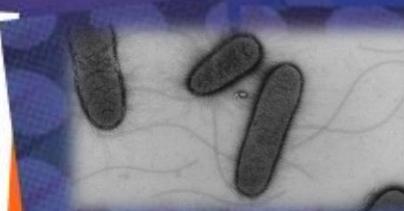


UNIVERSITAIR MEDISCH CENTRUM GRONINGEN

# Zoonosis from the ground

Alex W. Friedrich  
Medical Microbiology  
University Medical Center Groningen

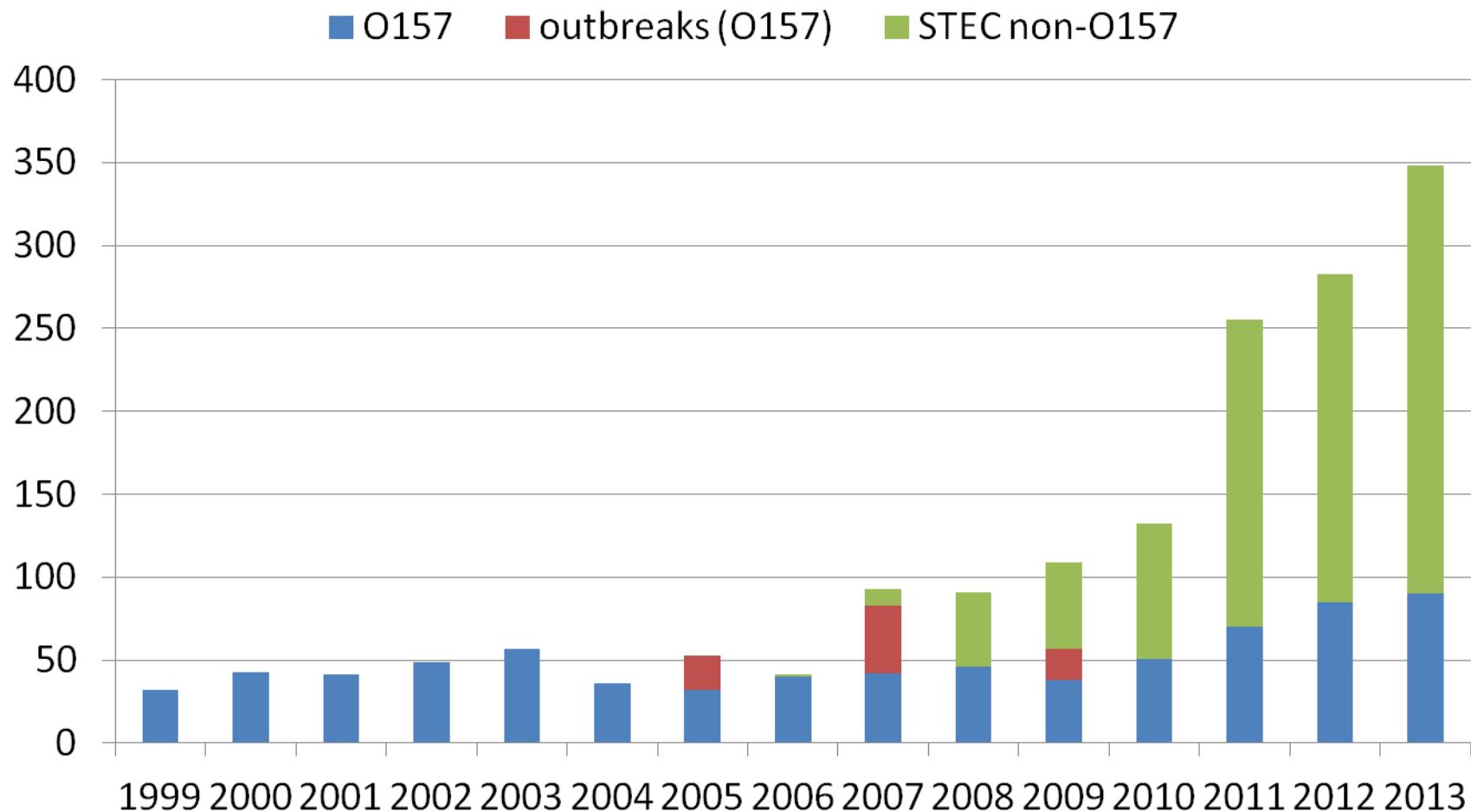
[alex.friedrich@umcg.nl](mailto:alex.friedrich@umcg.nl)



