

Regulation of Human Gene of Influenza A Virus-Associated NS1-Binding Protein by 2,3,7,8-TCDD: Mechanistic Data and Epidemiological Findings

Jerry L. Wu¹, Andre G. Pokrovsky², Vladimir S. Rumak³, Ilya B. Tsyrllov¹

¹XENOTOX Inc., Scarsdale, U.S.A. (e-mail: xenotoxit@optonline.net). ²Novosibirsk State University, Russia.

³Research Tropical Center, Hanoi, Vietnam

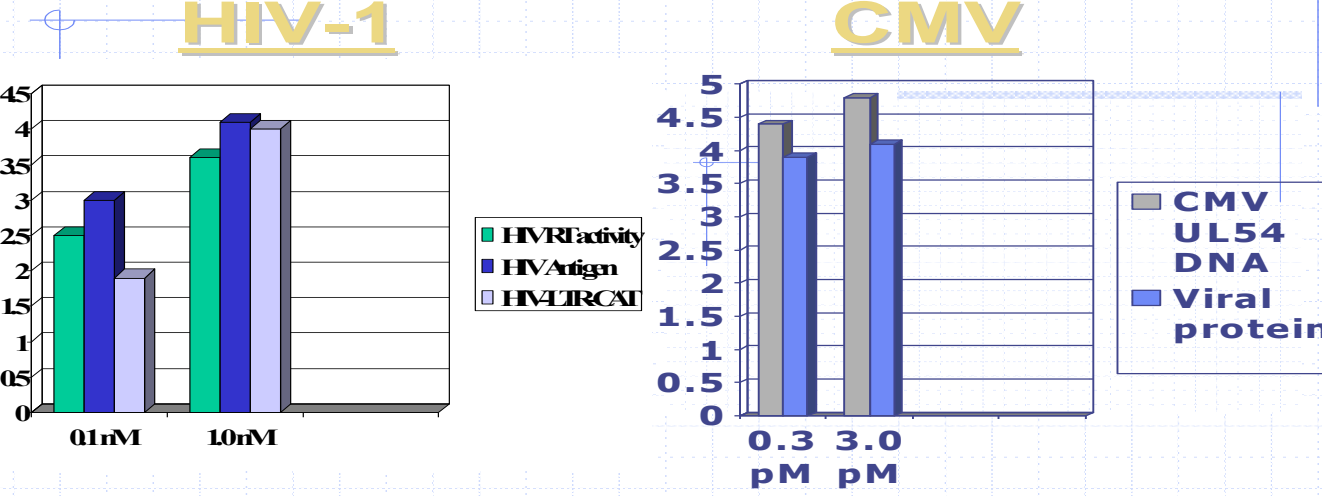
Earlier we postulated that 2,3,7,8-tetrachlorodibenzo-p-dioxin (TCDD) might transactivate HIV-1 gene expression in HIV-infected human cells. The results have been confirmed in cell models with HIV-1 and CMV, and a clinical study on HIV-AIDS patients.

Ah receptor (AhR)-mediated transcriptional response network was shown in HIV-1 and CMV activation by TCDD, however, mechanistic concept was formulated after identifying “dioxin response elements” (DRE), core nucleotide sequence 3’ A-CGCAC 5’, in promoter distal region of HIV-1, CMV and several cancer-associated common human viruses.

Here, the above mechanistic concept is applied to genetically similar genes encoding human and chicken influenza virus NS1 binding protein (NS1BP), as two DREs were revealed in 5’-flanking region (at positions - 7942 and - 687).

