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## **RPPR Final Report**

as of 10-Jul-2018

Agency Code:

Proposal Number: 70640MAII INVESTIGATOR(S):

Agreement Number: W911NF-17-1-0070

Name: Frederick R Adler Email: adler@math.utah.edu Phone Number: 8015816848 Principal: Y

Organization: University of Utah Address: 75 South 2000 East, Salt Lake City, UT 841128930 Country: USA DUNS Number: 009095365 Report Date: 01-Jan-2018 Final Report for Period Beginning 02-Jan-2017 and Ending 01-Oct-2017 Title: How complex systems cope with noise: Balancing centralized and decentralized control Begin Performance Period: 02-Jan-2017 Report Term: 0-Other Submitted By: Frederick Adler Email: adler@math.utah.edu Phone: (801) 581-6848

**Distribution Statement:** 1-Approved for public release; distribution is unlimited.

### **STEM Degrees:**

### **STEM Participants:**

**Major Goals:** We investigated two complementary topics: how living things respond to stochasticity and how we as modelers do the same. In particular, we examined the strategies organism use for maintaining and enhancing function in unpredictable environments, and the strategies used to model complexity in the face of stochasticity.

**Accomplishments:** Through extensive review and synthesis of the literature, we developed the first comprehensive outline of the theme of stochastic dynamics in biological systems. Our literature review and framework provides the foundation for a review paper. We here present our detailed outline of this work and a selected bibliography of papers we synthesized.

A. Deterministic vs. stochastic processes in biological systems

i. Broad examples of mechanisms dominated by deterministic processes
ii. Broad counter-examples of mechanisms in which stochasticity plays an important role
iii. Challenges to expanding our view of the world to include stochasticity and why doing so is vitally important.
iv. Brief description of the role of math modeling in understanding biological systems.

B. Coping with stochasticity

i. Scales of stochasticity: noise vs catastrophic events
ii. Realized overcapacity: Lung tissues, genetic redundancy/degeneracy, futile cycling, large broods.
iii. Potential overcapacity:
iv. Dangers of overcapacity:

C. Capitalizing on stochasticity

i. Trial and Error (Developmental Selection): neural Development, angiogenesis, oogenesis, T cells, Rag transposons/affinity maturation.

ii. Genetic Regulation: delayed reinitiation (ATF4), gene duplication(K3L),

iii. Ploidy Changes: liver, fungi, C. neoformans

## **RPPR Final Report**

as of 10-Jul-2018

iv. Resource Acquisition: ant foraging

v. Noise in protein production: peptide fragments as an immune signal

vi. Binding Promiscuity: detoxification and immune response

D. Communication in a stochastic world

i. Stochastically established coordination: Neural growth

ii. Regulation of capacity

- iii. Recognition of environmental state
- iv. Robustness: maintaining a phenotype despite environmental/cellular stochasticity
- v. Stochastically established coordination: immune system

E. The role of mathematical modeling in understanding stochastically driven systems

i. History and classic examples

ii. Challenges in determining when and how to use stochastic modeling iii. Call for future work

**Training Opportunities:** Two graduate students were trained. Laura Strube greatly advanced her thesis work on the integrated stress response in cells, and will be completing her dissertation in October 2018 based on that research. Emerson Arehart used this opportunity to build his knowledge of complex systems and will be writing a thesis proposal based on that work to be defended in September, 2018.

Results Dissemination: Nothing to Report

Honors and Awards: Nothing to Report

**Protocol Activity Status:** 

Technology Transfer: Nothing to Report

**PARTICIPANTS:** 

Participant Type: PD/PI Participant: Frederick R Adler Person Months Worked: 6.00 Project Contribution: International Collaboration: International Travel: National Academy Member: N Other Collaborators:

Funding Support:

 Participant Type: Graduate Student (research assistant)

 Participant: Laura Strube

 Person Months Worked: 9.00
 Funding Support:

 Project Contribution:

 International Collaboration:

 International Travel:

 National Academy Member: N

 Other Collaborators:

Participant Type:Graduate Student (research assistant)Participant:Emerson ArehartPerson Months Worked:3.00Funding Support:Project Contribution:

# RPPR Final Report

as of 10-Jul-2018

International Collaboration: International Travel: National Academy Member: N Other Collaborators:

## DISSERTATIONS:

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 Authors: Laura Strube

 Acknowledged Federal Support: Y

Statement of the problem studied

We investigated two complementary topics: how living things respond to stochasticity and how we as modelers do the same. In particular, we examined the strategies organsism use for maintaining and enhancing function in unpredictable environments, and the strategies used to model complexity in the face of stochasticity.