# Reported hospitalisation and case-fatality rates due to zoonoses in confirmed human cases in the EU, 2011

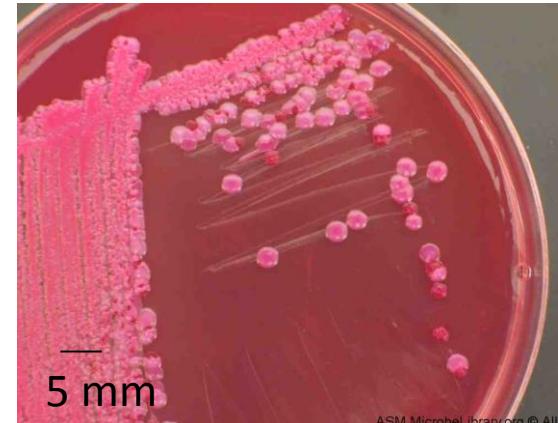
Zoonotic Diseases	Causative agent	Confirmed human cases	Reported hospitalised cases	Hospitalisation rate (%)	Reported deaths	Case fatality rate (%)
Campylobacteriosis	Campylobacter spp.	220,209	8,137	47.9	43	0.04
Salmonellosis	Salmonella spp.	95,548	4,557	45.7	56	0.12
STEC infections	Shiga toxin-producing <i>E. coli</i>	9,485	721	33.8	56	0.75
Yersiniosis	<i>Yersinia</i> spp.	7,017	427	55.2	1	0.02
Listeriosis	<i>L. monocytogenes</i>	1,476	604	93.6	134	12.7
Echinococcosis	<i>Echinococcus</i> spp.	781	96	67.6	2	0.90
Brucellosis	<i>Brucella</i> spp.	330	118	66.3	1	0.74
Trichinellosis	<i>Trichinella</i> spp.	268	153	74.3	1	0.49

# STEC reports in the Netherlands



Data: Ingrid Friesema, RIVM

# Infections due to *E. coli*

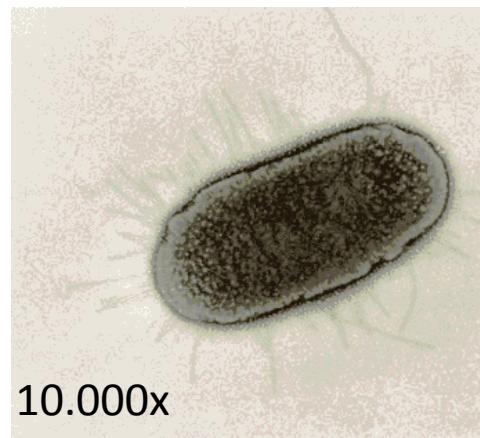


Urinary tract infections

Neonatal encephalitis

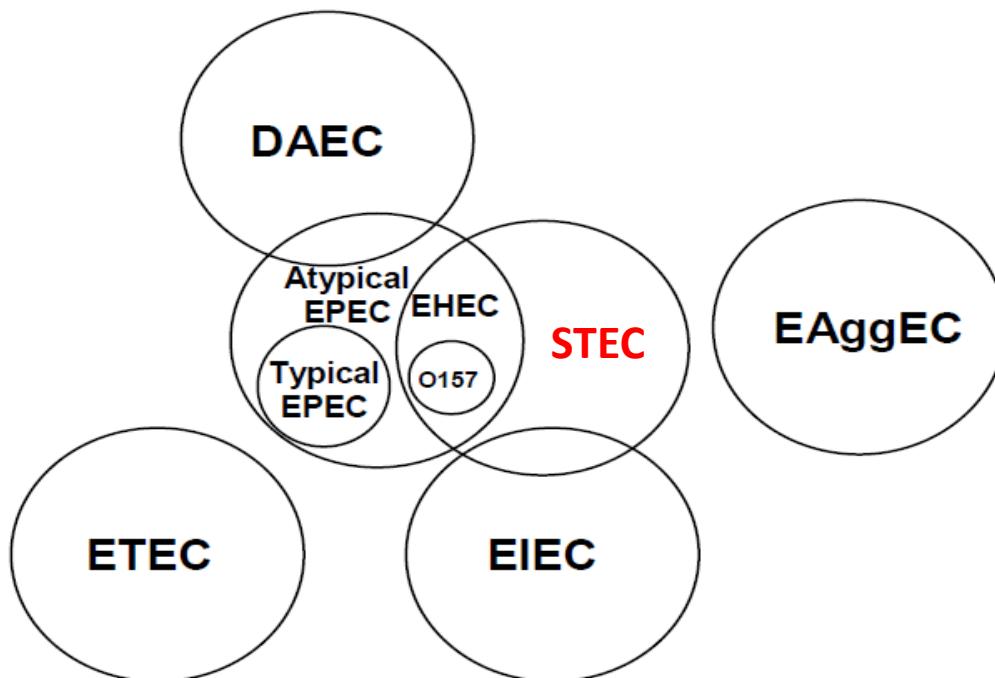
Enteropathogenic *E. coli*

Other opportunistic infections



# Enteropathogenic *E. coli*

- Enterotoxigenic *E. coli* (ETEC)
- Enteropathogenic *E. coli* (EPEC)
- Shiga toxin-producing (STEC) /Enterohaemorrhagic *E. coli* (EHEC)/ VTEC
- Enteroaggregative *E. coli* (EAEC)
  - children in 3rd world, in HIV patients, chron. Diarrhea. Malnutrition, highly antibiotic resistant
- Enteroinvasive *E. coli* (EIEC)