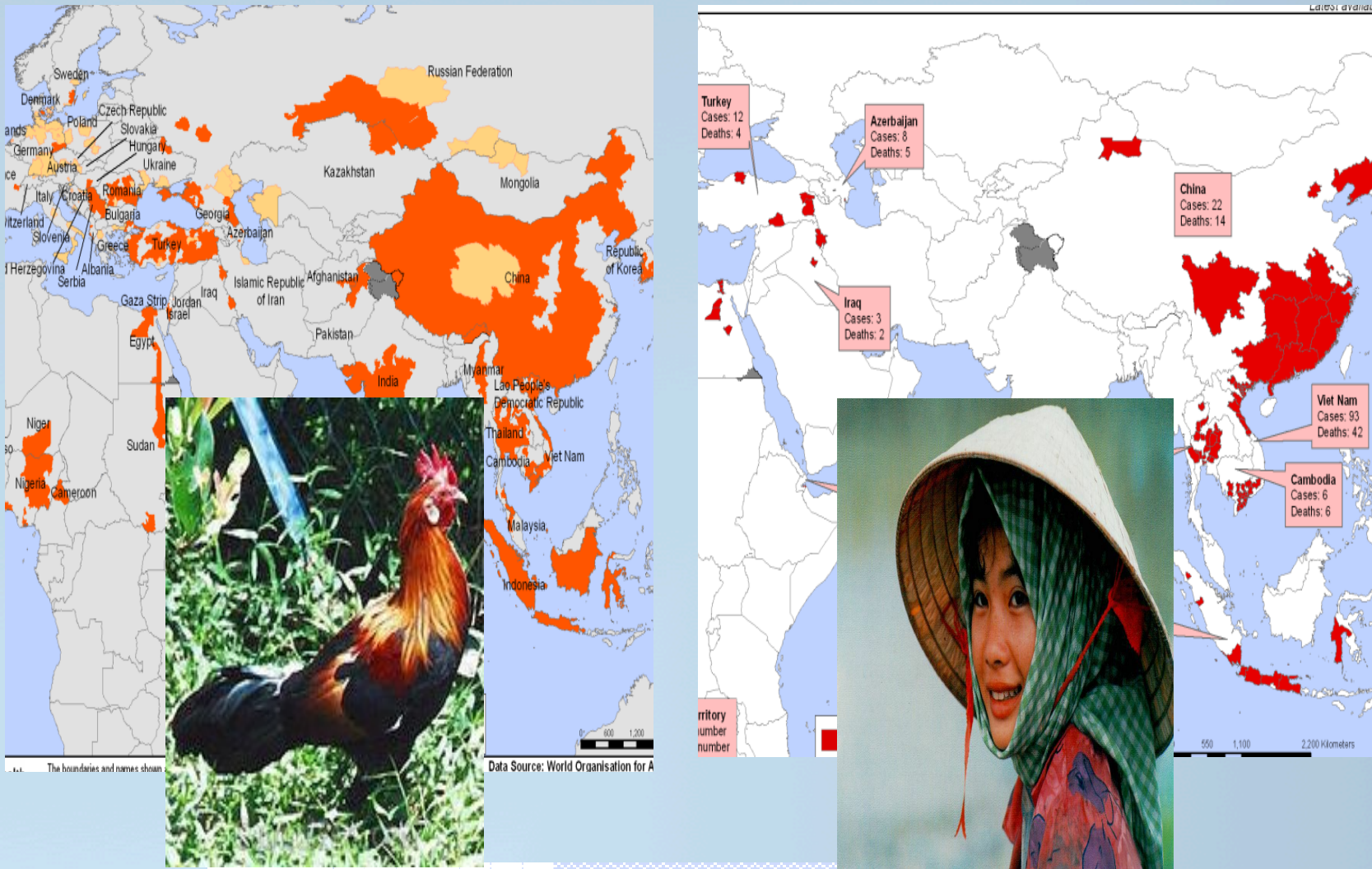
PREVIOUS DATA ON HUMAN VIRUSES

TCDD trans-activates replication of human viruses



From: Pokrovsky, Tsyrllov et al., *BBRC* 1991; Tsyrllov & Pokrovsky, *Xenobiotica* 1993; Gollapudi et al., *BBRC* 1996; Murayama et al., *BBRC* 2002; Ohata et al., *Microbiol. Immunol.* 2003; Tsyrllov, *Organohalogen Compounds* 2006