Summary of the most important results

Through extensive review and synthesis of the literature, we developed the first comprehensive outline of the theme of stochastic dynamics in biological systems. Our literature review and framework provides the foundation for a review paper. We here present our detailed outline of this work and a selected bibliography of papers we synthesized.

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E. The role of mathematical modeling in understanding stochastically driven systems

i. History and classic examples ii. Challenges in determining when and how to use stochastic modeling iii. Call for future work Bibliography

L. W. Ancel and W. Fontana, Plasticity, evolvability, and modularity in RNA, Journal of Experimental Zoology, 288 (2000), pp. 242--283.

A. Arkin, J. Ross, and H. H. McAdams, Stochastic kinetic analysis of developmental pathway bifurcation in phage \$\lambda\$-infected Escherichia coli cells, Genetics, 149 (1998), pp. 1633--1648.

W. M. Atkins, Biological messiness vs. biological genius: mechanistic aspects and roles of protein promiscuity, The Journal of steroid biochemistry and molecular biology, 151 (2015), pp. 3--11.

K. Axelrod, A. Sanchez, and J. Gore, Phenotypic states become increasingly sensitive to perturbations near a bifurcation in a synthetic gene network, eLife, 4 (2015), p. e07935.

A. K. Bershad, M. A. Fuentes, and D. C. Krakauer, Developmental autonomy and somatic niche construction promotes robust cell fate decisions, J. Theor. Biol., 254 (2008), pp. 408--416.

B. J. Binder, K. A. Landman, D. F. Newgreen, and J. V. Ross, Incomplete penetrance: The role of stochasticity in developmental cell colonization, J. Theor. Biol., 380 (2015), pp. 309--314.

J. D. Bloom, S. Labthavikul, C. R. Otey, and F. H. Arnold, Protein stability promotes evolvability, Proc. Nat. Acad. Sci., 103 (2006), pp. 5869--5874.

L. M. Boukhibar and M. Barkoulas, The developmental genetics of biological robustness, Annals Bot., 9 (2015), p. mcv128.

C. Briat, A. Gupta, and M. Khammash, Antithetic integral feedback ensures robust perfect adaptation in noisy bimolecular networks, Cell Systems, 2 (2016), pp. 15--26.

L. Castelli, R. Pesenti, and D. Segre, The cell as a decision-making unit, IEEE Life Sciences Letters, 2 (2016), pp. 27--30.

G. Chalancon, C. N. Ravarani, S. Balaji, A. Martinez-Arias, L. Aravind, R. Jothi, and M. M. Babu, Interplay between gene expression noise and regulatory network architecture, Trends in Genetics, 28 (2012), pp. 221--232.

M. F. Ciaccio, J. D. Finkle, A. Y. Xue, and N. Bagheri, A systems approach to integrative biology: an overview of statistical methods to elucidate association and architecture, Integrative and Comparative Biology, 54 (2014) pp. 296--306

B. S. Cooper, L. A. Hammad, N. P. Fisher, J. A. Karty, and K. L. Montooth, In a variable thermal environment selection favors greater plasticity of cell membranes in drosophila melanogaster, Evolution, 66 (2012), pp. 1976--1984.

E. Dacheux, H. Firczuk, and J. E. McCarthy, Rate control in yeast protein synthesis at the population and single-cell levels, Biochemical Society Transactions, 43 (2015), pp. 1266--1270.

J. Davila-Velderrain and E. R. Alvarez-Buylla Bridging genotype and phenotype, Frontiers in Ecology, Evolution and Complexity, (2014)

H. De Jong Modeling and simulation of genetic regulatory systems: a literature review, Journal of Computational Biology, 9 (2002)

S. A. Diaz and M. Viney, Genotypic-specific variance in caenorhabditis elegans lifetime fecundity, Ecology and evolution, 4 (2014), pp. 2058--2069.

