

## Název: ELECTROCHEMISTRY METHOD FOR IDENTIFICATION INFLUENZA VIRUSES:

### CombiMatrix ElectraSense<sup>тм</sup>

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Reg.č.projektu: CZ.1.07/2.4.00/31.0023

Název projektu: Partnerská síť centra excelentního bionanotechnologického výzkumu



#### CombiMatrix ElectroSense<sup>™</sup>: Microarray

MicroArray is a new technology to show the expression of genes.

MicroArray is a hybridization of a nucleic acid sample (target) to a very large set of <u>oligonucleotide probes</u> inside on the chip.



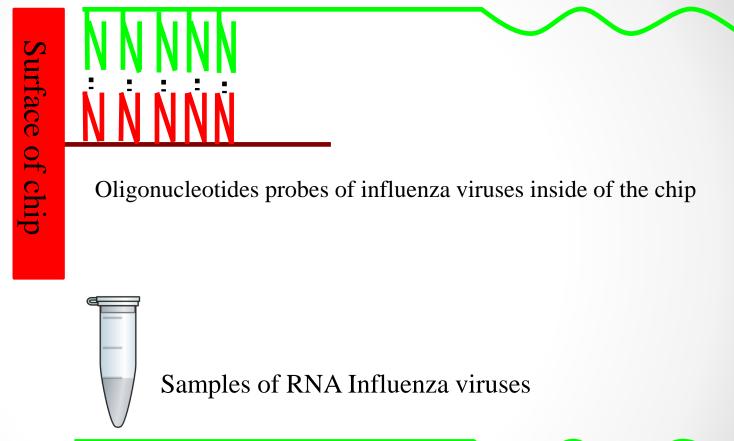
Array Reader

Chip

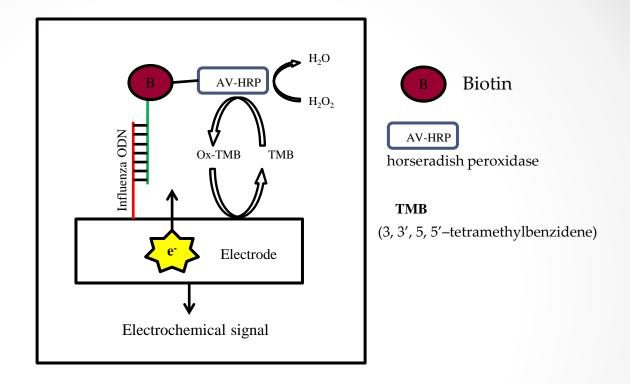
CombiMatrix ElectroSense<sup>TM</sup>: Microarray and hybridization

ΝN





#### CombiMatrix ElectroSense<sup>™</sup>: Electrochemical detection



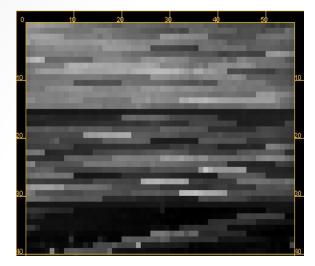
- The approach is based on the detection of redox active chemistries (such as horseradish peroxidase (HRP) and the associated substrate TMB) proximal to specific microarray electrodes.

- Microarray probes are hybridized to biotin-labeled targets.

- The HRP-streptavidin conjugate binds to biotin, and enzymatic oxidation of the electron donor substrate then occurs.

- The detection current is generated due to electro-reduction of the HRP reaction product, and it is measured with the CombiMatrix ElectraSense<sup>TM</sup> Reader.

