RELEVANCE OF STX2-PHAGE IN THE DEVELOPMENT OF HEMOLYTIC UREMIC SYNDROME (HUS)

Ms. Carla Ivanna Rivero

National University of José C. Paz.

Institute of Studies for Productive Development and Innovation

Buenos Aires, Argentina

email: carlairivero@gmail.com



BACTERIOPHAGES AND EUKARYOTIC CELLS

Bacteriophages: bacterial virus



What is happening in eukaryotic environment?



BACTERIOPHAGES AND EUKARYOTIC CELLS

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Bacterial Virus Gene Expression in Human Cells

CARL R. MERRIL & MARK R. GEIER

Laboratory of General and Comparative Biochemistry, National Institute of Mental Health, Bethesda, Maryland 20014

JOHN C. PETRICCIANI

Laboratory of Pathology, Division of Biologics Standards, National Institutes of Health, Bethesda, Maryland 20014

Human fibroblasts, from a patient with congenital lack of α -D-galactose-1-phosphate uridyl transferase activity, have been infected with transducing bacteriophage that harbours either wild type or defective transferase gene. Infection only by the former phage initiates transferase synthesis.

Gal-1-P it is possible to follow this reaction (Fig. 2). This activity was linear with time in the conditions used and proportional to the amount of cell lysate used (Fig. 3). All activity was lost by boiling the cell lysate. Uninfected galactosemic cells had an activity ≤ 0.2 nmol of UDP-Gal/60 min/mg of protein (Table 1). No difference in enzyme activity could be detected between uninfected galactosemic fibroblasts and fibroblasts infected with λ virus or λ pgal T⁺ DNA-infected in transferase structural gene. The λ pgal T⁺ DNA-infected λ pgal T⁺ whole virus-infected cells. Infected cell cultures have

Lambda Phage Transcription In Human Fibroblasts

MARK R. GEIER¹ AND CARL R. MERRIL

Laboratory of General and Comparative Biochemistry, National Institute of Mental Health, Bethesda, Maryland 20014

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Lambda bacteriophage was used to infect human fibroblasts and viral specific RNA was detected by hybridization with lambda DNA. A correlation was found between the amount of virus used in each infection and the percentage of lambda-specific RNA produced. The percentage of lambda-specific RNA (0.2%) reached a maximum after 4 days. Cells monitored for 41 days following infection continue to produce lambda-specific RNA.

Is able to express genes under eukaryotic promoter



So, once inside the cell, host cell is able to express genes under eukaryotic promotors

PREVIOUS IDEAS



THAT WE MAKE US ARRIVE OUR OBJETIVES

- Shiga toxins (Stx) are the main virulence factors in enterohemorrhagic Escherichia coli (EHEC) infections causing diarrhea, hemorrhagic colitis, and Hemolytic Uremic Syndrome (HUS)
- > Stx genes are located in the genomes of prophages that resemble the coliphage lambda
- > Under stress condition, the bacteriophage release and Stx is express
- > Bacteriophage could infect others bacterias present in the gut (Gamage *et al.*, 2003)
- > Bacteriophage induction is required for HUS development (Tyler *et al.*, 2013).



OUR HYPOTHESIS + PREVIOUS IDEAS

WE DECIDED OUR OBJETIVES



2 Studies focus in the expression of Stx in eukaryotic cells



Studies focused on the bacteriophage as therapeutic target



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Functional Capacity of Shiga-Toxin Promoter Sequences in Eukaryotic Cells

Leticia V. Bentancor¹*³, Marcos F. Bilen²³, María P. Mejías¹, Romina J. Fernández-Brando¹, Cecilia A. Panek¹, Maria V. Ramos¹, Gabriela C. Fernández¹, Martín Isturiz¹, Pablo D. Ghiringhelli², Marina S. Palermo¹

1 División Inmunología, Instituto de Medicina Experimental (IMEX) (CONICET), Academia Nacional de Medicina, Buenos Aires, Argentina, 2 Laboratorio de Ingeniería Genética y Biología Celular y Molecular, Universidad Nacional de Quilmes, Buenos Aires, Argentina

Evaluate the Ability of the eukaryotic machinery to recognize genetic sequences as Stx2 promoters



To transcribe a Stx2-like protein and produce the functionally active toxin.



