Ar*H* ortho to OH), 6.90 (2H, AA'XX', *J*_{AX}= 8.80, *J*_{AA'} = 2.76, Ar*H* ortho to OH), 7.07 (1H, m, Ar*H* para to pyrrole), 7.11 (2H, AA'XX', *J*_{AX}= 8.70, *J*_{XX'} = 2.73, Ar*H* meta to OH), 7.15 (4H, m, Ar*H* ortho and meta to pyrrole), 7.32 (2H, AA'XX', *J*_{AX}= 8.63, *J*_{XX'} = 2.51, Ar*H* meta to OH), 8.18 (1H, s, ArO*H*), 8.67 (1H, s, ArO*H*);. MS (EI, 70 eV) *m*/z 355.2 (M⁺). HRMS calcd. for C₂₄H₂₁NO₂: 355.1572, found: 355.1567.

Molecular Modeling

The protein structure used in the docking simulations was based on the X-ray crystallographic structure of the human estrogen receptor ligand binding domain bound to estradiol (entry lere in the Protein Data Bank). The crystal structure contains 3 homodimers with 244 residues each. For modeling purposes only one monomer (chain A) was chosen. The monomer contains several residues with missing atoms and five residues with alternate conformations. Missing atoms were added using InsightII, and one conformation was selected for each of the five residues. Several residues that are part of two loops were completely missing. These two loops were modeled based on the crystal structure of the wild type estrogen receptor ligand binding domain complexed to estradiol (entry 1qku) using InsightII. To eliminate bad contacts and constraints due to crystal packing and loop reconstruction, the protein was energy minimized for 1000 steps using the Powell algorithm in the presence of strong harmonic constraints on the backbone, followed by an additional 1000 steps without constraints. Minimization was done with the program SYBYL 6.6 and the MMFF94 force field. Furan **15b**

was docked into the minimized receptor using the FlexiDock routine, and the receptor-ligand complex put through a minimization protocol as previously described.²⁴

Biological Procedures

Relative Binding Affinities. Ligand binding affinities (RBAs) using lamb uterine cytosol as a receptor source were determined by a competitive radiometric binding assay using 10 nM [³H]estradiol as tracer and dextran-coated charcoal as an adsorbant for free ligand.³⁸ Purified ER α and ER β binding affinities were determined using a competitive radiometric binding assay using 10 nM [³H]estradiol as tracer, commercially available ER α and ER β preparations (PanVera Inc. Madison, WI), and hydroxylapatite (HAP) to adsorb bound receptor-ligand complex.³⁹ HAP was prepared following the recommendations of Williams and Gorski.⁴⁶ All incubations were done at 0 °C for 18-24h. Unlabeled competitors were prepared in 1:1 DMF:TEA to ensure solubility. Binding affinities are expressed relative to estradiol on a percent scale (i.e., for estradiol, RBA = 100%). All essays were run in separate, duplicate experiments, which were reproducible with a coefficient of variation of less than 0.3.

Transcriptional Activation Studies. Transactivation by ligands on ER α and ER β was tested in transfected human endometrial cancer (HEC-1) cells. HEC-1 cells, maintained in MEM containing 5% CS and 5% FCS, were seeded into 24-well plates in transfection media (IMEM containing 5% FCS, and transfected at about 50% confluency, using lipofectin-transferrin. For each well, 1 µg 4ERE-TATA-LUC, 2.5 µg of pCMV β Gal as internal control, 50-100 ng pCMV5-ER α or pCMV5-ER β were mixed with 5 µl lipofectin (GIBCO, BRL) and 1.6 µl lmg/ml transferrin in 150 µl HBSS. The mixture was applied to the cell with 350 µl serum-free IMEM media for each well. The cell was incubated at 37° C in the 5% CO₂ containing incubator for 6 h. Compounds were prepared as solutions in ethanol and were added to medium to give a final ethanol concentration of 0.1%. The cell culture media was replaced by transfection media containing different concentration of ligands. Cell were incubated for 24 h in the presence of

ligand. The luciferase reporter assay system (Promega) was used for the luciferase activity assay. The activity of E_2 (10⁻⁸ M) on ER α or ER β was set as 100%, and the relative activity was adjusted based on the transfection efficiency which was monitored by the internal control β -galactosidase activity, as previously described.^{16,47}

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Supporting Information Available:

Experimental detail for all precursors is available.

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Table 1. Synthesis of furans 14a-n.