# EHEC O157:H7 outbreak in Canada



7 died, 2000 with diarrhea  
cause: contaminated water

# Clinical importance and danger to public health

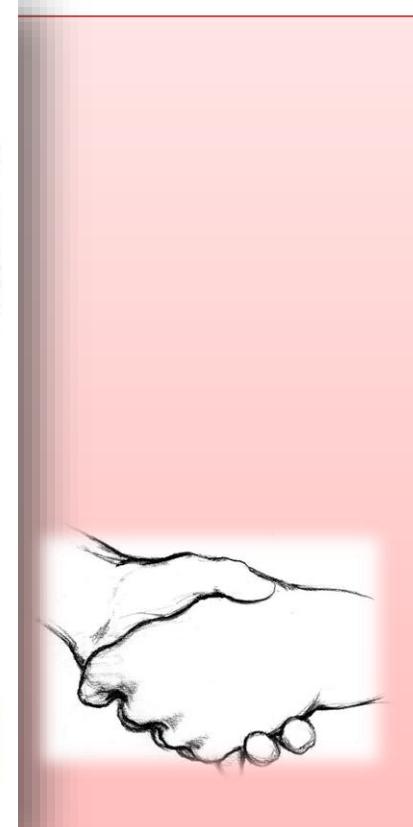
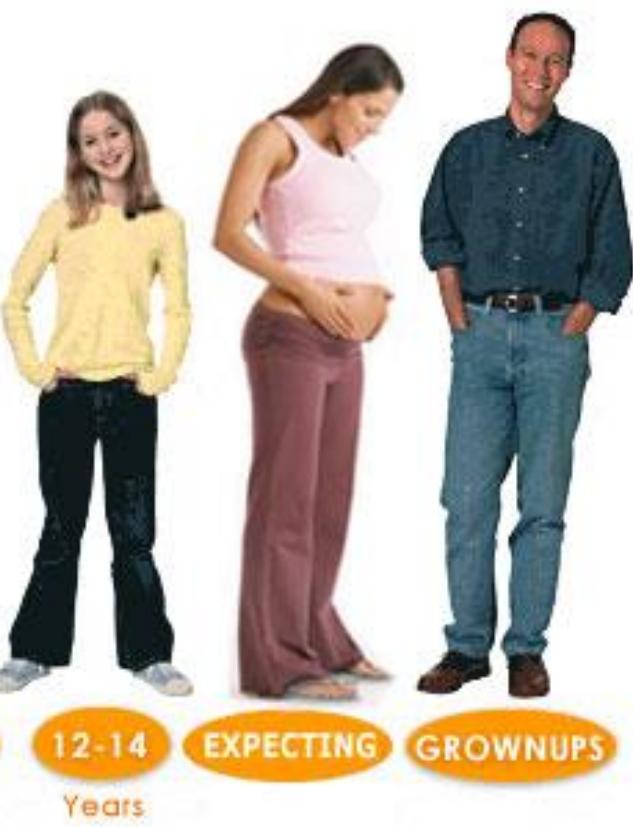
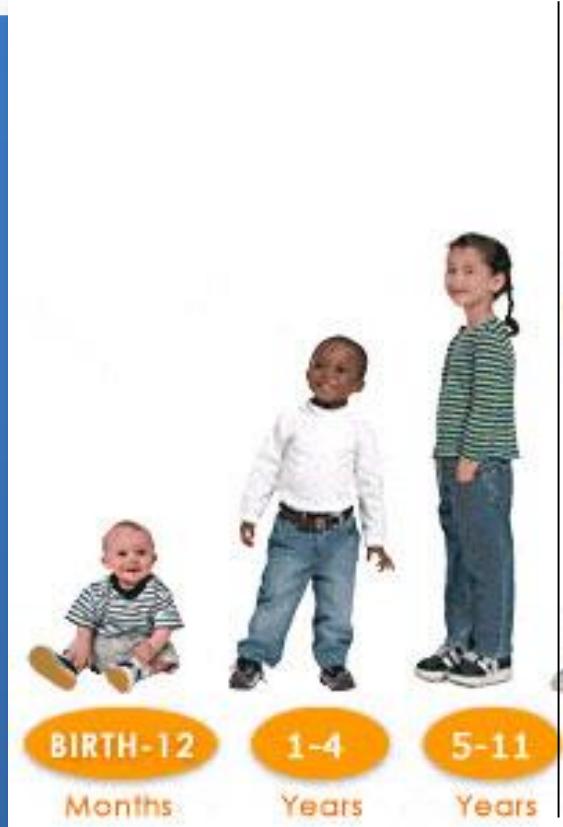
## → HUS (most frequent cause for acute renal failure)

**Table 7** Reported food- and waterborne *E. coli* outbreaks to EFSA in accordance with Directive 2003/99/EC in 2004–2009

STEC/VTEC/EHEC outbreaks	2009	2008	2007	2004–2006
Food-borne outbreaks	75	75	61	195
Waterborne outbreaks	5	4	4	5
Human cases in food-borne outbreaks	595	339	479 (includes only verified outbreaks)	2 345 (data missing from some outbreaks)
Human cases in waterborne outbreaks	12	22	62	26

- No specific therapy
- No specific prevention (e.g. vaccination)
- Complex diagnostics