Tested by GFP expression

PLASMID CONSTRUCTIONS

Plasmid pStx2



PLASMID CONSTRUCTIONS

GFP activity driven by the pr1 or pr7 regions



293T cells were transfected with the next plasmids - (Incubated for 48h)

pr1-eGFP Treatment groups pr7-eGFP

 $\Delta pr-eGFP \rightarrow$ As negative control

Numbers

- 1 correspond to images visualized with white light
- 2 correspond to images visualized with green filter
- 3 correspond to merge between DAPI and green filter
- 4 correspond to merge between white light and green filter

A. Cells transfected with the Δpr -eGFP plasmid.

B. Cells transfected with pr1-eGFP.

C. Cells transfected with pr7-eGFP.

PSTX2: CYTOTOXICITY ON SUSCEPTIBLE EUKARYOTIC CELLS

Cells were stained

with Crystal Violet

Cytotoxicity on Vero Cells

Vero cells were 48 h incubated with purified Stx2 or supernantants of cells transfected with pStx2 plasmid.



Analyzed by optical microscopy. Representative pictures using 200X original magnification are shown.



- A. Non-treated Vero cells.
- B. Cells transfected with the pGEM-T plasmid.
- C. Cells transfected with the pStx2 plasmid.
- D. Cells incubated with purified Stx2.



PSTX2: CYTOTOXICITY ON SUSCEPTIBLE EUKARYOTIC CELLS

OD of the samples with antibodies is more higher than the citotoxic samples.



This results demonstrated that Stx is expressed in eukaryiotic keeping this structure, function and the capacity to bind the receptor.

CONCLUSIONS - IN VITRO ASSAYS

- pr1 and pr7 sequences are recognized by eukaryotic cells
- Stx2 expressed by eukaryotic cells is biologically active
- Significant neutralization of the cytotoxic activity was observed using mouse polyclonal anti-Stx2 antibodies, confirming that the cytotoxicity was specifically induced by Stx2.
- TATA box depletion did not show GFP expression

What would happen *In vivo*?

So, for this reason we did the same but *In vivo* in a Murine model.



RESEARCH ARTICLE

Promoter Sequence of Shiga Toxin 2 (Stx2) Is Recognized *In Vivo*, Leading to Production of Biologically Active Stx2

Leticia V. Bentancor,^{a,b} Maria P. Mejías,^a Alípio Pinto,^c Marcos F. Bilen,^b Roberto Meiss,^a Maria C. Rodriguez-Galán,^d Natalia Baez,^d Luciano P. Pedrotti,^d Jorge Goldstein,^c Pablo D. Ghiringhelli,^b Marina S. Palermo^a

División Inmunología, Instituto de Medicina Experimental (IMEX) (CONICET), Academia Nacional de Medicina, Buenos Aires, Argentina^a; Laboratorio de Ingeniería Genética y Biología Celular y Molecular, Universidad Nacional de Quilmes, Buenos Aires, Argentina^b; Laboratorio de Neurofisiopatología, Departamento de Fisiología, Facultad de Medicina, Universidad de Buenos Aires, Ciudad Autónoma de Buenos Aires, Buenos Aires, Argentina^c; Centro de Investigaciones en Bioquímica Clínica e Inmunología, Consejo Nacional de Investigaciones Científicas y Técnicas, Departamento de Bioquímica Clínica, Facultad de Ciencias Químicas, Universidad Nacional de Córdoba, Córdoba, Argentina^d

PSTX2: HYDRODYNAMIC INOCULATION IN MOUSE



after 24 hours.

Letal doses depended of concentration

- 1ld100 the mice were dead after 72 hours
- 3ld100 the mice were dead after 96 hours

IN VITRO EVALUATION OF STX2 ACTIVITY IN PLASMA



In vitro evaluation of Stx2 activity in plasma.

(A) Vero cells were incubated with plasma from mice inoculated with different doses of pStx2 obtained at 24 h post-inoculation. Each bar represents the mean \pm SEM for 4 mice. *, P < 0.03, t test.

CONCLUSIONS - IN VIVO ASSAYS

- Stx2 is expressed in vivo after pStx2 HBT, driven by its own wild stx2 promoter.
- Active Stx2 thus produced is able to target the kidney and the brain and reproduces the lethal damage induced by purified Stx2 or secondary to EHEC infection.