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Dione	Furan	R ¹	R ²	R ³	R ⁴	Yield
13a	14a	OCH ₃	OCH ₃	CH3	OCH ₃	74%
13b	14b	OCH ₃	OCH ₃	C ₂ H ₅	OH	70%
13c	14c	OCH ₃	OCH ₃	C_3H_7	OCH ₃	85%
13d	14d	OCH ₃	OCH ₃	C ₄ H ₉	OCH ₃	98%
13e	14e	OCH ₃	OCH ₃	pCH ₃ OC ₆ H ₄	OCH ₃	85%
13f	14f	Н	OCH ₃	C_2H_5	OCH ₃	87%
13g	14g	Н	OCH ₃	C ₃ H ₇	OCH ₃	83%
13h	14h	OCH ₃	Н	C_2H_5	OCH ₃	95%
13i	14i	OCH ₃	OCH ₃	C_2H_5	Н	73%
13j	14j	Н	Н	C ₂ H ₅	OCH ₃	42%
13k	14k	Н	OCH ₃	C_2H_5	Н	75%
131	141	OCH ₃	Н	C_2H_5	Н	25%
10a	14m	OCH ₃	Н	Н	OCH ₃	88%
10b	14n	Н	OCH ₃	Н	OCH ₃	92%

Table 2. Demethylation of furans 14a-n to 15a-n.

F		R ⁴	BF3•SMe2 CH2Cl2	R ¹ R ²	-0 	R ⁴
Furan	Deprot. Furan	\mathbf{R}^{1}	R ²	R ³	R ⁴	Yield
14a	15a	OH	ОН	CH ₃	ОН	77%
14b	15b	ОН	OH	C ₂ H ₅	ОН	93%
14c	15c	ОН	ОН	C_3H_7	ОН	93%
14d	15d	OH	ОН	C ₄ H ₉	OH	88%
14e	15e	OH	OH	pHOC ₆ H ₄	OH	85%
14f	15f	Н	ОН	C_2H_5	ОН	94%
14g	15g	Н	OH	C_3H_7	OH	88%
14h	15h	OH	Н	C_2H_5	OH	93%
14i	15i	OH	OH	C_2H_5	Н	84%
14j	15j	Н	Н	C_2H_5	OH	55%
14k	15k	Н	OH	C_2H_5	Н	70%
141	151	OH	Н	C_2H_5	Н	78%
14m	15m	OH	Н	Н	OH	75%
14n	15n	Н	OH	Н	OH	77%

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Ligand	Cytosol	ER-a	ER-β	α/β Selectivity
15a	2.9 ± 0.09	40 ± 6.5	0.62 ± 0.02	65 fold
15b	6.5 ± 0.8	140 ± 38	2.9 ± 0.1	48 fold
15c	$\textbf{3.6} \pm \textbf{0.6}$	100 ± 14	1.8 ± 0.65	56 fold
15d	1.3 ± 0.7	21 ± 0.6	3.9 ± 1.1	5.4 fold
15e	$\textbf{0.84} \pm \textbf{0.07}$	8.7 ± 1.1	0.25 ± 0.01	36 fold
15f	4.6 ± 0.5	82 ± 20	7.1 ± 1.2	12 fold
15g	3.6 ± 0.3	140 ± 13	15 ± 4.1	9.5 fold
15h	0.7±0	16.5 ± 1.9	3.0 ± 0.6	5.5 fold
15i	0.45 ± 0	14.8 ± 3.1	4.5 ± 1.2	3.3 fold
15j	0.44 ± 0.09	10.8 ± 2.6	3.4 ± 1.2	3.8 fold
15k	$\boldsymbol{0.02\pm0.01}$	0.15 ± 0.01	$\textbf{0.07} \pm \textbf{0.02}$	2.1 fold
151	$\textbf{0.10} \pm \textbf{0.04}$	6.8 ± 2.1	2.0 ± 0.4	3.4 fold
15m	0.13	NA	NA	NA
15n	0.04	NA	NA	NA
17a	0.03	NA	NA	NA
17b	0.04	NA	NA	NA
19	0.50	NA	NA	NA

Table 3. Relative Binding Affinity (RBA)^a data for furans 15a-n, thiophenes17a-b, and pyrrole 19.

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^aDetermined by a competitive radiometric binding assay with [³H]estradiol using methods described in the Experimental Section. Where indicated, values represent the average (\pm SD or range) of multiple determinations. In our hands, RBA values are reproducible with a coefficient of variation of less than 0.3.



Figure 1. Structures and binding affinity data pyrazoles 1 and 2.



3 $R^{1}=R^{2}=H$, $R^{3}=OCH_{2}CH_{2}NCH_{2}(CH_{2})_{2}CH_{2}$, $R^{4}=C_{6}H_{5}$ **4** $R^{1}=R^{2}=R^{4}=H$, $R^{3}=OH$ **5** $R^{1}=R^{3}=R^{4}=H$, $R^{2}=OCH_{2}CH_{2}NCH_{2}(CH_{2})_{2}CH_{2}$





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Figure 3. Transcriptional activation data for furans **15a-c** and **15g** through ER α and ER β . Human endometrial cancer (HEC-1) cells were transfected with expression vectors for ER α or ER β and an estrogen responsive reporter gene and were treated with indicated concentrations of estradiol or ligand for 24h to assay for agonism (diagonal slashed bars) or antagonism (cross-hatched bars). Transcriptional activity was normalized to an internal β -gal control plasmid using the luciferase reporter assay system. Values are expressed as a percent of the ER α or ER β response with 10 nM E₂, which is set at 100% (open bars).²⁹ For details, see Experimental Section.



