Award Number: W81XWH-09-2-0109

TITLE: Innovative Computational Waterborne Pathogen Research for Chemical/Biological Detection

PRINCIPAL INVESTIGATOR: David M. Mosser, Ph.D.

CONTRACTING ORGANIZATION: University of Maryland, College Park, MD 20742

REPORT DATE: September 2010

TYPE OF REPORT: Annual

PREPARED FOR: U.S. Army Medical Research and Materiel Command Fort Detrick, Maryland 21702-5012

DISTRIBUTION STATEMENT:

Approved for public release; distribution unlimited

The views, opinions and/or findings contained in this report are those of the author(s) and should not be construed as an official Department of the Army position, policy or decision unless so designated by other documentation.

-					Form Approved
		UMENTATIO	-		OMB No. 0704-0188
data needed, and completing a this burden to Department of D 4302. Respondents should be	and reviewing this collection of in befense, Washington Headquart aware that notwithstanding any	nformation. Send comments regarders Services, Directorate for Info	arding this burden estimate or an rmation Operations and Reports n shall be subject to any penalty t	y other aspect of this co (0704-0188), 1215 Jeff	ching existing data sources, gathering and maintaining the ollection of information, including suggestions for reducing erson Davis Highway, Suite 1204, Arlington, VA 22202- h a collection of information if it does not display a currently
1. REPORT DATE (DD 01-09-2010	D-MM-YYYY)	2. REPORT TYPE Annual	xess.		DATES COVERED (From - To) 5 August 2009 - 14 August 2010
4. TITLE AND SUBTIT		AIIIuai			CONTRACT NUMBER
				5b.	GRANT NUMBER
Innovative Con Chemical/Biolo	-	-	en Research for	-	81XWH-09-2-0109 PROGRAM ELEMENT NUMBER
Chemical/Bioit	gical Decection	511			
6. AUTHOR(S) David Mosser, Ph.D				5d.	PROJECT NUMBER
dmosser@umd.e	edu			5e.	TASK NUMBER
				5f.	WORK UNIT NUMBER
7. PERFORMING ORG	GANIZATION NAME(S)	AND ADDRESS(ES)		-	PERFORMING ORGANIZATION REPORT
University of	_				
College Park,	MD 20742				
9. SPONSORING / MC	NITORING AGENCY N	IAME(S) AND ADDRES	S(ES)	10.	SPONSOR/MONITOR'S ACRONYM(S)
		Ind Materiel Com			
Fort Detrick, M	D 21702-5012			11	SPONSOR/MONITOR'S REPORT
					NUMBER(S)
12. DISTRIBUTION / A	VAILABILITY STATEM	IENT		• • • • • • • • • • • • • • • • • • •	
Approved for pu	ıblic release; dis	stribution unlimite	ed		
13. SUPPLEMENTAR	Y NOTES				
14. ABSTRACT					
					ntilevers that will be used ated devices will allow
water samples	to be surveyed	d and monitored	in real time.	They will	be highly sensitive and
_	-		-		at we are developing is
					g software that we have Aquisite sensitivity to
-	-		2		the detection of small
					bined expertise of life
	ngineers, and o versee this pro	_	iologists, MPRI	I is unique	ely positioned to develop,
15. SUBJECT TERMS					
16. SECURITY CLASS	SIFICATION OF:		17. LIMITATION OF ABSTRACT	18. NUMBER OF PAGES	19a. NAME OF RESPONSIBLE PERSON USAMRMC
a. REPORT	b. ABSTRACT	c. THIS PAGE	UU	10	19b. TELEPHONE NUMBER (include area
U	U	U			code)
				1	Standard Form 298 (Pov. 8-98)

Table of Contents

Page

ntroduction1	
3ody1	
Key Research Accomplishments5	
Reportable Outcomes 5	
Conclusion5	
References6	
Appendices	

Annual Report:

Introduction:

We are constructing a prototype device to detect waterborne pathogens in real time, with a high degree of sensitivity, and low false positives. In the next year, we will continue refining this device with the goal of β -testing it against a limited number of known bacterial species. This will allow us to determine the feasibility of this approach, the sensitivity of this instrumentation, and the specificity of detection. The milestones that have been completed thus far include: –DNA probe chemically immobilized on microchip surface.

- -Target sequences in sample hybridize at surface with probes.
- -Enzyme-labeled sandwich probe hybridizes to target sequence.