WHICH IS THE ROLE OF BACTERIOPHAGE 933W IN HUS DEVELOPMENT?

IS BACTERIOPHAGE 933W ENOUGH TO DEVELOP THE SYNDROME?



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Relevance of Bacteriophage 933W in the Development of Hemolytic Uremic Syndrome (HUS)

Manuel E. Del Cogliano^{1†‡}, Alipio Pinto^{2†‡}, Jorge Goldstein², Elsa Zotta³, Federico Ochoa³, Romina Jimena Fernández-Brando⁴, Maite Muniesa⁵, Pablo D. Ghiringhelli¹, Marina S. Palermo⁴ and Leticia V. Bentancor^{1*}



We worked with a non pathogenic *Eschericia coli* strain carrying the lysogenic bacteriophage 933W (C600:933W) and we induced the release of the bacteriophage using antibiotics.

SURVIVAL CURVE AND PHYSIOLOGICAL PARAMETERS OF HUS



Survival rates of mice infected with E. coli C600 Φ 933W are shown. Mice infected with E. coli C600 were used as controls.

we analyze urea levels

DETECTION OF STX2 IN BRAINS



We performed an immunofluorescence assay using a polyclonal anti-Stx2 antibody.

Were observed inmunopositive cells in mice infected with C600:933W \rightarrow Like in + Control

Immunopositive cells for Stx2 were not found in control mice infected with C600

Detection of Stx2 in brains of mice infected with E. coli C600Ф933W.

- (A–C) A representative brain section from control mouse infected with *E. coli* C600.
- (D–F) Brain section from mouse infected with *E. coli* C600Ф933W.
- (G–I) Brain section from a positive control mouse after 4 days of treatment with 1 μg of Stx2.

(J) Negative control.

(K) Number of Stx2-immunopositive cells in the mouse motor cortex (mc) per micrograph. Stx2 immunopositive cells are indicated by arrows in (F,I). (L)

The area observed is located in the mouse motor cortex. The scale bar in "J" applies to all 20 micrographs.

BACTERIOPHAGE DETECTION IN BRAINS

We used antibodies for phage detection

Bacteriophage 933W was observed in mice infected with C600:933W

The controls were negatives

Bacteriophage detection in brains of mice infected with E. coli C600Ф933W.

(A,C,D) Brain from control mouse infected with E. coli C600.

(E,L) Brain from mouse infected with E. coli C600Ф933W 72 h after inoculation.

(E–H) External capsule;

(I-L) hypothalamus.

(B) Negative control omitting the secondary antibody.

(N) Bacteriophage were detected in the mouse corpus callosum (cc) and hypothalamus (hy) (the red square shows where the micrographs were taken).

(M) Quantification of bacteriophages in the cerebral parenchyma by integral optical density (IOD). Iv, lateral ventricle; gl, glia limitans. The scale bar in "B" applies to all micrographs.

CONCLUSIONS BACTERIOPHAGE IN VIVO

- In absence of other pathogenic bacterial factors, bacteriophage 933W provoked intestinal, renal and brain damage associated with HUS development.
- It is the first detection of bacteriophage in brain.
- The new animal model described here is proposed as a useful tool to study bacteriophages as a therapeutic target, independently of the bacterial background, and opens an innovative approach to treating infections caused by toxin-encoding bacteriophages.

Universidad Nacional de José C. Paz Leticia Bentancor Alejandra Capozzo Carla Rivero

Academia Nacional de Medicina Marina Palermo María Pilar Mejías Romina J. Fernández-Brando

LIGBCM-UNQUI Manuel Del Cogliano Pablo D. Ghiringhelli

В

Universitat de Barcelona

Laboratorio de Desarrollo de Vacunas Luis Carlos de Souza Ferreira Rita Ferreira Jaime H. Amorim Monica J.R. Rodrigues

Lab. de Neurofisiopatología Fac. de Medicina, UBA Jorge Goldstein Alipio Pinto

CONICET

A G E N C I A NACIONAL DE PROMOCION CIENTIFICA Y TECNOLOGICA

Maite Muniesa Pérez

Pablo Quiróz

Departamento de Microbiología

http://redargentinadebacteriofagos.web.unq.edu.ar