Figure 4. Dose-response curve for transcriptional activation by furan 15b though ERα (solid squares) and ERβ (solid triangles). Antagonist activity through ERβ (circles) was also assayed. The activation curves for estradiol on those two receptors is shown for reference. (open squares and triangles, respectively; dotted lines) For details, see Experimental Section.



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Figure 5. Three possible mimics for the A-ring of estradiol. Note, with each case A-ring mimic, there are two possible orientations of the furan core (only one is shown).



Figure 6. Summary of ER α binding data for furan monophenols and bis-phenols for comparative binding analysis. The binding affinity of the three monophenols is shown in the shadowed box by each of the three hydroxyl groups, and the affinity of the three bis-phenols is indicated by the italicized number in parentheses on braces linking two hydroxyl groups. For a discussion, see text.



Figure 7. (A) Model of furan 15b in ER α ligand binding pocket. The surface of the ligand is shown as a continuous green shape; the surface of the ER α pocket is shown as purple dots. (B) Comparison of the orientation of the furan (gray) with respect to that of estradiol (purple) in the ER α ligand binding pocket. For details, see text and Experimental Section.



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Scheme 1. Preparation of 1,2,4-trisubstituted diones 10a-c.



Scheme 2. Preparation of 1,2,3,4-tetrasubstituted diones 13a-l.

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Scheme 3. Preparation of thiophenes 17a-b.



Scheme 4. Preparation of pyrrole 19.

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Furans with Basic Side Chains: Synthesis and Biological Evaluation of a Novel Series of Antagonists with Selectivity for the Estrogen Receptor Alpha

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Abstract – 3-Alkyl-2,4,5-triarylfurans with basic side chain substituents were prepared as ligands for the estrogen receptor. Those analog having the basic side chain on the C(4) phenol were high affinity, $ER\alpha$ -selective antagonists.

We have recently described the synthesis and biological evaluation of a series of 3-alkyl-2,4,5-triaryl-substituted furans, several of which proved to be ligands for the estrogen receptor (ER) with very high selectivity for ER α over ER β , both in terms of binding affinity and potency of transcriptional activation.¹ Because there is great interest in the development of *antagonists* as well as *agonists* that are ER subtype-selective,^{2,3} we have investigated a way to convert these ER α -selective furan agonists into ER α -selective antagonists.

The antagonist character of the antiestrogens raloxifene and hydroxy-tamoxifen (now more properly designated "selective estrogen receptor modifiers" or SERMs)⁴ relies on a tertiary amino ethoxy substituent, termed a basic sidechain (BSC). When these ligands are bound to $ER\alpha$, this bulky BSC projects outward from the ligand binding pocket and effects a structural reorganization of the receptor surface, repositioning helix 12 in a manner that blocks the binding of coactivator proteins important in mediating transcriptional activity.^{5,6} Thus, we surmised that we might be able to impart antiestrogenic character to the furan series of agonists ligands, by appending a BSC through one of the three phenols of our furan ligands (Figure 1). It is not immediately evident which attachment site would be preferred, nor that the product BSC-furans would retain the ERa selectivity of the parent furans, because neither raloxifene nor tamoxifen is an ER subtype selective antagonist. We were able to adapt the synthetic route originally used to prepare the tetra-substituted furan systems for the preparation of the BSC-analogs,¹ by differential protection of the phenol functions in the starting materials. We recently used a similar approach to attach BSC groups to pyrazoles.² Here we report the results of our efforts to identify the preferred furan position for attachment of the BSC, to obtain ER antagonists that have some selectivity for ER α over ER β , and to elucidate the orientation with which these furans bind to the receptor.



Figure 1. Three potential antiestrogenic furan basic side chain (BSC) derivatives.

Results and Discussion

Synthesis of Ligands

Treatment of the differentially protected 1,2-diarylethanones 1-2, ketone 3, or desoxyanisoin (4) with potassium hexamethyl-disilyl-amide (KHMDS), followed by the addition of an α -bromoketone 5a-e, provided the desired, differentially protected 1,3,4-triaryl-2-alkylbutane-1,4-diones 6a-g in 66-98% yields (Scheme 1). Although these diones were mixtures of diastereomers, no separation was required, because these stereocenters become trigonal in the furan products.



Scheme 1. Synthesis of basic side-chain furans 9a-f (BSC = CH_2CH_2 -piperidine).