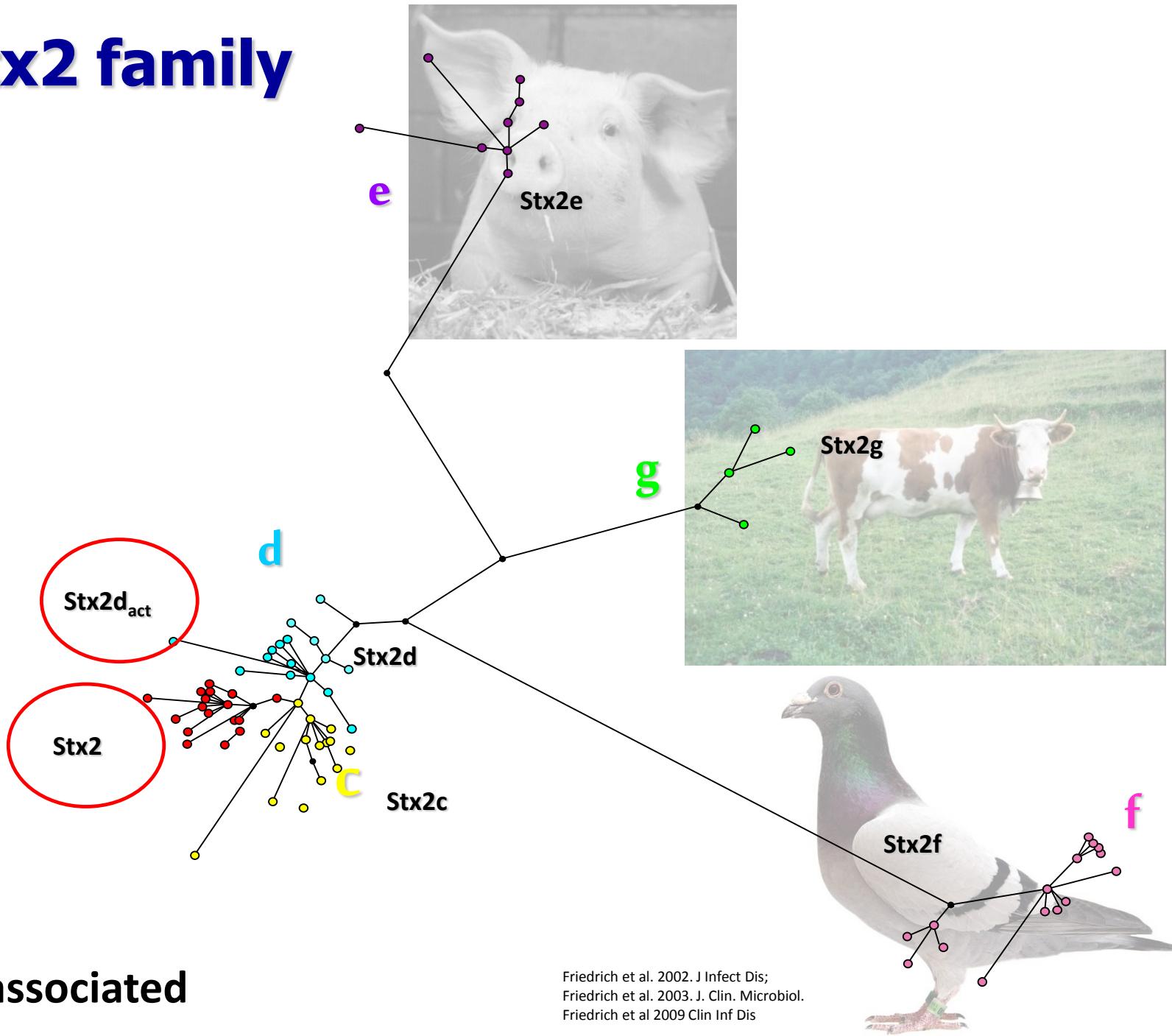
# (Zoonotic) STEC Infection sources



# HUS-association of molecular markers in EHEC diagnostics

PCR results	Percentage of pos. PCR on pat with HUS	<i>E. coli</i> isolates
<i>sfpA</i> <i>rfbO157</i> <i>eae</i> <i>stx</i>		
+	81%	SF O157:HNM
-	60%	NSF O157:H7
-	25%	non O157 EHEC
-	13%	STEC
-	0%	EPEC

# The Stx2 family



## EHEC: major concern of food born outbreak

- ✓ Common O-serogroups reported to cause foodborne illness are
- O157:H7/HNM**

O26,

O45

O103,

**O104:H4**

O111,

O121 and

O145

All eae +, except ...



- Association with raw fruits & vegetables?

# EHEC on fruits and veg

1997	<i>E. coli</i> O157:H7	Sprouted seeds (alfalfa)	Sivaplasingham et al. 2004
1997	<i>E. coli</i> O157:H7	Salad	Anon (2005a)
1998	<i>E. coli</i> O157:H7	Salad	Anon (2001a,b)
1998	<i>E. coli</i> O157:H7	Fruit salad	Anon (2001a,b)
1998	<i>E. coli</i> O157:H7	Coleslaw	Anon (2001a,b)
1998	<i>E. coli</i> O157:H7	Sprouted seeds (clover/alfalfa)	Taormina et al. 1999
1998	<i>E. coli</i> O157:H7	Unpasteurized apple juice	Anon (2001a,b)
1998	<i>E. coli</i> O157:H7	Parsley	Sivaplasingham et al. 2004
1999	<i>E. coli</i> O157:H7	Coriander (cilantro)	Campbell et al. 2001
1999	<i>E. coli</i> O157:H7	Unpasteurized apple juice	Anon (2001a,b)
2003	<i>E. coli</i> O157:H7	Cucumber	Duffell et al. (2003)
2003	<i>E. coli</i> O157:H7	Lettuce	Anon (2005a)
2005	<i>E. coli</i> O157:H7	Lettuce	Söderström et al. (2005)
2006	<i>E. coli</i> O157:H7	Spinach	CDC (2006b)
2006	<i>E. coli</i> O157:H7	Lettuce	CDC (2006c)