J. Draghi and M. Whitlock, Robustness to noise in gene expression evolves despite epistatic constraints in a model of gene networks, Evolution, 69 (2015), pp. 2345--2358.

M. B. Elowitz, A. J. Levine, E. D. Siggia, and P. S. Swain, Stochastic gene expression in a single cell, Science, 297 (2002), pp. 1183--1186.

M.-A. Felix and M. Barkoulas, Pervasive robustness in biological systems, Nature Reviews Genetics, 16 (2015), pp. 483--496.

H. B. Fraser, A. E. Hirsh, G. Giaever, J. Kumm, and M. B. Eisen, Noise minimization in eukaryotic gene expression, PLoS Biol, 2 (2004), p. e137.

C. Furusawa and K. Kaneko, A generic mechanism for adaptive growth rate regulation, PLoS Comp. Biol., 4 (2008), p. e3.

A. B. Gjuvsland, J. O. Vik, D. A. Beard, P. J. Hunter, and S. W. Omholt Bridging the genotype--phenotype gap: what does it take?, The Journal of Physiology, 591 (2013)

P. Godfrey-Smith and M. Martinez, Communication and common interest, PLOS Comput Biol, 9 (2013), p. e1003282.

D. M. Gordon, The Evolution of the Algorithms for Collective Behavior, Cell Systems, 3 (2016) pp 514--520

N. Hao, B. A. Budnik, J. Gunawardena, and E. K. O'Shea, Tunable signal processing through modular control of transcription factor translocation, Science, 339 (2013), pp. 460--464.

K. D. Harris, M. Weiss, and A. Zahavi, Why are neurotransmitters neurotoxic? an evolutionary perspective, F1000Research, 3 (2014).

Y. Hart and U. Alon, The utility of paradoxical components in biological circuits, Molecular cell, 49 (2013), pp. 213--221.

E. Jablonka, Do cells show off? somatic selection and the nature of intercellular signalling, Trends Ecol. Evol., 11 (1996), pp. 395--396.

Q. A. Justman, An explicit source for extrinsic noise, Cell systems, 1 (2015), pp. 308--309.

A. Kashiwagi, I. Urabe, K. Kaneko, and T. Yomo, Adaptive response of a gene network to environmental changes by fitness-induced attractor selection, PLoS One, 1 (2006), p. e49.

E. Fox Keller, Revisiting "scale-free" networks, BioEssays, 27 (2005), pp. 1060--1068.

F. D. Klironomos, J. Berg, and S. Collins, How epigenetic mutations can affect genetic evolution: model and mechanism, Bioessays, 35 (2013), pp. 571--578.

D. C. Krakauer and M. Pagel, Selection by somatic signals: the advertisement of phenotypic state through costly intercellular signals, Phil. Trans. Roy. Soc. B, 351 (1996), pp. 647--658.

E. Kussell and S. Leibler, Phenotypic diversity, population growth, and information in fluctuating environments, Science, 309 (2005), pp. 2075--2078.

B. Lehner, Conflict between noise and plasticity in yeast, PLoS Genet, 6 (2010), p. e1001185.

M. A. Lemmon, D. M. Freed, J. Schlessinger, and A. Kiyatkin, The dark side of cell signaling: Positive roles for negative regulators, Cell, 164 (2016), pp. 1172--1184.

I. Lestas, G. Vinnicombe, and J. Paulsson, Fundamental limits on the suppression of molecular fluctuations, Nature, 467 (2010), pp. 174--178.

Liu, J., Martin-Yken, H., Bigey, F., Dequin, S., François, J. M., & Capp, J. P. (2015). Natural yeast promoter variants reveal epistasis in the generation of transcriptional-mediated noise and its potential benefit in stressful conditions. Genome biology and evolution, 7(4), 969-984.

J. Liu, J.-M. Francois, and J.-P. Capp, Use of noise in gene expression as an experimental parameter to test phenotypic effects, Yeast, 33 (2016), pp. 209--216.

L. Lopez-Maury, S. Marguerat, and J. Bahler, Tuning gene expression to changing environments: from rapid responses to evolutionary adaptation, Nature Reviews Genetics, 9 (2008), pp. 583--593.