Diones **6a-f** were refluxed in toluene with a slight excess of *p*-toluenesulfonic acid (TsOH) to effect furan formation and desilylation, giving mono-deprotected furans **7a-f**. The piperidinyl-ethoxy side-chain was added using Mitsunobu chemistry,¹ and the intermediates **8a-f** were demethylated using AlBr₃/EtSH to afford BSC-furans **9a**f.⁴

In contrast to the behavior of diones **6a-f**, when refluxed with a catalytic amount of TsOH in toluene, dione **6g** cyclized to the furan but did not undergo TIPS deprotection (Scheme 2). However, treatment of furan 10 with tetrabutylammonium fluoride at 0 °C did result in desilylation, giving the mono-deprotected furan 11 that was converted to BSC-furan 13, as described above.

Biological Studies

Binding Affinity and Ligand Orientation Studies: The affinity of the BSC-furans **9a-f** and **13** for the estrogen receptor was assayed by a competitive radiometric binding assay, using purified, full-length human ER α and ER β from recombinant sources.^{7,8} The results of these assays are expressed as relative binding affinity (RBA) values, where the affinity of estradiol is considered to be 100%, and they are summarized in Table 1.

By comparison of the ER α affinities for the isomeric series of 3-ethyl-substituted BSC-furans 9b, 9f, and 13, it appears that the preferred position for the attachment of the BSC is on C(4) phenol (9b); much lower affinities were observed when it was attached to the C(2) or C(5) phenols (13 and 9f, respectively). The BSC-modified furans did retain much of the selective affinity for ER α shown by their parent furans,¹ the most selective BSC-furan 9a having a 67-fold ER α /ER β binding selectivity. Even the analogs with low affinities for ER α maintained some of this selectivity.

The initial series of BSC-furans we prepared have two free phenols, either of which could be serving as the analog of the A-ring of estradiol, a group that functions as a crucial hydrogen-bonding partner in the ligand binding pocket and



Scheme 2. Synthesis of basic side-chain furan 13.

 Table 1. Estrogen receptor relative binding affinity^a data for basic sidechain furans 9a-f and 13.

Ligand	ER-a	ER-β	α/β
9a	32 ± 9	0.47 ± 0.1	67.2 fold
9b	75 ± 20	3.1 ± 0.1	24.3 fold
9d	60 ± 1	20 ± 3.5	3.0 fold
9e	3.8 ± 0.9	0.43 ± 0.04	8.8 fold
9f	2.5 ± 0.3	0.91 ± 0.02	2.7 fold
13	0.36 ± 0.11	0.06 ± 0.02	6.1 fold

^aDetermined by a competitive radiometric binding assay with [³H]estradiol.

is essential for the high affinity of the natural estrogen.⁹ We sought to determine which of these phenols in the C(4)-BSC-furans was playing this important role in binding by deleting each of them in turn, and then evaluating the effect that this had on their ER α binding affinity.⁹



Figure 2. Two possible A-ring mimics of estradiol for furan 9b.

The X-ray crystal structures for ER α complexed with either raloxifene⁶ or 4-hydroxy-tamoxifen⁵ show that the BSC on these ligands have a common orientation that places them in roughly in the 11 β direction with respect of a standard steroid core, where a salt bridge can then form between the teriary amine and aspartate 351 in helix-3. Thus, as indicated in Figure 2, there are likely to be only two binding modes for a C(4)-BSC-furan, such as 9b, mode A, with the C(2) phenol in the A-ring binding pocket, and mode B, with C(5) phenol mimicking the A-ring.

Comparison of the ER α binding affinity of BSC-furans 9d and 9e, mono-phenol analogs of the high affinity BSCfuran diol 9b (Table 1), clearly shows that the phenol attached at position C(5) serves as the A-ring mimic: There was a 20-fold loss of affinity when this phenol deleted (9e), whereas deletion of the C(2) phenol caused less than a 1.3-fold drop (9d). Thus, we conclude that the highest affinity furan with the basic side chain, ligand 9b, binds to ER in orientation B (Figure 2), and that in this orientation, having a second phenolic hydroxyl on the C(2) phenyl ring, does not seem to be very important for high affinity binding (9b vs 9d); this second phenol does, however, play a significant role in the selective affinity of these BSCfurans for ER α .

Transcriptional Activation Assays: We selected the BSCfurans that combined high affinity with good ER α binding selectivity (namely 9a-c) to be assayed for their capacity to activate transcription through either ER α or ER β , or to antagonize the transcriptional activity of estradiol through these ERs. These assays were done by cotransfection in human endometrial carcinoma (HEC-1) cells using a luciferase reporter gene system;¹⁰ dose-response studies with the BSC-furans alone were conducted to measure agonist activity, and in the presence of 1 nM estradiol to determine antagonist activity. The results are summarized in Figures 3 and 4.

