

Immobilized CRISPR Chemistry for Massively Multiplexed Target Detection

Elaine Bradford¹, Ryan Howard², Alyssa Jespersen¹, Julie L. Lucas¹, Kaylee Mathiason², Isaac Moran³, Logan Rubio³, Chelsey Smith¹, Michael Turo³, Richard A. Winegar¹

¹MRIGlobal, Gaithersburg, Maryland

²MRIGlobal, Kansas City, Missouri

³The Charles Stark Draper Laboratory, Cambridge, Massachusetts



Abstract

Current biosurveillance and diagnostic systems aim to detect a broad range of biothreats and pathogens of interest in a timely and user-friendly manner. Traditional PCR-based detection devices can be used to detect relatively small panels of assays. Typically, it takes weeks to months to develop and transition new assays onto closed platform devices. However, as Ebola virus, SARS-CoV-2 and Monkeypox virus outbreaks have shown, novel, emerging, and re-emerging pathogens present an ever evolving challenge. We are developing a Massively Multiplexed Detection (MMD) device to detect >500 pathogenic targets in a microfluidic platform. By using CRISPR technologies, we aim to be able to quickly reconfigure the open platform to accommodate new pathogens and/or new sequence variations within 24 hours of discovery.

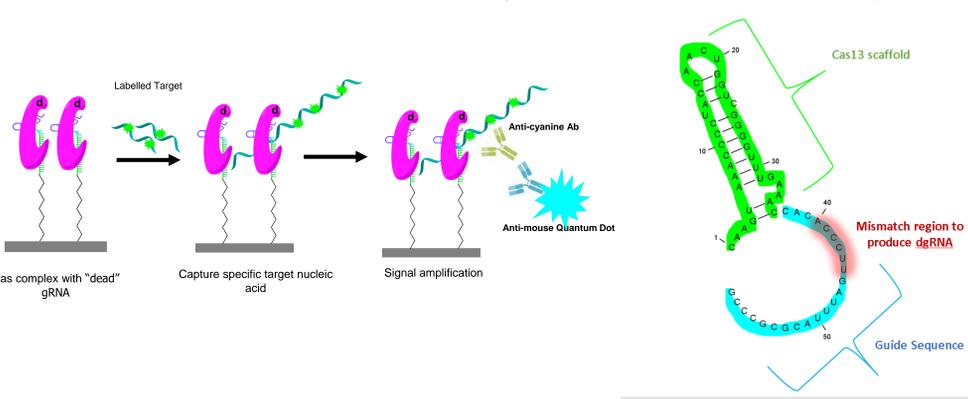
The current panel for the MMD device includes upper and lower respiratory bacteria and viruses, bacterial anti-microbial resistance genes, fungal pathogens, select agents and their near neighbors, vector-borne viruses, food-borne and water-borne pathogens, zoonotic pathogens, biomarkers for disease severity, and endogenous controls. In addition, several variant assays are included, since alterations in CRISPR guide RNA can distinguish single-nucleotide polymorphisms (SNPs) with specificity. The MMD platform requires advances in automated sample preparation, microfluidics, optics, and immobilized CRISPR-based detection chemistry. In this work we will describe strategies and progress on developing optimized CRISPR-based detection chemistry.

Overview of Chemistry

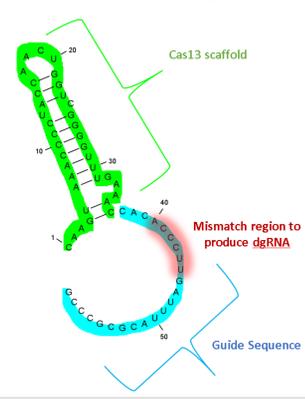
This strategy relies on the use of “dead gRNA” (d-gRNA) sequences, which are modified to include mismatches at defined sites within the guide region.

- gRNA is altered so that the Cas complex captures target specifically (and stays bound), but does not cleave the target.
- Captured nucleic acid is detected directly by pre-labelling with dye.
- Cyanine label is amplified through quantum dot-labeled anti-cyanine antibodies

Schematic of Immobilized Assay



Cas13 “dead” gRNA



Disclaimer: The views, opinions and/or findings expressed are those of the authors and should not be interpreted as representing the official views or policies of the Department of Defense or the U.S. Government.

Distribution Statement "A": Approved for Public Release, Distribution Unlimited

500-plex Panel

Targets for the MMD 500-plex panel are shown below. The number in each box is the estimated number of spots on the array for each target.