B. B. Machta, R. Chachra, M. K. Transtrum, and J. P. Sethna, Parameter space compression underlies emergent theories and predictive models, Science, 342 (2013), pp. 604--607.

T. T. Marquez-Lago and J. Stelling, Counter-intuitive stochastic behavior of simple gene circuits with negative feedback, Biophysical journal, 98 (2010), pp. 1742--1750.

M. D. McDonnell and L. M. Ward, The benefits of noise in neural systems: bridging theory and experiment., Nature Reviews Neuroscience, 12 (2011).

I. A. Mellis and A. Raj, Half dozen of one, six billion of the other: What can small-and large-scale molecular systems biology learn from one another?, Genome Research, 25 (2015), pp. 1466--1472.

H. M. Meyer and A. H. Roeder, Stochasticity in plant cellular growth and patterning, Frontiers in plant science, 5 (2014).

K. J. Mitchell, What is complex about complex disorders, Genome Biol, 13 (2012), p. 237.

A. P. Moczek, S. Sultan, S. Foster, C. Ledon-Rettig, I. Dworkin, H. F.

Nijhout, E. Abouheif, and D. W. Pfennig, The role of developmental plasticity in evolutionary innovation, Proc. Roy. Soc. B, 278 (2011), pp. 2705--2713.

F. Moss, L. M. Ward, and W. G. Sannita, Stochastic resonance and sensory information processing: a tutorial and review of application, Clinical neurophysiology, 115 (2004), pp. 267--281.

D. Muzzey, C. A. Gomez-Uribe, J. T. Mettetal, and A. van Oudenaarden, A systems-level analysis of perfect adaptation in yeast osmoregulation, Cell, 138 (2009), pp. 160--171.

D. F. Newgreen, S. Dufour, M. J. Howard, and K. A. Landman Simple rules for a "simple" nervous system? molecular and biomathematical approaches to enteric nervous system formation and malformation, Developmental biology, 382 (2013)

J. R. S. Newman, S. Ghaemmaghami, J. Ihmels, D. K. Breslow, M. Noble, J. L. DeRisi, and J. S. Weissman, Single-cell proteomic analysis of S. cerevisiae reveals the architecture of biological noise, Nature, 441 (2006), pp. 840--846.

H. F. Nijhout, Metaphors and the role of genes in development BioEssays, 12, 441--446, 1990,

D. Noble, A biological relativity view of the relationships between genomes and phenotypes, Progress in Biophysics and Molecular Biology, 111 (2013), pp. 59--65.

S. W. Omholt From sequence to consequence and back, Progress in Biophysics and Molecular Biology, 111 (2013)

E. M. Ozbudak, M. Thattai, I. Kurtser, A. D. Grossman, and A. Van Oudenaarden, Regulation of noise in the expression of a single gene, Nature genetics, 31 (2002), pp. 69--73.

V. Pancaldi, Biological noise to get a sense of direction: an analogy between chemotaxis and stress response, Frontiers in genetics, 5 (2014).

S. Patalano, A. Vlasova, C. Wyatt, P. Ewels, F. Camara, P. G. Ferreira, C. L. Asher, T. P. Jurkowski, A. Segonds-Pichon, M. Bachman, et al., Molecular signatures of plastic phenotypes in two eusocial insect species with simple societies, Proc. Nat. Acad. Sci., 112 (2015), pp. 13970--13975.

R. Pinho, V. Garcia, and M. W. Feldman, Phenotype accessibility and noise in random threshold gene regulatory networks, PloS one, 10 (2015), p. e0119972.

D. M. Posadas and R. W. Carthew, Micrornas and their roles in developmental canalization, Current opinion in genetics & development, 27 (2014), pp. 1--6.

J. M. Raser and E. K. O'Shea, Control of stochasticity in eukaryotic gene expression, Science, 304 (2004), pp. 1811--1814.

M. Richard and G. Yvert, How does evolution tune biological noise?, Frontiers in Genetics, 5 (2014), p. 374.

M. A. Riley and M. T. Turvey, Variability and determinism in motor behavior, Journal of motor behavior, 34 (2002), pp. 99--125.