As expected, none of the three BSC-furans showed appreciable agonist activity on either ER α or ER β . The methyl analog (9a) was a relatively weak antagonist on both ERs, showing no selectivity; the propyl analog (9c) was somewhat more potent and showed a small degree of selectivity in antagonizing ER α in preference to ER β . The most potent and selective antagonist was the ethyl BSC-furan, 9b, and a more complete dose response curve for this compound (Figure 4) shows that at 0.1 μ M, it almost fully suppresses the transcriptional activity of estradiol through ER α , yet it has no effect on ER β at this concentration. The IC₅₀ values for 9b on ER α and ER β are approximately 6.5 x 10⁻⁸ M and 4.8 x 10⁻⁷ M, respectively, which corresponds to a nearly 10-fold antagonist potency selectivity for ER α over ER β .

We have shown that one can obtain an ER α selective antagonist by adding a basic side chain (BSC) to the C(4) phenol of 3-ethyl-2,4,5-tris(4-hydroxylphenyl)furan, a ligand we have previously shown to be a very selective ligands.¹² However, the BSC-pyrazoles retained a greater degree of ER α -subtype selectivity in terms of both affinity and antagonist potency than did the BSC-furans.¹¹ Through binding affinity correlations, we concluded that



Figure 3. Transcriptional activity of the BSC-furans 9a-c and their antagonism of estradiol activity through ERa



Figure 4. Full dose-response, showing that the ER α -selective antagonism of BSC-furan 9b.

these BSC-furans bind to ER α with the C(5) phenol playing the role of the important A-ring of estradiol; we had reached a similar conclusion with the BSC-pyrazoles.³ It is of note in both the furan and pyrazole series that the parent ligands appear to bind in a different orientation than do the BSC-derivatives, placing the C(2) phenol of the furan (and the corresponding phenol of the pyrazole) in the A-ring binding pocket. The BSC derivatives of ER α selective ligands should prove to be useful agents to investigate the biological functions of the two ER subtypes.

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Abstract

ANTAGONISTS SELECTIVE FOR ESTROGEN RECEPTOR-ALPHA

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By adding basic side chains typically found in non-steroidal antiestrogens, to pyrazoles that are very selective estrogen receptor alpha (ER α) agonists, we have developed compounds that are high affinity, potent antagonists on ER α , and are essentially inactive on ER β . When such compounds are used together with estradiol, which activates both ER α and ER β , one can achieve selective stimulation of ER β . These compounds should be useful in studying the biological functions of ER β and in selectively blocking responses that are mediated through ER α .

Introduction

The action of estrogens in regulating gene transcription is mediated through two estrogen receptor subtypes, $ER\alpha$ and $ER\beta$.^{1.2} While both of these receptors function as ligand-modulated transcription factors, their tissue distributions are quite different.^{1.2} and the activity of estrogens of varying structure is not always the same on the two subtypes.^{3.4} Thus, it is likely that the biological roles of ER α and ER β are different, but the nature and importance of these differences are still open questions.

The generation of ER α and ER β knockout animals has provided much information about the function of these ER subtypes.⁵ Another approach has been the development of ER subtype-selective ligands. Coming to a good understanding of the key biological roles of ER α and ER β will be important for understanding the action of natural estrogens and for developing synthetic selective estrogen receptor modifiers (SERMs) ^{6,7} that are needed to optimally regulate fertility and menopausal hormone replacement, and to prevent and treat hormone-responsive breast cancer. We have developed compounds that are capable of stimulating ER α very selectively.

Members of the triary lpyrazole class, such as propylpyrazole triol (**PPT**), are more than 1000-fold

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more potent on ER α than on ER β .⁸ and certain tetrahydrochrysenes. such as R.R-diethyl-THC that we have developed. are nearly full to full agonists on ER α but pure antagonists on ER β .^{9.10} These two ligands could be classified as "ER α potency selective" and "ER α efficacy selective" compounds. respectively; in both cases, however, these compounds activate ER α in preference to ER β .

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From what has been published, it appears that it is more difficult to develop ligands that stimulate ER β to a greater extent than ER α . Certain phytoestrogens, such as genistein, are somewhat more potent on ER β than ER α .^{3,4} but their selectivity towards ER β is more limited than is the selectivity of **PPT** towards ER α , for example.⁸ We have also developed some synthetic ligands, bisbenzylnitriles and related compounds, that have up to 50-fold potency selectivity on ER β .¹⁰ However, as far as we are aware, no compounds thus far described have higher efficacy on ER β than ER α , that is, are more agonistic on ER β than they are on ER α .