EPIDEMIOLOGIC CONCEPT



Gallus gallus versus Homo sapiens Influenza Virus A NS1 BP Genomic Alignment:



DREs IN PROMOTER REGION OF VIRUSES AND NS1BP

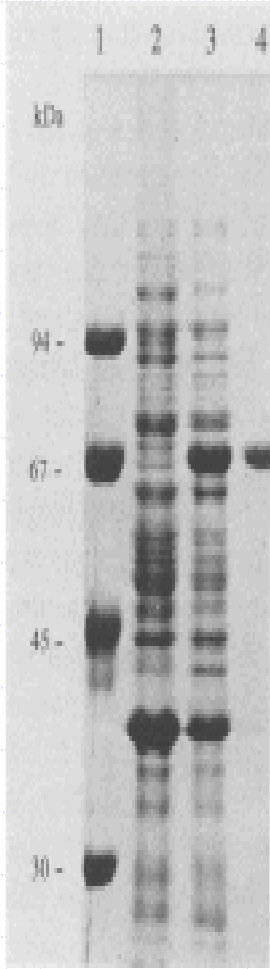
Organism summary of the dioxin response element (DRE) core sequence (5'-GCGTG-3') found in viral promoters in the Eukaryotic Promoter Database (from T. Zacharewski, 2002)		
Species	# DREs Located	# Promoters Represented
Human adenovirus type 12	10	4
Human adenovirus type 2	36	9
Human adenovirus type 5	19	5
Human adenovirus type 7	12	3
Human adenovirus type 12	3	1
Human herpesvirus 4	154	22
Human herpesvirus 8	6	2
Human herpesvirus 13	6	2
Human cytomegalovirus	102	10
Human herpes simplex virus type 1	345	30
Human herpes simplex virus type 2	38	8
Human cytomegalovirus	3	1
Bovine papillomavirus type 1	15	6
Human Papillomavirus type 16	3	1
Human Papillomavirus type 18	9	2
Mouse polyoma virus	1	1
Human immunodeficiency virus type 1	4	2
Human immunodeficiency virus type 2	9	1
Human immunodeficiency virus type 3	1	1
Human immunodeficiency virus type 2	2	1
Human immunodeficiency virus type 1	7	1
Human immunodeficiency virus type 2	2	1
Human immunodeficiency virus type 1	7	1
Bovine leukemia virus	1	1
Gibbon ape leukemia virus	1	1
Human T-cell leukemia virus type I	4	1

Positions of Dioxin Response Elements (DRE) in INNS1BP Gene			
Genomic Information: Gene title: Influenza virus NS1A binding protein Genomic full length in this probe set: 18,762 bp Exon number: 15			
Type	Length (bp)	DRE number	DRE Position
5'-flanking	10,000	2	-7942, -687
Exon 1	200	2	416, 1465
Exon 2	128	2	416, 1465
Exon 3	1576	1	569
Exon 4	173	0	
Exon 5	102	0	
Exon 6	275	0	
Exon 7	174	0	
Exon 8	108	0	
Exon 9	129	0	
Exon 10	129	0	
Exon 11	129	0	
Exon 12	129	0	
Exon 13	129	0	
Exon 14	129	0	
Exon 15	129	0	
3'-flanking	10,000	2	381, 2041

OVERLAPPING SPOTS OF H5N1 OUTBREAKS AND DIOXIN CONTAMINATION IN CHINA AND VIETNAM

NEW FINDINGS ON TCDD-NS1BP

Immunoblotting of TCDD-Induced human NS1BP



Confluent monolayers of HeLa cells, pretreated with 0.1 nM TCDD for 36 h (column 3), were lysed in RIA-precipitation assay buffer. Soluble proteins from equivalent volumes of extract corresponding to 10⁴ cells were separated by SDS gel electrophoresis, transferred to a nitrocellulose membrane, and probed with affinity-purified NS1BP-specific antibodies. The marker proteins - column 1; non-treated HeLa cells - column 2; the recombinant NS1BP - column 4.