#### CombiMatrix ElectroSense<sup>™</sup>: Results





Viruses	Genes	GenBank accession manber	Sequence	Number of position inside of Array
Seasonal A(HIN l)	neuraminidase	CY076797	5'-TCCTCATAATGAAAATTGGG-3'	
Pandemic A(HIN I)	ne uraminidase	GQ377078	5'-TCCTCATAGTGATAATTAGG-3'	
HZN2	neuraminidase	CY125896.1	5'-TAGCAATTGCAGGGATCCTA-3'	
H2N8	ne uraminidase	HM589207.1	5'-TGGATCAAGAGGCCATGTTT-3'	
A(H3N2)	neuraminidase	GQ293082	5'-ACACATAAGGTTCTCTTGT-3'	
HINS	neuraminidase	JO433881.1	5'-GTIGAAATCAGTTTCAATGG-3'	
Seasonal A(H5N I)	neuraminidase	CY053327.1	5'-ACCGTTAAAGACAGAAGC C-3'	
HPAI A(H5N1)	neurominidase	FI1263982.1	5'-AGGGAAAGTAGTTAAATCAG-3'	
HPALA(HDN1) H7N7	ne uraminidase	AY340079.1	5'-CCAGAAACATTCTGAGGACT-3'	
HPN2	neuraminidase	NC 004909.1	5'-TACATGATAGGAGTCCCCAT-3'	
Seasonal A(HIN I)	hemaggiutinine	JN017181.1	5'-AGAGAAGAAATAGATGGGGTA-3'	1
Pandemic A(HIN I)	hemaggistinine	JX625498.1	5'-ACCCAAAGCTCAGCAAATC-3'	
H2N2	he maggiotinine	CY125918.1	5'-TGTGTGGACATACAATGCTG-3'	
H2N8	hemagglutinine	HM589205.1	5'-ATCAACACTAAATCAGAGGTC-3'	
A(H3N2)	he maggiutinine	JX844665.1	5'-TTTIGTIGAACGCAGCAAAG-3'	
HINR	hemaggiptinine	JQ433879.1	5'-GACCTTAAAAGCACTCAAGC-3'	
Seasonal A/H5N1)	hemagglotinine	CY053325.1	5'-GGGAATACACCACCCTAAT-3'	
HPAI A(H5N1)	hemaggiotinine	EU263981.1	5'-CTCCAGAATATGCATACAAAA-3'	
H7N7		AY338459.1	5'-GAAACGGTGGAACGAACAAA-3'	
	hemaggiotinine			
H9N2	hemagglutinine	NC_004908.1	5'-GTAGTGCAATGTCAGACTGA-3'	
Seasonal A(HIN l)	nucleocapsid protein	CY087027.1	5'-ATAGACGGAAAATGGATGAG-3'	
Pandemic A(HIN I)	nucleo: apsid protein	JX625388.1	5'-GACGACTAATCCAGAATAGC-3'	
H2N2	nucleo: apsid protein	CY125897.1	5'-AGAGAGTAAATGGAAAGTGGA-3'	2
H2N8	nucleocapsid protein	HM589206.1	5'-AATGGAGAAGACGCAACC-3'	
A(H3N2)	nucleo: apsid protein	CY113840.2	51-GATTATGA AGGGCGGTTGAT-31	
HEN8	nucleo: apaid protein	JO433880.1	5'-CTICAGICTTATCAGACCAAA-3'	
		CV0533281	5'-GCGAATAATGGAGAAGACG-3'	
Seasonal A(H5N l)	nucleocapsid protein			
HPAI A(H5N1)	nucleocapsid protein	EU263985.1	5'-ACATATCAGAGAACGAGAGC-3'	1
H7N7	nucleocapsid protein	AY342425.1	5'-CTAATICGGATGATAAAGCG-3'	
H9N2	nucleo: apsid protein	AF255743.1	5'-GGAAGGTTGATCCAGAAC-3'	
Seasonal A(HIN I)	matrix protein 1	CY087025.1	5'-ACTATACAAGAAGCTCAAAAG-3'	
Pandemic A(HINI)	matrix protein 1	136253901	5'-ATGAAAACAGAATGGTGCTG-3'	
HZN2	matrix protein 1	CY125895.1	5'-TTAAGAGGGAGATAACATTC-3'	
HZN8	matrix protein 1	HM589208 1	5'-CTTIGTCCAAAATGCCCTTAA-3'	
A(H3N2)		CY113838.2		
HINS	matrix protein 1	0433882.1	5'-AACATGTGAACAGATTGCTG-3'	
	matrix protein 1		5'-TACAGGAAGCTTAAAAGGGA-3'	
Seasonal A(H5N l)	matrix protein 1	CY053326.1	5'-TCCTGTCACCTCTGACTA A-3'	1
HPAI A(H5N1)	matrix protein 1	EU263984.1	5'-ACTGCAGCGTAGACGTTTT-3'	1
H7N7	matrix protein 1	GU053111.1	5'-OCCACTAATCAGGCATGAA-3'	
H9N2	matrix protein 1	AJ278646.1	5'-CAAGAAGCTGAAGAGGGAAA-3'	