-Sensitivity has been improved so that as little as 0.5 nM of target DNA sequences can be recognized in microchannels with interdigitated microelectrodes

Body:

The goal of this project is to construct an on-chip DNA biosensor that will be used to detect pathogens in contaminated water. The on-chip format enables integration with other functions, such as sample processing, DNA extraction and PCR. The unique design will have several biosensors fabricated in parallel so that it will be possible to detect multiple signatures from a single species, as well as multiple species/strains with a single device (Figure 1). The novel aspects of the pathogen detection device that we propose to develop are that i) they have a high degree of pathogen specificity, ii) they are highly sensitive, iii) they can detect pathogens in real time, iv) and they have virtually no false positives.

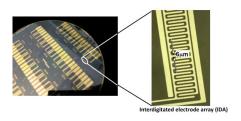


Figure 1. Schematic of biosensor device designed to detect multiple pathogens at the same time.

Signature selection. For the initial proof of concept studies, the bacterium *P. aeruginosa* is being used for detection, because it is a well-understood model organism with both pathogenic and non-pathogenic subtypes. Using the Insignia pipeline (Figure 2) we have selected signature sequences for our initial testing. We have generated signatures that distinguish pathogenic from nonpathogenic strains. We have also designed and had synthesized targets with one or two mismatches and the probes that are complementary to those targets (Figure 3). Experiments using these oligos will further test the specificity of the probes binding to the targets.

We have also made significant updates in bioinformatics, specifically to the Insignia Pipeline database. In the past year, Dr. Steven Salzberg, one of the main contributors on this project and the developer of Insignia, has increased the number of pathogens and their related signatures in the database from 8,341 to 13,928. The implications of this significant increase are that we will be able to adapt the biosensor to recognize an extensive range of pathogens.

Navigation » Home	Has been increased to 13928 organisms
» Run Insignia	Insignia Pipeline 11274 viruses/phages, 2653 non-viruses)
» Validated Signatures	There are 8341 organisms in Insignia (5506 viruses/phages , 1139 non-viruses)
» Examples	Click to select, ctrl+click to unselect or multi-select, shift+click to range-select
» Help	* The numbers in parentheses indicate the number of assembly contigs for each organism.
» Contact Us	—
 » Gemina Database Stats » Download 	🗐 Show Viruses/Phages 🗹 Show Nos-Viruses/Phages
	Exclude ALL organisms with greater than (2) contigs
Collaborators » ICS	
» IGS » Canon Life Sciences	- REFERENCE -
	Select one genome to act as a reference coordinate system.
RelatedLinks	narrow your search
>> GEMINA	Pseudomonas aeruginosa PA7 (1) Pseudomonas aeruginosa PACS2 (1) Pseudomonas aeruginosa PACS2 (1)
» NCBI Microbial	Pseudomonas aeruginosa PAO1 (1) Would you like to include them in the result set?
	Pseudomonas aertuophila L48 (1) Include 1-unique 18-mers
	Pseudomonas fluorescens Pf-5 (1) Include 1-unique 19-mers
	Pseudomonas mendocina ymp (1) Pseudomonas putida F1 (1)
	Pseudomonas putida CB-1 (1)
	Pseudomonas putida KT2440 (1)
	- TARGET -
	Optionally select additional genomes to target. All signatures returned for the reference genome will be shared by these genomes.
	Show all Show only organism(s) on the same taxonomy branch
	Ê-Pseudomonadales
	⊕ Moraxellaceae
	⊖Pseudomonadaceae
	Cellvibrio ⊖Pseudomonas
	Preudomonas Develomonas aeruginosa group
	· · · · · · · · · · · · · · · · · ·

Figure 3. Signatures designed for specificity testing in prototype waterborne pathogen detection device.

Wildtype: ^{5'}GTTGCCCTGGACATTGATCTGGATGTTGTTGCTTTCCATCG^{3'}

Target 1: 5'GTTGCCCTGGACATTGATCTGGATGTTGTTGGTTTCCATCG^{3'}

Target 2: ^{5'}GTTGCCCTG<mark>C</mark>ACATTGATCTGGATGTTGTTGCTTTCCATCG^{3'}

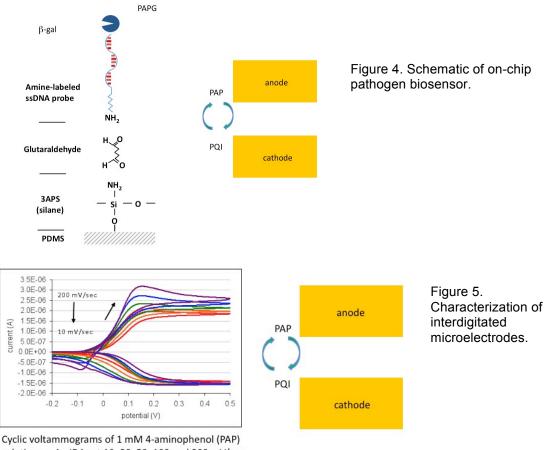
Target 3: 5'GTTGCCCTGCACATTGATCTGGATGTTGTTGGGTTTCCATCG^{3'}