In this report, we take a different approach to the development of ER β -selective ligands, the investigation of *antagonists that are more potent on ER\alpha than on ER\beta.* Such compounds, when used together with a subtype non-selective agonist such as estradiol, should result in selective ER β stimulation. because they would block the agonism of estradiol on ER α , but not on ER β . The compounds we have investigated are derivatives of the very ER α potency-selective triaryl-substituted pyrazole ligands, of which **PPT** is an example, into which we have introduced basic side chain (BSC) substituents, typical of those found in antiestrogens such as tamoxifen and raloxifene. In this report, we describe the transcriptional activity of these BSC-pyrazoles and the optimization of their selectivity as ER α potency-selective antagonists.

Results

Structure and Synthesis of Basic Side Chain Pyrazoles – In earlier reports, we identified a single position on the triarylpyrazoles where the BSC substitution is well tolerated by the ERs, namely, the C(5) phenol group,¹¹ and we developed a regioselective route to prepare these compounds efficiently.¹²

The structures of the BSC pyrazoles that we have studied are shown in **Figure 1** (See Fig. 1 legend for a simple nomenclature nemonic for these pyrazole systems). Those that we prepared first (**I**-(**A**-**G**)-**Et**) exemplified various basic side chains placed on our initially discovered 4-ethyl-triaryl ER α -selective pyrazole **I**.^{8,13} Once we had identified the most promising basic side chains (**A**, **C**, and **E**, see below), we explored other C(4) substituents on this core system (methyl (**A**,**C**,**E**) and in one case propyl (**A**).

ERa and ERB Binding Selectivity of Basic Side Chain Pyrazoles – The binding affinities of the BSC-pyrazoles for ER α and ER β , determined in a radiometric competitive binding assay.¹⁴ are listed in Table 1. All of the affinities are expressed as <u>Relative Binding Affinity</u> (RBA) values. where the RBA of estradiol is 100. As seen in Table 1, all of the compounds investigated had ER α binding affinity selectivity. with some showing ER α /ER β selectivity ratios of 100-400. It is instructive to compare the ER α binding affinity and selectivity of these BSC-pyrazoles to those of the parent triaryl pyrazoles (i.e., R = H instead of the basic side chain): The free phenols have ER α RBA values of 5.4 to 51 and ER α /ER β ratios of 80 to 390; interestingly, the BSC analogs can reach similar levels of ER α selectivity (ER α /ER β ratio of 220), and they show similar binding affinity for ER α . Although we have not explored these series exhaustively, where comparisons can be made, the ER α /ER β affinity ratio reaches a maximum with a smaller C(4) substituent in the BSC analogs

than in the triphenol series (compare pyrazole triols I, where the propyl analog **PPT** has the highest ratio, with BSC-pyrazoles I-A, where the methyl analog has the highest). This may reflect differences in the way in which triphenolic and BSC-pyrazoles are oriented in the ligand binding pockets of the ERs (see Discussion).

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ERa and ER β Transcriptional Selectivity of Basic Side Chain Pyrazoles – To evaluate which basic side chains were effective in converting the pyrazoles into ER antagonists, we first screened the activity of the seven BSC-pyrazoles having C(4)-ethyl substituents as agonists or as antagonists of estradiol-induced transcription through ER α and ER β by a cotransfection assay, using a simple single-concentration regimen. Expression plasmids for ER α and ER β (pCMV5-ER α and pCMV5-ER β), together with an estrogen-responsive reporter gene plasmid (3ERE-pS2-CAT) containing three estrogen response elements and the promoter from the pS2 gene, and a control plasmid (pCMV5- β gal), were transfected into human endometrial cancer (HEC-1) cells. The results of these single-concentration assays are summarized in Figure 2.

None of the seven C(4)-ethyl BSC-pyrazoles was a transcriptional agonist on either ER α or ER β ; all were transcriptional antagonists. Several of them showed considerable ER-subtype selectivity, being more potent on ER α , and we selected three of these (piperidinyl (A), pyrrolidino (C), diethylamino (E)) for more extensive dose-response studies, together with their analogs having different C(4)-alkyl substituents. These data are shown in Figures 3 and 4.

Antagonist dose response curves for the three BSC-pyrazoles with C(4)-ethyl and C(4)methyl substitutents are shown in Figures 3A and 3B, respectively. Comparison of the C(4)-ethyl series (Fig. 3A) with the C(4)-methyl series (Fig. 3B), shows, in each case, that the methyl series is more selective. Within the C(4)-methyl series (Fig. 3B), the piperidinyl analog (A) appears to be both a more potent and a more complete antagonist with ER α selectivity than the two other analogs.

Thus, of these six compounds, **I-A-Me** appears to be most selective as an ER α antagonist (cf. Fig. 3B). A final evaluation of piperidinyl congeners (A) having three different C(4) substituents (methyl, ethyl and propyl), shown in Figure 4, demonstrates again that in terms of both antagonist potency and ER α selectivity, the best compound compound is **I-A-Me**. It is of note that, overall, there is a reasonable correlation between the selectivity of these BSC pyrazoles for ER α in terms of their binding affinity and their potency as antagonists.