Target	Target	Target	Target	Target	Target
<i>Bordetella bronchoseptica</i> 1	D	Influenza A-H11 1	D	Influenza A-H8 1	D
<i>Bordetella pertussis</i> 1	D	Influenza A-H10 1	D	Influenza A-H9 1	D
<i>Bordetella pertussis & Bordetella parapertussis</i> 1	D	Influenza A-H11 1	D	Influenza A 1	D
<i>Chlamydia pneumoniae</i> 1	D	Influenza A-H12 1	D	Influenza B 1	D
Exogenous control MS2 1	C	Influenza A-H13 1	D	Influenza A-N1 1	D
hCoV-229E 1	D	Influenza A-H14 1	D	Influenza A-N2 1	D
hCoV-HKU1 1	D	Influenza A-H15 1	D	Influenza A-N3 1	D
hCoV-NL63 1	D	Influenza A-H16 1	D	Influenza A-N4 1	D
hCoV-OC43 1	D	Influenza A-H2 1	D	Influenza A-N5 1	D
Human adenovirus Type A 1	D	Influenza A-H3 1	D	Influenza A-N6 1	D
Human adenovirus Type B 1	D	Influenza A-H4 1	D	Influenza A-N7 1	D
Human adenovirus Type C 1	D	Influenza A-H5 1	D	Influenza A-N8 1	D
Human adenovirus Type D 1	D	Influenza A-H6 1	D	Influenza A-N9 1	D
Human adenovirus Type E 1	D	Influenza A-H7 1	D	Human parainfluenza 4 1	D
Human adenovirus Type F 1	D	Argentinian mammarenavirus 1	D	Human metapneumovirus 1	D
Human adenovirus Type G 1	D	AlphaCoV genus 1	D	Enterobacter aerogenes 1	D
<i>Acinetobacter baumannii</i> 1	D	Lassa mammarenavirus 1	D	BetaCoV genus 1	D
<i>Acinetobacter calcoaceticus-baumannii</i> complex 1	D	Machupo mammarenavirus 1	D	DeltaCoV genus 1	D
<i>Acinetobacter</i> spp. 1	D	Francisella tularensis - pan 1	D	GammaCoV genus 1	D
<i>Actinomyces</i> spp. 1	D	Andes Orthohantavirus 3	D	Enterobacter cloacae subsp. cloacae 1	D
<i>Aspergillus flavus</i> 1	D	Dobrava-Begrade orthohantavirus 1	D	Enterobacter cloacae subsp. hormaechei 1	D
<i>Aspergillus fumigatus</i> 1	D	Puumala orthohantavirus 3	D	Enterobacter cloacae subsp. kobei 1	D
<i>Aspergillus</i> spp. 3	D	Seoul orthohantavirus 3	D	Enterobacter cloacae complex 2	D
<i>Bacillus anthracis</i> 4	D	Sin Nombre orthohantavirus 3	D	<i>Escherichia coli</i> - pan 2	D
<i>Bacillus atrophaeus</i> 1	D	Hantaan orthohantavirus 3	D	<i>Escherichia coli</i> O157:H1 2	D
<i>Bacillus thuringiensis</i> 2	D	Influenza A oseltamivir resistance 2	D	<i>Pneumocystis jirovecii</i> 1	D
<i>Brucella</i> spp. 1	D	Influenza A/H1-2009 1	D	Chikungunya virus 1	D
<i>Burkholderia cepacia</i> 2	D	Influenza B-Victoria 1	D	Rabies virus 1	D
<i>Burkholderia mallei</i> 1	D	Influenza B-Yamagata 1	D	African Swine Fever Virus 1	D
<i>Burkholderia pseudomallei</i> 1	D	Human Rhinovirus/Enterovirus 6	D	Yellow Fever Virus 1	D
<i>Campylobacteraceae</i> spp. 3	D	Extended spectrum beta-lactamase (CTX-M, TEM) 7	D	Tick-borne encephalitis virus 1	D
<i>Candida albicans</i> 1	D	Human Endogenous controls 35	C	<i>Listeria monocytogenes</i> 1	D
<i>Candida lusitanae</i> 1	D	<i>Klebsiella pneumoniae</i> complex 4	D	Lumpy Skin Disease Virus 1	D
<i>Candida</i> spp. 4	D	<i>Chronobacter</i> spp. 1	D	Sheep pox virus 1	D