Probe A: 5'-NH2 CGATGGAAAGCAACAACATC3'

Probe A variant: ^{5'-NH2}CGATGGAAA<mark>C</mark>CAACAACATC^{3'}

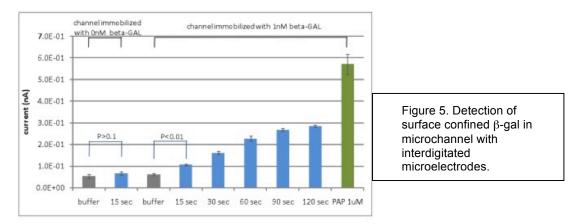
Probe B: ^{5'}AGATCAATGTCCAGGGCAAC^{3'-Thiol}

Probe B variant: 5'AGATCAATGTGCAGGGCAAC^{3'-Thiol} In the first year, we have developed and then continued refining the pathogen biosensor. Figure 4 depicts the binding of the oligonucleotide probe to the biosensor and the signal detection methodology. This includes that the DNA probe chemically immobilized on microchip surface and then target sequences in each sample hybridize at surface with probes. An enzyme-labeled sandwich probe hybridizes to target sequence. Key to the detection is that the substrate (PAPG) is cleaved by enzyme (β -gal) and forms PAP. Since PAP and PQI are reversible states of redox active molecules, each redox event (PAP -> PQI, PQI -> PAP) results in a current signal thereby amplifying the signal (Figure 5).



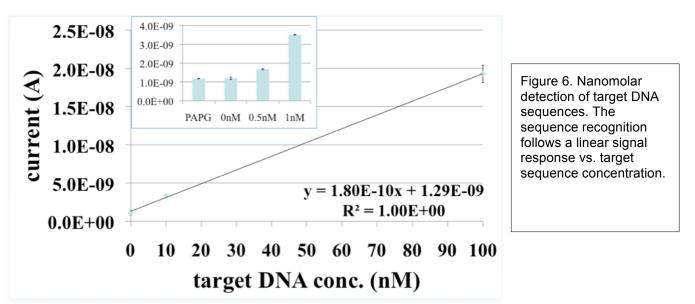
Cyclic voltammograms of 1 mM 4-aminophenol (PAP) solution on Au IDAs at 10, 20, 50, 100 and 200 mV/sec in 35- μ L sample droplets. Collecting electrode (cathode) was kept at -0.15 V.

Initial experiments in Q1 that 100 nM of target DNA can bind to specific labeled probe and elicit biosensor current above background. We have recently demonstrated that as little as 0.5 nM of target DNA sequences can be recognized in microchannels with interdigitated microelectrodes (Figure 6). This represents a 200-fold improvement since this project began in August of 2009. We are continuing to refine the detection system to increase the sensitivity.



Oxidation current recorded on IDA in microchannel immobilized with 0 and $1nM \beta$ -GAL.

Next steps will include empirically determining optimal signature length. Several signatures that are unique to a selected pathogen and of the appropriate length will be used to develop a panel of pathogen-specific oligonucleotide probes. We aim to perform the entire biosensing experiment on electrode microchip and determine detection limits for the specific DNA sequence. We also plan to test strategies for optimizing selective hybridization.



Problems:

1) Sample preparation: Samples may need to be concentrated in order to maximize exposure of the devices to large quantities of water.

Dr. Ian White in the Bioengineering Department is currently working on methods to filter and concentrate samples efficiently so that pathogens that may be contaminating the samples can effectively be detected.

2) Vibration. Handheld devices and even stationary devices are subject to mechanical vibration and physical movement that can significantly impact the system's performance and reliability.

Drs. Bentley and Payne and others in the Bioengineering Department are currently working on this issue, and will make every effort to design the proposed detection devices with maximum reliability and stability.

Key Research Accomplishments:

- Developed DNA signatures for *P. aeruginosa*
- Developed first generation of microchannel biosensor with interdigitated microelectrode
- Made significant upgrades to Insignia Pipeline

Reportable outcomes:

Upgraded Insignia Pipeline (open source searchable database, http://insignia.cbcb.umd.edu/)

Conclusion:

We have made progess in developing a biosensor that can detect pathogens in contaminated water. This biosensor will be extremely sensitive, inexpensive to produce and have negligible false positive or negative results. We are in the process of further improving the detection limits and determining the specificity. In the next year, we will continue refining the device for β -testing.