# EHEC O104:H4 outbreak In Germany

- >5922 cases
- >850 HUS
- 53 patients died

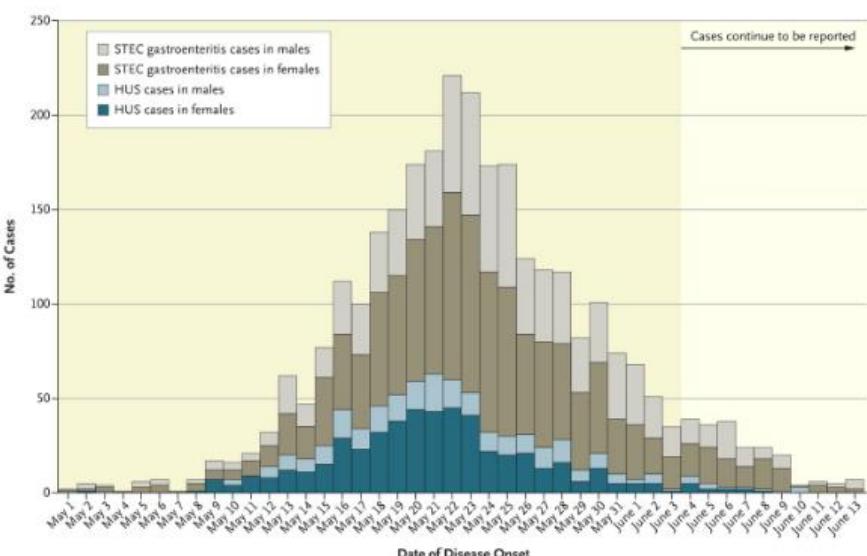
➤ E. coli O104:H4 HUSEC41/ESBL

ID: O104:H4

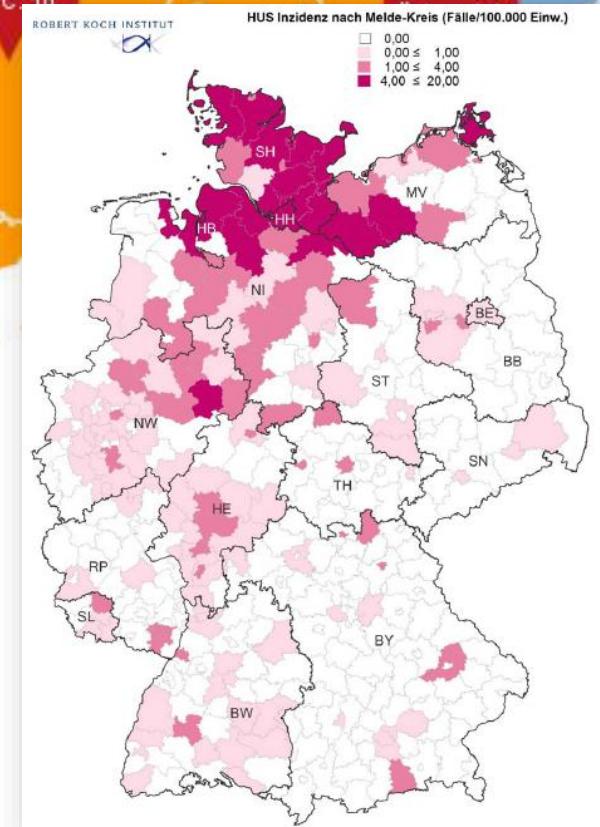
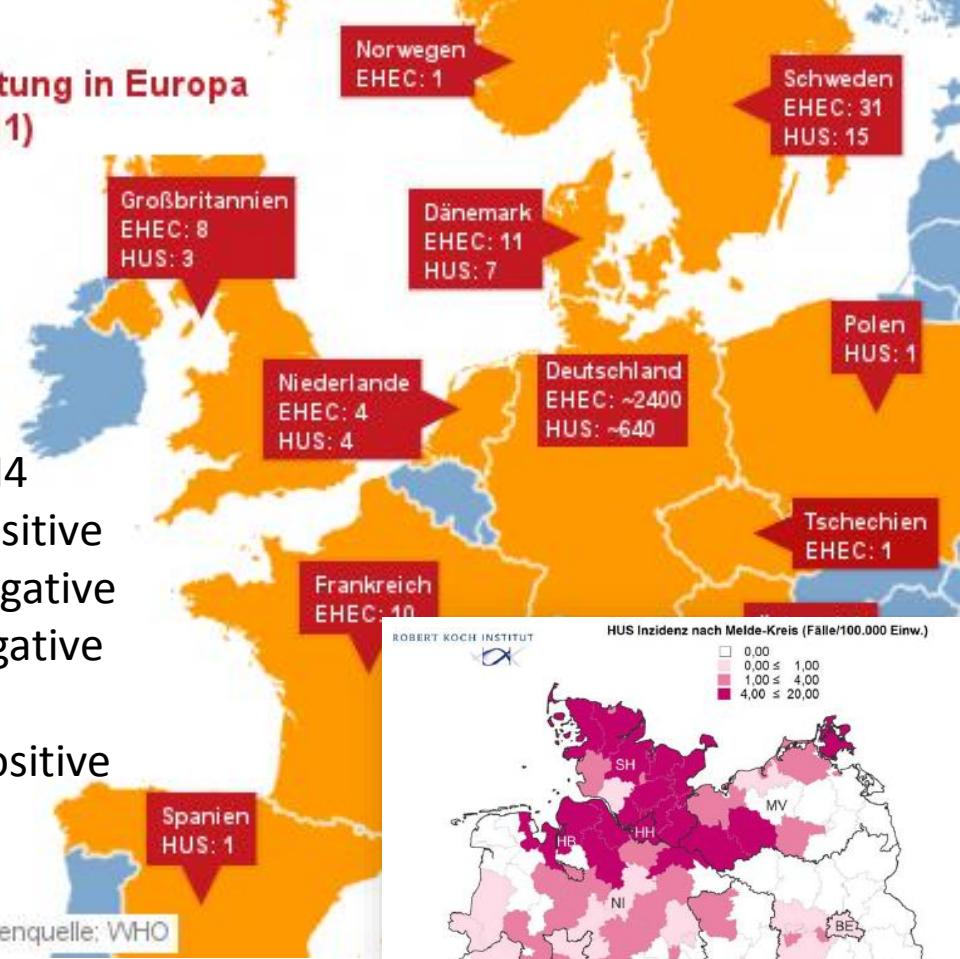
Stx2 positive Stx1 negative