Seasonal A(HIN I)	matrix protein 2	CY087025.1	5'-TCGCTTTAAATACGGTTTGAA-3'	
Pandemic A(HINI)	matrix protein 2	D0625390.1	5'-TCATTGGGATCTTGCACCT-3'	
H2N2	matrix protein 2	CY125895.1	5'-AACACGGTCTGAAAAGAGGG-3'	
H2N2 H2N8	matrix protein 2	HM589208.1	5'-ATCATEGGATCTEGCACTT-3'	
A(H3N2)	matrix protein 2	CY113838.2	5'-TICAAACACGGTCTGAAAAGA-3'	
HBN8	matrix protein 2	JQ433882.1	5'-GCCTTAAATACGGTTTGAA-3'	
Seasonal A(H5N l)	matrix protein 2	CY053326.1	5'-TATCATTGGGATCTTGCACT-3'	
HPAI A(H5N1)	matrix protein 2	EU263984.1	5'-TCGCCTTAAATACGGTTTGAA-3'	
H7N7	matrix protein 2	GU053111.1	5'-CGCCTTAAATACGGTTTGAA-3'	
H9N2	matrix protein 2	AJ278646.1	5'-CTICAAATGCATITATCGTC-3'	
Seasonal A(HIN I)	polymerase PB1	CY125673.1	5'-AAGATGATGACCAACTCCCA-3'	
Pandenzi (HINI)	polymerase PB1	HQ240702.2	5'-AGTGGAATGAAAATCAAAATC-3'	
H2N2	polymerase PB1	CY125900.1	5'-TGGAATGAGAATCAAAATCCT-3'	
H2N2		HM589203.1		
	polymerase PB1		5'-GGATAAAGAGGAAATGGAAAT-3'	
A(H3N2)	polymerase PB1	CY112971.2	5'-TGAATCTTGGGCAAAAGAAA-3'	
HBN8	polymerase PB1	JQ433877.1	5'-GATGACTAATTCACAAGACA-3'	
Seasonal A(H5N l)	polymerase PB1	JF758819.1	5'-AGCATTGACACTGAACACTA-3'	:
HPAI A(H5N1)	polymerase PB1	EU263987.1	5'-AGAAAATGGTAACACAAAGAA-3'	
H7N7	polymerase PB1	AY340083.1	5'-AATTACTGGAGACAACACCA-3'	
H9N2	polymerase PB1	NC 004911.1	5'-AGTAAGAGCATGAAGCTACG-3'	
Seasonal A/HIN I)	polymerase PA	CY125671.1	5'-AAACCAGACTATTCACCATAA-3'	
Pandemic A(HIN I)	polymerase PA	JO433878.1	5'-ATTCACTGGAGAGAGATGG-3'	
H2N2	polymerase PA	CY125923.1	5'-AGAACATICTITIGGATGGAA-3'	
H2N8	polymerase PA	HM589204.1	5'-GAAAGCCAACAAGATAAAATC-3'	
A(H3N2)	polymerase PA	CY113858.2	5'-CAGAGCCACTGAGTACATAAT-3'	
HBN8	polymerase PA	JQ433878.1	5'-AGGCGAAGAGACAATTGAAG-3'	
Seasonal A(H5N l)	polymerase PA	CY053330.1	5'-TGCATTGAGGGCAAGCTTT-3'	
HPAI A(HSN1)	polymerase PA	EU263986.1	5'-GGGCAAGCTTICTCAAATGT-3'	
H7N7	polymerase PA	AY342418.1	5'-ACAACACCACGCCCTCT-3'	
HPN2	polymerase PA	NC 004912.1	5'-CTGAAATTAAGCATIGAGGAC-3'	
Seasonal A(HIN I)		CV125670.1	SUGATIBA AGA AGTBAGACACA A.3/	
	nuclear export protein (nep)			
Pandemic A(HIN I)	nuclear export protein (nep)	HQ240288.2	5'-TGGTTAATTGAAGAAATGCGG-3'	
H2N2	nuclear export protein (nep)	CY125898.1	5'-ACGGAAAATGGCGAGAACAA-3'	
H2N8	nuclear export protein (nep)	HIM589209.1	5'-TGATTGAGGAAGTACGACATA-3'	
A(H3N2)	nuclear export protein (nep)	CY103967.1	5'- AGAACAGTTAGGTCAAAAGT-3'	
H3N8	nuclear export protein (nep)	JQ433883.1	5'-TGGCTGATTGAAGAAGTGC-3'	
Seasonal A(H5N I)	nuclear export protein (nep)	CY053329.1	5'-ACCTCCACTCCCTCCAAAA-3'	
HPAI A(H5N1)	naciaal export protect(fiep)	CY093529.1 CY098518.1	5'-GCTGATTGAAGAAGTACGACA-3'	
	nuclear export protein (nep)			
H7N7	nuclear export protein (nep)	GU053125.1	5'-AAGTGCGACATAGGTTGAAG-3'	
H9N2	nuclear export protein (nep)	FJ793288.1	5'-TGGCTGATTGAAGAAGTGC-3'	