Dioxin Exposure Level in Guangdong Province, China's First H5N1 Outbreak

“The Asian H5N1 virus was first detected in Guangdong Province, China, in 1996, but it received little attention until it spread through live-poultry markets in Hong Kong to humans in May 1997, killing 6 of 18 infected persons. From 1997 to May 2005, H5N1 viruses were largely confined to Southeast Asia, but after they had infected wild birds in Qinghai Lake, China, they rapidly spread westward”.

“According to the Chinese Ministry of Health, human bird flu infection confirmed in Guangdong Province. A 32-year-old man died of bird flu infection, H5N1. The man was from the province of Guangdong”.

“Mothers born in Chinese Guangdong province, have had a significantly higher CALUX-TEQ (a luciferase expression bioassay for maternal exposure to dioxin and dioxin body load). Higher seafood consumption was associated with a higher maternal CALUX-TEQ level”.



Content of Dioxins in Humans and Chicken from Non-Treated Northern (■) and Agent Orange-Treated Southern (■) Regions of Vietnam (2005)

Dioxin Level (TCDD-based TEQ, ppt):

Northern Provinces (number on map):

- Ha Noi (17)

- Thanh Hoa (27)

- Nghe An (28)

Human blood (n = 168): 1.20 ± 0.29

Chicken fat tissues: 2.02 ± 0.22

Chicken eggs: 0.015 ± 0.002

Southern Provinces (number on map):

- Tay Ninh (48)

- Binh Duong (49)

- Can Tho (60)

Human blood (n = 1210): 3.53 ± 1.04

Chicken fat tissues: 3.13 ± 0.43

Chicken eggs: 0.35 ± 0.04

On January 6, 2004, WHO was informed of suspicious chicken deaths in southern Vietnam, in provinces shown on this map with beige arrow beneath, namely *Gong Thap* (number on map - 53), *Tien Giang* (54), and *Ben Tre* (59). By January 8, Vietnam informed WHO that birds on two farms of these southern provinces were infected with H5N1. 70,000 birds died or were destroyed to prevent the spread of the disease.

On January 11, Vietnam advised WHO that humans had been admitted to a hospital with a severe respiratory infection. On the same day, it was announced that tests on two samples from two fatal cases of respiratory infections in Vietnam proved positive for the H5N1 virus.

The southern cases had a case fatality rate approaching 100%, while the fatality rate in northern Vietnam fell to 10-20%.

MECHANISTIC CONCEPT

Non-structural protein 1



NS1, a dimeric protein consisting of six helical folds, is encoded by the influenza A viruses. NS1 is a RNA-binding protein required for virus replication. Its RNA-binding activities are:

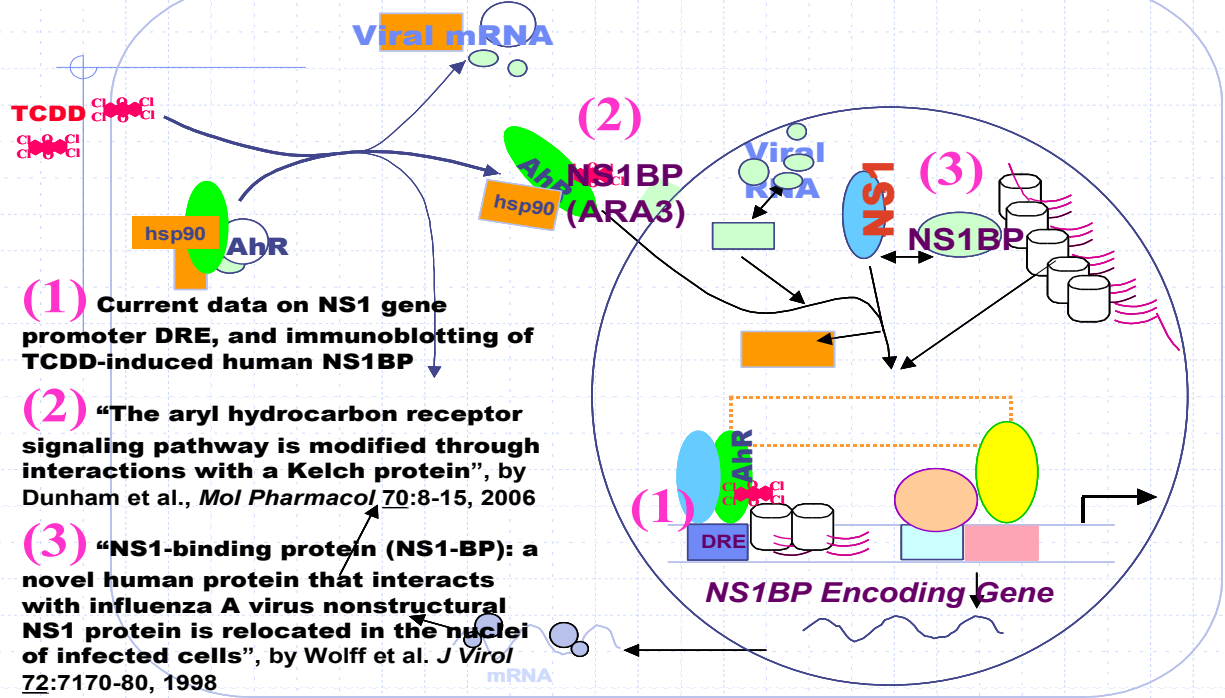
NS1 binds to the poly A tails of cellular pre-mRNAs. Thus it retains these pre-mRNAs within the cell's nucleus by blocking their export into the cytoplasm.

NS1 blocks pre-mRNA splicing by binding to a stem-bulge in U6 small nuclear RNA (snRNA) thus inactivating any U6-U2 or U6-U4 interactions that are required for splicing.

NS1 can bind double-stranded RNA (dsRNA) with a low sequence specificity while maintaining a high affinity which blocks the activation of PKR kinase.

Three steps body burden TCDD might facilitate flu viremia:

Transactivation of NS1BP gene thru promoter DRE (1); Physical association of NS1BP (ARA3) with AhR (2); antiviral defense-dismantling interaction of induced NS1BP with viral NS1 (3)



ESTIMATED RELATIVE SUSCEPTIBILITY to TCDD of VIRAL (HIV-1 and CMV) and HUMAN (NS1BP and CYP1A1) GENES

Name (symbol) of the Gene	HIV-1	CMV	NS1BP	CYP1A1
Quantity of DRE(s) in gene enhancer	1	10	2	2
Minimal gene - activating concentration of TCDD	0.1 nM or 32 ng/kg	0.3 pM or 0.1 ng/kg	33 pM or 10 ng/kg	33 pM or 10 ng/kg

CONCLUDING REMARKS

The above information on domestic poultry dying from H5N1 in Guangdong province of China, and Long An, Tieng Giang and Ben Tre provinces of Vietnam might relate to the fact that water and soil in these very regions in China and Vietnam are highly contaminated with dioxin-like compounds.

As human NS1BP gene promoter contains two DRE, an extrapolation from the data on HIV-1 (1 DRE) and CMV (10 DRE) genes suggested that TCDD level upregulating NS1BP gene should be moderately above current TCDD blood levels in general population. The data revealed here show that NS1BP might be induced in human HeLa cells with 0.1 nM TCDD. The same concentration of TCDD has been shown earlier to induce human microsomal cytochrome P4501A1, encoded by CYP1A1 gene, the promoter region of which also possesses two DRE.

Thus in human subgroups slightly exposed to TCDD, its body level is a possible promotional factor for seasonal influenza virus A outbreaks, and may strongly facilitate spreading of the H5N1 if avian flu pandemic were to occur. This is because the resistance of highly virulent H5N1 to antiviral effects of IFN-β and TNF-α is directly associated with viral NS1.