Eae negative

ESBL positive

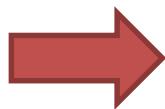


## EHEC-Vorbreitung in Europa (Stand: 5.6.2011)





# Mission uncompleted...



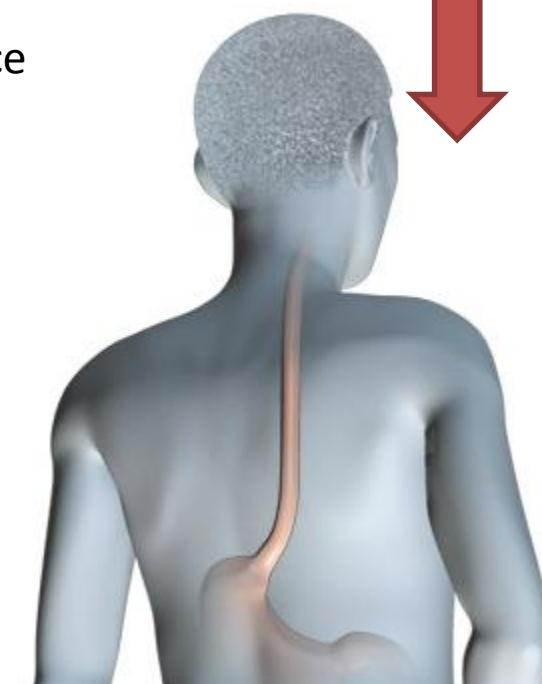
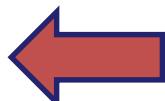
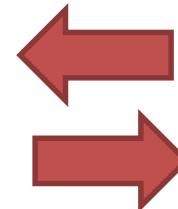
Primary source  
„fennugreek“ et al.



outbreak source



2nd source



# Why Sprouts ??

- Seeds and beans need warm and humid conditions to sprout and grow
- These sprouting conditions are also ideal for the growth of bacteria such as *E. coli*



# Tenacity of EHEC

Pathogen	Environment	Survival (day)	References
<i>Escherichia coli</i> O157:H7	Soil + animal manure	30	Nicholson et al. (2005)
<i>E. coli</i> O157:H7	Soil + animal manure	99	Nicholson et al. (2005)
<i>E. coli</i> O157:H7	Animal manure	60	Avery et al. (2005)
<i>E. coli</i> O157:H7	Slurries	60	Avery et al. (2005)
<i>E. coli</i> O157:H7	Abattoir waste	60	Avery et al. (2005)
<i>E. coli</i> O157:H7	Sewage sludge	60	Avery et al. (2005)
<i>E. coli</i> O157:H7	Nonaerated ovine manure	>365	Kudva et al. (1998)
<i>E. coli</i> O157:H7	Aerated ovine manure	120	Kudva et al. (1998)
<i>E. coli</i> O157:H7	Nonaerated slurry	600	Kudva et al. (1998)
<i>E. coli</i> O157:H7	Aerated slurry	30	Kudva et al. (1998)
<i>E. coli</i>	Slurry + dirty water	90	Nicholson et al. (2005)

# Prevention of Gastrointestinal Zoonotic diseases

## Keep cleaning



## Food safety



Raw Milk  
Harbors  
Dangerous  
Pathogens

## Animal Contact



## Surface structures involved in plant stomata and leaf colonization by Shiga-toxigenic *Escherichia coli* O157:H7

Zeus Saldaña<sup>1</sup>, Ethel Sánchez<sup>2</sup>, Juan Xicohtencatl-Cortés<sup>3</sup>, José Luis Puente<sup>4</sup> and Jorge A. Girón<sup>\*1</sup>

<sup>1</sup> Department of Molecular Genetics and Microbiology, Emerging Pathogens Institute, University of Florida, Gainesville, FL, USA

<sup>2</sup> Centro de Investigación en Estructuras Microscópicas, Universidad de Costa Rica, San José, Costa Rica

<sup>3</sup> Laboratorio de Bacteriología Intestinal, Hospital Infantil de México Federico Gómez, México D.F., México

<sup>4</sup> Departamento de Microbiología Molecular, Instituto de Biotecnología, Universidad Nacional Autónoma de México, Cuernavaca, Morelos, México