The activity of the most ER α -selective antagonist, pyrazole **I-A-Me**, that we have termed **MPP**. as an abbreviation for "methyl-piperidino-pyrazole", was evaluated further in a different estrogen-responsive reporter gene construct, NHERF/EBP50-Luc (which containing the 5'-flanking region of the sodium-hydrogen exchanger regulatory factor/ezrin-radixin-moesin binding protein 50; Fig. 5).^{15,16} Even at high (1 μ M) concentrations, **MPP** had no stimulatory activity on ER α , and it fully inhibited ER α activity by estradiol in a concentration-dependent manner, while having little to no suppressive activity on ER β stimulation by estradiol.

To evaluate the ability of **MPP** to suppress ER α stimulation of an endogenous gene. we monitored its effectiveness in blocking estradiol induction of pS2 mRNA production in MCF-7 breast cancer cells, which are known to contain high levels of ER α and minimal if any ER β . As seen in Figure 6, **MPP** was effective in suppressing pS2 mRNA induction by 1 nM estradiol. Full suppression required rather high concentrations of MPP (10 μ M); however, at this concentration. induction was suppressed to the control level and was equivalent to the suppression by 1 μ M of ICI 182.780. Thus, MPP is effective in blocking the stimulation of endogenous genes by estradiol through ER α .

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Discussion

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Typical estrogen antagonists, such as tamoxifen (Nolvadex^{*}), raloxifene (Evista^{*}), and ICI 182.780 (Faslodex^{*}), block estrogen action with equal or rather equal potency through the two estrogen receptor subtypes. ER α and ER β . We have been able to develop estrogen antagonists that are ER α selective by starting with triarylpyrazoles, a class of ER agonists that we have previously shown to be highly ER α -selective agonists, by introducing basic side chain (BSC) substituents on the C(5) phenol. These BSC-pyrazole antagonists all maintain significant affinity and potency selectivity for ER α , which in the best case (MPP) is very high. Thus, as shown in Figure 3 and 4, with MPP at 1 μ M we can antagonize the effect of 1 nM estradiol in activating ER α essentially completely, with only ca. 15% inhibition of the activation of ER β . The development of these antagonists that are more potent on ER β represents a significant advance in the preparation of ER automore these compounds are used together with estradiol, they would allow estendial stimulation only via ER β .

Within the BSC pyrazole series, there is a reasonably good correlation between their ER α selectivity in terms of binding affinity vs antagonist potency. What is interesting in this comparison is that the C(4)-alkyl substitutent that gives the highest selectivity for the basic side chain *antagonists*, namely a *methyl* group as in MPP, is smaller than the C(4)-*propyl* group needed to get the highest ER α /ER β selectivity in the triarylpyrazole *agonist* series. We believe that this is due to differences in how these two classes of ligands are oriented in the ligand binding pocket of ER α , 8.11.17 so that this alkyl substituents project into different regions of the pocket in the triaryl pyrazoles than in the BSC pyrazoles. As a result, groups of different size are required to obtain the highest selectivity in binding in the agonist series.

It is of note that we also prepared BSC analogs of pyrazoles in an isomeric series (i.e., 1.3.4triary1-5-alky1, with the BSC on the 4-ary1 group: not shown)¹⁷. As was true for parent pyrazoles in this series, the corresponding BSC derivatives were selective for ER α in terms of affinity and antagonist potency, but they are not as selective as **MPP** (data not shown).

The availability of a compound such as the ER α -selective antagonist MPP provides a new tool for investigating the biological function of the two ER subtypes. It can be used by itself to selectively turn off or block responses that might be mediated through ER α , or it can be used together with an ER-subtype non-selective agonist, such as estradiol, to achieve selective stimulation of ER β . This compound could also be paired with an ER β agonist, even if the latter is only moderately selective. to achieve highly selective stimulation via only ER β . We anticipate that this compound will be useful as an agent to investigate the biological functions of the less well understood estrogen receptor, ER β , in diverse cells and tissues in which ER β is expressed.

Materials and Methods

Chemicals, materials, and plasmid constructions

Cell culture media were purchased form Gibco BRL (Grand Island, NY). Calf serum was obtained from Hyclone Laboratories, Inc. (Logan, UT), and FCS was purchased from Atlanta Biologicals (Atlanta, GA). ¹⁴C-Chloramphenicol (50-60 Ci/mmol) was from DuPont, NEN Research Products (Boston, MA). ICI 182,780 was kindly provided by Dr. Alan Wakeling. Zeneca Pharmaceuticals, Macclesfield, UK. Luciferase reporter assay system was from Promega (Madison. WI). The expression vector for human ER α (pCMV5-hER α) was constructed previously as described. The expression vector pCMV5-ER β was constructed by inserting the cDNA encoding the full length human ER β (530 residues) or the short form of human ER β (477 residues), into the