#### CombiMatrix ElectroSense<sup>™</sup>: Review of Results

<u>Kristian M. Roth et al</u>, demonstrated very good detection of avian influenza subtype H5N1 by application of electrochemical detection of Combimatrix microarrays



#### **Electrochemical Detection of Short DNA Oligomer Hybridization** Using the CombiMatrix ElectraSense Microarray Reader

Kristian M. Roth,\* Kia Peyvan, Kevin R. Schwarzkopf, Andrei Ghindilis Combimatrix Corp., 6500 Harbour Heights Pkwy Mukilteo, WA 98275 \*e-mail: kroth@combimatrix.com

<u>Michael J. Lodes et al</u>, identified of upper respiratory tract pathogens by electrochemical detection on an oligonucleotide microarray, including A and B influenza viruses



Identification of Upper Respiratory Tract Pathogens Using Electrochemical Detection on an Oligonucleotide Microarray

Michael J. Lodes<sup>1</sup>", Dominic Suciu<sup>1</sup>, Jodi L. Wilmoth<sup>1</sup>, Marty Ross<sup>1</sup>, Sandra Munro<sup>1</sup>, Kim Dix<sup>1</sup>, Karen Bernards<sup>1</sup>, Axel G. Stöver<sup>1</sup>, Miguel Quintana<sup>2</sup>, Naomi lihoshi<sup>2</sup>, Wanda J. Lyon<sup>3</sup>, David L. Danley<sup>1</sup>, Andrew McShea<sup>1</sup>

1 CombiMatrix Corporation, Mukilteo, Washington, United States of America, 2 United States Army Center for Health Promotion and Preventive Medicine-West, Fort Lewis, Washington, United States of America, 3 Air Force Research Laboratory/Human Effectiveness Directorate, Applied Biotechnology Branch, Wright-Patterson Air Force Base, Ohio, United States of America CombiMatrix ElectroSense<sup>™</sup>: Review of Results

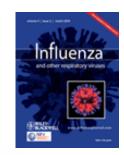
<u>Shelly Bolotin et al</u>, showed that the sensitivity of the CombiMatrix influenza detection system was 95.2% and the specificity was 100% for influenza A subtype during the 2007–2008 influenza season in Toronto, Canada



Verification of the Combimatrix influenza detection assay for the detection of influenza A subtype during the 2007–2008 influenza season in Toronto, Canada Shelly Bolotin<sup>\*1</sup>, Ernesto Lombos<sup>1</sup>, Rani Yeung<sup>1</sup>, AliReza Eshaghi<sup>1</sup>,

Joanne Blair<sup>1</sup> and Steven J Drews<sup>1,2,3</sup>

<u>Straight, T.M. et al</u>, correctly identified 23 of 24 samples of laboratory-confirmed pandemic (H1N1) 2009 Influenza by the ElectraSense Influenza A assay.



## A novel electrochemical device to differentiate pandemic (H1N1) 2009 from seasonal influenza

T. M. Straight1, G. Merrill1,L. Perez1, J. Livezey1, B. Robinson1,M. Lodes2,D. Suciu2,B. Anderson2

#### CombiMatrix ElectroSense<sup>TM</sup>: Conclusions

The CombiMatrix influenza detection system is an effective methodology for influenza A and B subtype analysis

Rapid identification pathogens will significantly decrease the time and cost for the identification of potential lethal virus and bacterial strains and lead to better treatment and management of infections

> Microarray and biosensor technologies show great promise for virus and bacteria detection and genotyping and are needed for rapid effective treatment, environmental monitoring and the detection of bioterrorism agents.

#### CombiMatrix ElectroSense<sup>TM</sup>: Acknowledgements



I want to thank all colleagues of the laboratory

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# Thank you for attention

NanoBioMetalNet

Reg.č.projektu: CZ.1.07/2.4.00/31.0023

Název projektu: Partnerská síť centra excelentního bionanotechnologického výzkumu