**Edited by:**

Adel M. Talyar, University of Wisconsin

Madison, USA

**Reviewed by:**

Mark A. Kotter, Washington State

University, USA

Jay C. D. Hinton, Trinity College Dublin,

Ireland

**\*Correspondence:**

Jorge A. Girón, Department of

Molecular Genetics and Microbiology,

Emerging Pathogens Institute,

University of Florida, 2055 Mowry

Road, PO Box 100099, Gainesville, FL,

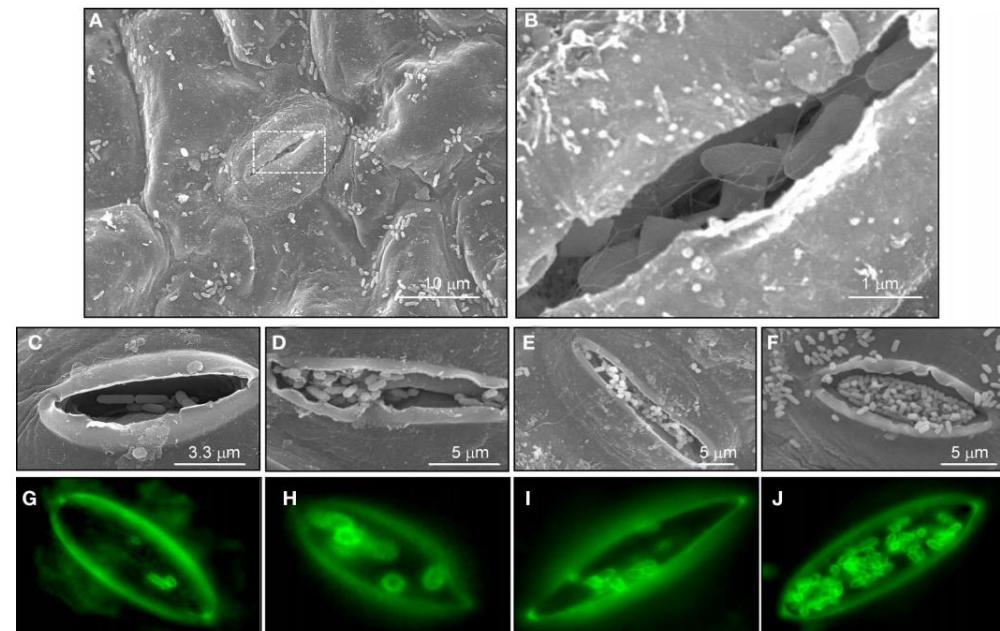
32610, USA.

e-mail: jagiron@ufl.edu

Shiga-toxigenic *Escherichia coli* (STEC) O157:H7 uses a myriad of surface adhesive appendages including pili, flagella, and the type 3 secretion system (T3SS) to adhere to and inflict damage to the human gut mucosa. Consumption of contaminated ground beef, milk, juices, water, or leafy greens has been associated with outbreaks of diarrheal disease in humans due to STEC. The aim of this study was to investigate which of the known STEC O157:H7 adherence factors mediate colonization of baby spinach leaves and where the bacteria reside within tainted leaves. We found that STEC O157:H7 colonizes baby spinach leaves through the coordinated production of curli, the *E. coli* common pilus, hemorrhagic coli type 4 pili, flagella, and T3SS. Electron microscopy analysis of tainted leaves revealed STEC bacteria in the internal cavity of the stomata, in intercellular spaces, and within vascular tissue (xylem and phloem), where the bacteria were protected from the bactericidal effect of gentamicin, sodium hypochlorite or ozonated water treatments. We confirmed that the T3S *escN* mutant showed a reduced number of bacteria within the stomata suggesting that T3S is required for the successful colonization of leaves. In agreement, non-pathogenic *E. coli* K-12 strain DH5α transformed with a plasmid carrying the locus of enterocyte effacement (LEE) pathogenicity island, harboring the T3SS and effector genes, internalized into stomata more efficiently than without the LEE. This study highlights a role for pili, flagella, and T3SS in the interaction of STEC with spinach leaves. Colonization of plant stomata and internal tissues may constitute a strategy by which STEC survives in a nutrient-rich microenvironment protected from external foes and may be a potential source for human infection.

**Keywords:** plant colonization, stomata, STEC, O157:H7, pathogenesis, spinach, T3SS, pilus

# In and not on plants



**FIGURE 4 | Evidence of STEC in stomata. (A)** Scanning electron micrograph showing bacteria on leaf epidermis at 6 h of infection. **(B)** High magnification of boxed area in (A) showing flagellate bacteria internalized in the stomata. **(C–F)** Micrographs (60X) of time-course EDL933 infection experiments between 3, 6, 12, and 24 h showing progressive association of bacteria with stomata. **(G–J)** Same experiment as before employing IFM and anti-O157 antibodies to stain bacteria (green).

(60X) of time-course EDL933 infection experiments between 3, 6, 12, and 24 h showing progressive association of bacteria with stomata. **(G–J)** Same experiment as before employing IFM and anti-O157 antibodies to stain bacteria (green).