Carbapenemase (KPC, NDM, Oxa-23, Oxa 24/40, Oxa-48, Oxa48-like, Oxa-58, VIM, IMP) 18	D	Porcine Reproductive and Respiratory Syndrome (PRRS) - Eastern Europe 1	D	Nipah virus 1	D
<i>Citrobacter freundii</i> 1	D	Ravn Marburgvirus 1	D	Monkeypox virus 1	D
<i>Clostridium botulinum</i> 1	D	Marburg Marburgvirus 2	D	Omsk hemorrhagic fever virus 1	D
<i>Clostridium botulinum</i> toxin A 5	D	Measles morbillivirus 3	D	Foot and Mouth Disease virus 3	D
<i>Clostridium botulinum</i> toxin B 1	D	Methicillin resistance (mecA, mecC) 6	D	Porcine Reproductive and Respiratory Syndrome (PRRS) - North America 1	D
<i>Clostridium botulinum</i> toxin C/D 1	D	<i>Moraxella catarrhalis</i> 1	D	Host Biomarker - ARL17B 1	D
<i>Clostridium botulinum</i> toxin E 1	D	<i>Morganella morganii</i> 1	D	Host Biomarker - CCL20 1	D
<i>Clostridium botulinum</i> toxin F 1	D	Mumps virus 1	D	Host Biomarker - CLECSA 1	D
<i>Clostridium botulinum</i> toxin G 1	D	Murray Valley Virus 1	D	Host Biomarker - CNTN1 1	D
<i>Clostridium difficile</i> 1	D	<i>Mycobacterium bovis</i> 1	D	Host Biomarker - CREB3L3 1	D
<i>Clostridium perfringens</i> 1	D	<i>Mycobacterium tuberculosis</i> 2	D	Host Biomarker - IGF1 1	D
<i>Clostridium tetani</i> 1	D	TB - fluoroquinolones resistance 11	V	Host Biomarker - IL1R2 1	D
<i>Corynebacterium diphtheriae</i> 1	D	TB - isoniazid resistance 6	V	Host Biomarker - ILK5 1	D
<i>Corynebacterium striatum</i> 1	D	TB - rifampicin resistance 6	V	Host Biomarker - MMP19 1	D
<i>Corynebacterium ulcerans</i> 1	D	TB - SLID resistance 6	V	Host Biomarker - MUC3A 1	D
<i>Corynebacterium pseudotuberculosis</i> 2	D	<i>Mycoplasma capricolum capripneumoniae</i> 1	D	Host Biomarker - PLAUI 1	D
<i>Corynebacterium</i> spp. 3	D	<i>Mycoplasma mycoides mycoides</i> 1	D	Host Biomarker - TNFRSF12A 1	D
<i>Coxiella burnetii</i> 2	D	<i>Neisseria meningitidis</i> 1	D	<i>Rickettsia</i> spp. 2	D
<i>Cryptococcus</i> spp. 5	D	Newcastle disease virus 1	D	<i>Salmonella enterica</i> Choleraesuis 2	D
Cytomegalovirus 1	D	<i>Nocardia</i> spp. 1	D	<i>Salmonella enterica</i> Paratyphi 2	D
Ebola virus - Bombali 1	D	Orthopox virus 2	D	<i>Salmonella enterica</i> Enteritidis 4	D
Ebola virus - Bundibugyo 1	D	Peste des petits ruminants virus 2	D	<i>Salmonella enterica</i> Typhimurium 5	D
Ebola virus - Reston 1	D	Poliovirus 1	D	<i>Salmonella enterica</i> Typhimurium 5	D
Ebola virus - Sudan 1	D	<i>Proteus</i> spp. 4	D	<i>Staphylococcus aureus</i> 2	D
Ebola virus - Tai Forest 1	D	<i>Pseudomonas aeruginosa</i> 1	D	<i>Stenotrophomonas maltophilia</i> 2	D
Ebola virus - Zaire 1	D	<i>Pseudomonas</i> Virus 1	D	<i>Streptococcus agalactiae</i> 2	D
				<i>Streptococcus pneumoniae</i> 2	D
				<i>Streptococcus pyogenes</i> 1	D
				<i>Streptococcus</i> spp. 4	D
				Togaviridae EEEV 1	D
				Togaviridae VEEV 3	D
				Togaviridae WEEV 1	D
				Varicella Zoster virus 1	D
				<i>Yersinia enterocolitica</i> 2	D
				<i>Yersinia pseudotuberculosis</i> 2	D
				<i>Yersinia pestis</i> 6	D