*Bam*HI site of pCMV5. The estrogen responsive reporter plasmids were (ERE)₃-pS2-CAT, constructed as described previously. (ERE)₂-pS2-Luc, constructed by inserting (ERE)₂-pS2 fragment from (ERE)₂-pS2-CAT into the *Mlul/BgIII* sites of pGL3-Basic vector (Promega, Madison, WI), NHERF-Luc, constructed by inserting human NHERF promoter fragment (containing –2985 to +496 bp of human NHERF gene) into the *Kpul/Xhol* sites of pGL3-Basic vector. The plasmid pCMVβ (Clontech, Palo Alto, CA) which contains the β-galactosidase gene, was used as an internal control for transfection efficiency. *Cell culture and transient transfections*

Human endometrial cancer (HEC-1) cells were maintained in culture as described. Transfection of HEC-1 cells in 24-well plate used a mixture of 0.35 ml of serum-free IMEM media and 0.15 ml of HBSS containing 5 μ l of lipofectin (Life Technologies. Rockville. MD), 1.6 μ g of transferrin (Sigma, St. Louis, MO), 0.5 μ g of pCMV β as internal control, 1 μ g of the reporter gene plasmid, 100 ng of ER expression vector, and carrier DNA to a total of 2.5 μ g DNA, applying them to each well. The cells were incubated at 37°C in a 5% CO₂ containing incubator for 6 h. Fresh media containing desired concentrations of ligands were added to each well. Reporter gene activity was assayed after 24 h incubation. CAT and luciferase activity, normalized for the internal control β -galactosidase activity, were assayed as described. MCF-7 cells were maintained in culture as described. About $0.3 \times 10^{\circ}$ cells were seeded into each well of a 6-well plate with phenol red free media. After 3 days, fresh media containing desired concentration of ligands were added to each well. 24 h later, the cells were harvested for RNA

extraction

Real-time PCR analysis

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The total RNA from MCF-7 cells was isolated using TRIzol® reagent (Life Technologies. Rockville, MD) according to the manufacturer's suggestion. 1 μ g of total RNA was reverse transcribed using GeneAmp® RNA PCR kit (Applied Biosystem, Foster City, CA). 1/100 of each cDNA products were subjected to the real-time PCR analysis on ABI PRISM 7700 sequence detection system using TaqMan® universal PCR master mix (Applied Biosystem, Foster City, CA). The pS2 cDNA was amplified with the primers pS2f (5' GCGCCTGGTCCTGGTGTCCAT 3') and pS2r (5' GAAAACCACAATTCTGTCTTTCAC3') and was detected by the probe 6FAM-CCCAGACAGAGGTGTACAGTGGCCC-TAMRA. The GAPDH was used as the internal control which was amplified with the primers GAPDHf (5' GAAGGTGAAGGTCGGAGTC 3') and GAPDHr (5' GAAGATGGTGATGGGATTTC 3'). The signal was detected by the probe 6FAM-CAAGCTTCCCGTTTCA3'). The signal was detected by the probe the relative expression level of pS2 mRNA.

Acknowledgments

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Pyrazoles Series

			RBA	RBA			
Cpd.	No. R	X	ERα	ERβ	ERα/ERβ		
I	H-	Me- Et- Pr- [PPT] Bu-	5.4 36 51 14	0.039 0.15 0.13 0.18	140 240 390 78		
I-A		Me- [MPF Et- Pr-	?] 11 12 6.8	0.05 0.65 0.54	220 18 13		
I-B		Et-	9.5	0.08	120		
I-C		Me- Et-	1.2 16	0.05 0.17	24 94		
I-D	Me.N~~~	Et-	9.1	0.64	14		
I-E	Et.N	Me- Et-	7.8 17	0.05 0.17	156 100		
I-F		Et-	13	0.43	30		
I-G	HO	Et	7.0	0.25	28		

Ar = p-hydroxyphenyl

Figure 1. Structures and estrogen receptor alpha and beta (ER α and ER β) binding affinities of pyrazoles and basic side chain (BSC) pyrazoles. Binding affinities are determined in a radiometric competitive binding assay, using [³H]estradiol as tracer and hydroxylapatite to adsorb ligand-receptor complex. Receptor preparations were human ER α and ER β , expressed in baculovirus and purified (PanVera). Values represent the mean of 2-3 repeat determinations (CV 0.3). Note that a simple compound numbering neumonic is derived from the pyrazole series (designated with a boldface Roman numeral I), the nature of the basic side chain (designated as a boldface uppercase letter A-E, as noted in the figure), and the nature of the C(4) substituent (boldface Me, Et, Pr)

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Figure 2.



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Concentration (Log M)





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Figure 4.



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Figure 5.



Figure 6.



### DEPARTMENT OF THE ARMY

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