A. Sanchez and J. Kondev, Transcriptional control of noise in gene expression, Proc. Nat. Acad. Sci., 105 (2008), pp. 5081--5086.

V. Shahrezaei and S. Marguerat, Connecting growth with gene expression: of noise and numbers, Current opinion in microbiology, 25 (2015), pp. 127--135.

M. S. Sherman, K. Lorenz, M. H. Lanier, and B. A. Cohen, Cell-to-cell variability in the propensity to transcribe explains correlated fluctuations in gene expression, Cell Systems, 1 (2015), pp. 315--325.

V. Siciliano, I. Garzilli, C. Fracassi, S. Criscuolo, S. Ventre, and D. di Bernardo, MiRNAs confer phenotypic robustness to gene networks by suppressing biological noise, Nature Communications, 4 (2013).

O. K. Silander, N. Nikolic, A. Zaslaver, A. Bren, I. Kikoin, U. Alon, and M. Ackermann, A genome-wide analysis of promoter-mediated phenotypic noise in Escherichia coli, PLoS genetics, 8 (2012), p. e1002443.

G. P. Singh, Coupling between noise and plasticity in e. coli, G3: Genes, Genomes, Genetics, 3 (2013), pp. 2115--2120.

E. C. Snell-Rood, Selective processes in development: Implications for the costs and benefits of phenotypic plasticity., Integrative & Comparative Biology, 52 (2012), pp. 31--42.

C. Sokolik, Y. Liu, D. Bauer, J. McPherson, M. Broeker, G. Heimberg, L. S. Qi, D. A. Sivak, and M. Thomson, Transcription factor competition allows embryonic stem cells to distinguish authentic signals from noise, Cell systems, 1 (2015), pp. 117--129.

S. R. Starck and N. Shastri, Nowhere to hide: unconventional translation yields cryptic peptides for immune surveillance, Immunological reviews, 272 (2016), pp. 8--16.

D. J. Stekel and D. J. Jenkins, Strong negative self regulation of prokaryotic transcription factors increases the intrinsic noise of protein expression, BMC systems biology, 2 (2008), p. 6.

N. Stergiou and L. M. Decker, Human movement variability, nonlinear dynamics, and pathology: is there a connection?, Human movement science, 30 (2011), pp. 869--888.

T. Szekely and K. Burrage, Stochastic simulation in systems biology, Computational and structural biotechnology journal, 12 (2014), pp. 14--25.

M. Thattai and A. Van Oudenaarden, Intrinsic noise in gene regulatory networks, Proc. Nat. Acad. Sci., 98 (2001), pp. 8614--8619.

C. J. Tomlin and J. D. Axelrod Biology by numbers: mathematical modelling in developmental biology Nature Reviews Genetics, 8 (2007)

L. S. Tsimring, Noise in biology, Reports on Progress in Physics, 77 (2014), p. 026601.

J. J. Tyson and B. Novak, Control of cell growth, division and death: information processing in living cells, Interface Focus, 4 (2014), p. 20130070.

M. Varlet, R. C. Schmidt, and M. J. Richardson, Influence of internal and external noise on spontaneous visuomotor synchronization, Journal of motor behavior, 48 (2016), pp. 122--131.

M. Viney and S. E. Reece, Adaptive noise, Proc. Roy. Soc. B, 280 (2013), p. 20131104.

G. Vogt, Stochastic developmental variation, an epigenetic source of phenotypic diversity with far-reaching biological consequences, Journal of biosciences, 40 (2015), pp. 159--204.

G. P. Wagner and J. Zhang, The pleiotropic structure of the genotype--phenotype map: the evolvability of complex organisms, Nature Reviews Genetics, 12 (2011), pp. 204--213.

Q. Wang, W. R. Holmes, J. Sosnik, T. Schilling, and Q. Nie, Cell sorting and noise-induced cell plasticity coordinate to sharpen boundaries between gene expression domains, PLoS Comp. Biol., 13 (2017), p. e1005307.

Z. Wang and J. Zhang, Impact of gene expression noise on organismal fitness and the efficacy of natural selection, Proc. Nat. Acad. Sci., 108 (2011), pp. E67--E76.