Target Type	RNA	DNA	Both	Assay Type	D	V	C
					Detection	Variant	Control

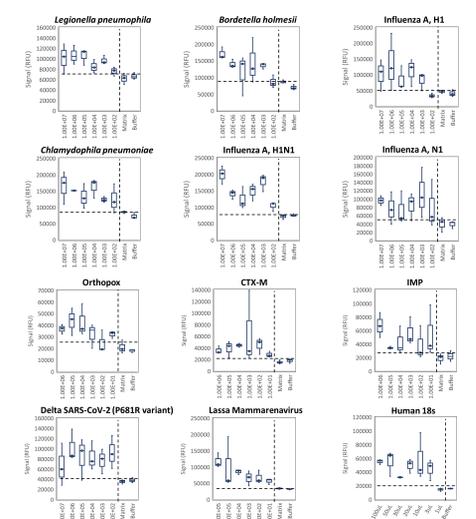
Results

Results for a subset of 12 assays are shown. Testing was done with a background of artificial nasal matrix. Target material was quantified by droplet digital PCR prior to testing. All plate-based assay experiments were in triplicate.

Assay Sensitivity

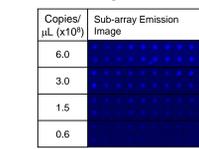
Assay	Target Type	Limit of Detection
<i>Bordetella holmesii</i>	Bacteria	1000 copies
<i>Legionella pneumophila</i>	Bacteria	1000 copies
<i>Chlamydia pneumoniae</i>	Bacteria	100 copies
Human 18S	Host internal control	3mL of nasal matrix
Orthopox	DNA virus	10,000 copies
FluA	RNA virus	100 copies
Influenza A-H1	RNA virus	1000 copies
Influenza A-N1	RNA virus	1000 copies
Lassa mammarenavirus	RNA virus	10 copies
IMP	AMR marker	1000 copies
CTX-M-2	AMR marker	10 copies
P681R SARS-CoV-2 Delta	Variant assay	100 copies

Dynamic Range

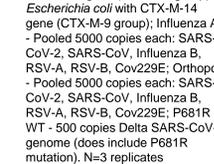
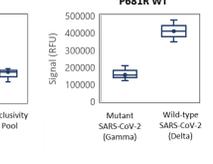
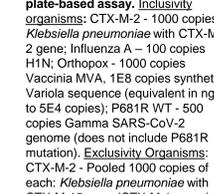
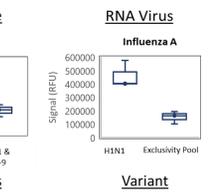
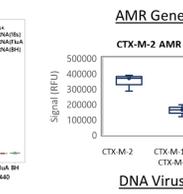
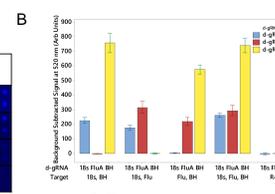


Inclusivity and Exclusivity

A Example Subarray Images for 18s



Inclusivity and Exclusivity Testing using printed microarray assay. Microarray results only used cyanine dye without signal amplification. A. Example Subarray image B. Specificity test (top); Sensitivity test (Bottom)



Inclusivity and Exclusivity in plate-based assay. Inclusivity organisms: CTX-M-2 - 1000 copies *Klebsiella pneumoniae* with CTX-M-2 gene; Influenza A - 100 copies H1N1; Orthopox - 1000 copies Vaccinia MVA, 1E8 copies synthetic Variola sequence (equivalent in ng to 5E4 copies); P681R WT - 500 copies Gamma SARS-CoV-2 genome (does not include P681R mutation). Exclusivity Organisms: CTX-M-2 - Pooled 1000 copies of each: *Klebsiella pneumoniae* with CTX-M-15 gene (CTX-M-1 group), *Escherichia coli* with CTX-M-14 gene (CTX-M-9 group); Influenza A - Pooled 5000 copies each: SARS-CoV-2, SARS-CoV, Influenza B, RSV-A, RSV-B, Cov229E; Orthopox - Pooled 5000 copies each: SARS-CoV-2, SARS-CoV, Influenza B, RSV-A, RSV-B, Cov229E; P681R WT - 500 copies Delta SARS-CoV-2 genome (does include P681R mutation). N=3 replicates

Summary

- Dead-gRNA capture assay with signal amplification has shown highly promising results
- Current LoDs indicate 1000 copies and fewer can be detected
- Dynamic range for detection covers over 4 logs
- Specific detection show for DNA virus, RNA, viruses, AMR genes, and sequence variants.

Acknowledgements

Support from the DARPA Biological Technologies Office as part of the Detect It with Gene Editing Technologies (DIGET) program funded under the Naval Information Warfare Center contract N66001-21-1-4048 which is awarded to MRIGlobal. The authors thank Craig Willis, Pamela Winegar, Landon Adebisi, and Sarah Pope for programmatic support, Charles Stark Draper Laboratories for their work on MMD device development and Mammoth Biosciences for reagent and assay design support.

Contact Information

Julie Lucas, Ph.D.
(240) 361-4012
julucas@mriglobal.org

MRIGlobal
65 West Watkins Mill Rd
Gaithersburg, MD 20878

The science you expect. The people you know.