References

- 1. Comprehensive DNA Signature Discovery and Validation. Phillippy AM, Mason JA, Ayanbule K, Sommer DD, Taviani E, Huq A, Colwell RR, Knight IT, Salzberg SL. PLoS Comput Biol. 2007 May 18;3 (5):e98.
- Mechano-transduction of DNA hybridization and dopamine oxidation through electrodeposited chitosan network. S. T. Koev, M. A. Powers, H. Yi, L. Q. Wu, W. E. Bentley, G. W. Rubloff, G. F. Payne and R. Ghodssi. Lab Chip, 2007, 7, 103–111.
- Chitosan Biotinylation and Electrodeposition for Selective Protein Assembly. X.-W. Shi, Y. Liu, A. T. Lewandowski, L.-Q. Wu, H.-C. Wu, R. Ghodssi, G.W. Rubloff, W. E. Bentley, G. F. Payne. 2008. Lab Chip. 2008. 8 (3):420-30.
- 4. Lewandowski, H. Yi, G. F. Payne, R. Ghodssi, W. E. Bentley, G. W. Rubloff. 2007. Biotechnol Bioeng. Jul 11. Protein assembly onto patterned microfabricated devices through enzymatic activation of fusion pro-tag.
- A. T. Lewandowski, H. Yi, X. Luo, G. F. Payne, R. Ghodssi, G. W. Rubloff, William E Bentley. Lab Chip. 2007. Jan ;7 (1):103-11. Zhang, X., Edwards, J.P., and Mosser, D.M. 2006. Dynamic and transient remodeling of the macrophage IL-10 promoter during transcription. J. Immunology 177:1282-88.
- Gemina: A Web-Based Epidemiology and Genomic Metadata System Designed to Identify Infectious Agents. Schriml, L., A. Gussman, K. Phillippy, S. Angiuoli, K. Hari, A. Goates, R. Jain, T. Davidsen, A. Ganapathy, E. Ghedin, S. Salzberg, O. White, N. Hall. Lecture Notes in Computer Science, Publisher Springer Berlin /Heidelberg. 2007 4506: 228-229.
- 7. Cao, S. Zhang, X. Edwards, J.P. and Mosser, D.M. 2006. NF-κB1 homodimers

differentially regulate pro- and anti-inflammatory cytokines in macrophages. J. Biological Chemistry 281:26041-50.

- 8. CcpA-mediated repression of streptolysin S expression and virulence in the group A streptococcus. Kinkel TL, McIver KS. 2008. Infect Immun.76(8):3451-63.
- D. Taylor, J. Ladd, Q. Yu, S. Chen, J. Homola and S. Jiang, Quantitative and simultaneous detection of four foodborne bacterial pathogens with a multi-channel SPR sensor, Biosensors and Bioelectronics 22, 752–758, 2006.
- 10. M. Piliarik, L. Párová and J. Homola, High-throughput SPR sensor for food safety, Biosensors and Bioelectronics 24, 1399–1404, 2009.
- A. Ymeti, J. Greve, P. V. Lambeck, T. Wink, S. W. F. M. v. Hovell, T. A. M. Beumer, R. R. Wijn, R. G. Heideman, V. Subramaniam and J. S. Kanger, Fast, Ultrasensitive Virus Detection Using a Young Interferometer Sensor, Nano Letters 7, 394-397, 2007.

	Start Date C	Completed	Remaining	
Develop detection device and sensors	11/1/09	305	240	
Identify pathogen signatures and generate oligonucleotid es	8/15/09	180	0	
Optimize linking oligos to cantilevers	3/15/10	180	185	Under out the first and to the out of sole of sole
Validate pathogen signatures	1/1/11	0	365	Develop detection device and sensors
Culture bacterial				Identify pathogen signatures and generate oligonucleotides
strains and prepare DNA				Optimize linking oligos to cantilevers
for detection testing	3/1/11	0	180	Validate pathogen signatures
Test detecition devices and				Culture bacterial strains and prepare DNA for detection testing
modify as needed	8/15/11	0	365	Test detection devices and modify as needed
Quantitate sensitivity				Quantitate sensitivity and specificity of devices
and specificity of devices	8/15/11	0	365	Develop and optimize sample preparation tenchniques
Develop and optimize sample preparation				
tenchniques	1/1/12	0	365	

Research and Development Timeline:

Page 7 of 7