# Stx-negative variants of EHEC O157

Outbreak no.	Year	Disease	No. of isolates		
			<i>stx</i> -positive	<i>stx</i> -negative	Total
1	2003	HUS	27	2	29
2	1995/96	HUS	12	1	13
3	2002	HUS	10	1	11
4	1993	D	1	6	7
5	1994	D	0	4	4
6	1994	D	0	3	3

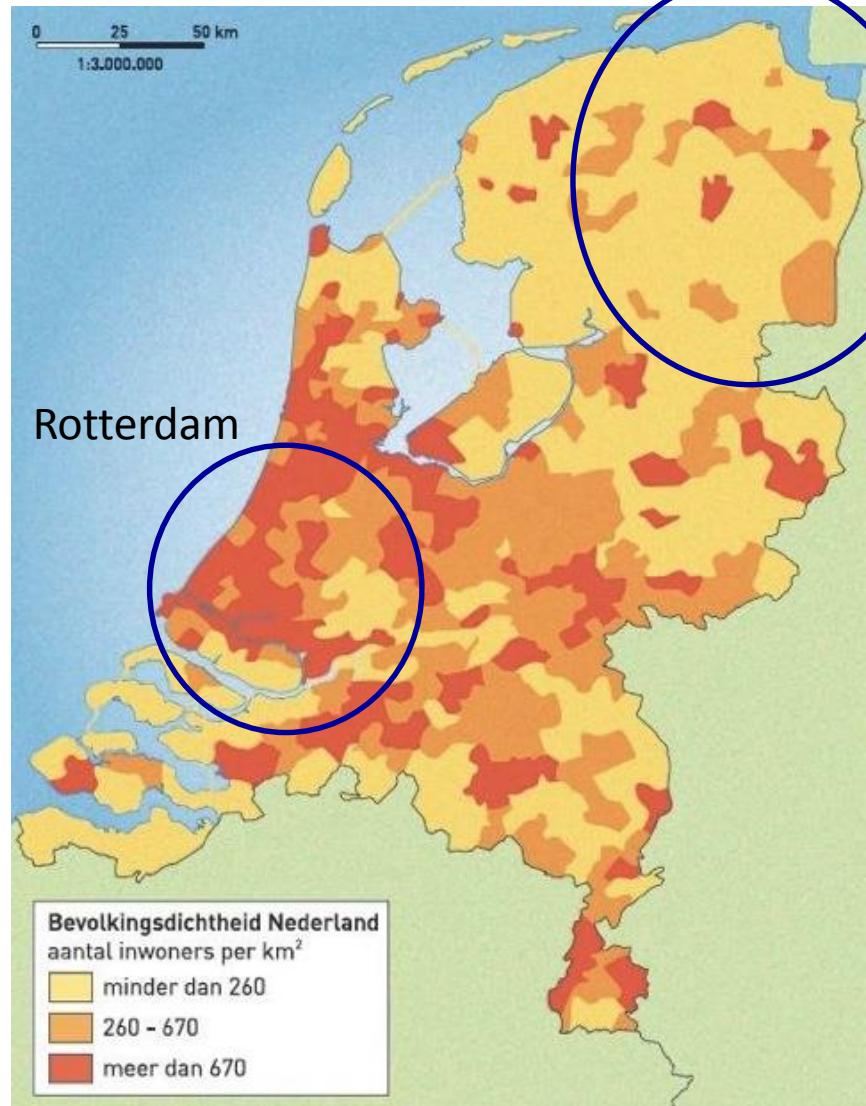
Groningen

# STEC-ID-net

- Running multicenter study in NL
- 25.000 patients from GP with Diarrhea/BD/HUS
- Identify prevalence of STEC-EHEC/HUSEC
- Capacity building of STEC-diagnostics
- Create an expert network
- Microbiological risk assessment for Public Health
- Create Dutch HUSEC collection



John Rossen, Mirjam Kooistra, CERTE  
UMCG



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Rijksinstituut voor Volksgezondheid  
en Milieu  
Ministerie van Volksgezondheid,  
Wetgeving en Sport



# STEC-ID net: direct PCR-detection

April 2013 t/m Augustus 2013

Included patients: n= 9582 (Groningen n=6358; Rotterdam n=3224)

Target(s)	Groningen		Rotterdam	
	n	%	n	%
<i>stx1</i>	23	0.4	15	0.5
<i>stx1,eae</i>	30	0.5	22	0.7
<i>stx2</i>	23	0.4	12	0.4
<i>stx2,eae</i>	30	0.5	16	0.5
<i>stx1,stx2</i>	6	0.1	4	0.1
<i>stx1,stx2,eae</i>	13	0.2	7	0.2
<i>eae</i>	638	10.0	410	12.7
<b>TOTAAL</b>	<b>763</b>	<b>12.1</b>	<b>468</b>	<b>15.1</b>
STEC-prevalence		0,8%		0,4%
EHEC-prevalence		1,1%		1,4

Shiga toxin 2-encoding bacteriophages in human fecal samples from healthy individuals.

Alexandre Martinez-Castillo, Pablo Quirós, Ferran Navarro, Elisenda Miró, Maite Muniesa

Department of Microbiology, University of Barcelona, Diagonal 643, Annex, Floor 0, 08028 Barcelona, Spain.

[Applied and environmental microbiology](#) (Impact Factor: 3.69). 06/2013;  
DOI:10.1128/AEM.01158-13

Source: PubMed

**ABSTRACT** Shiga toxin-converting bacteriophages (Stx phages) carry the stx gene and convert non-pathogenic bacterial strains into Shiga toxin-producing bacteria. Previous studies have shown that high densities of free and infectious Stx phages are found in environments polluted with feces and also in food samples. Taken together, these two

## Direct-PCR

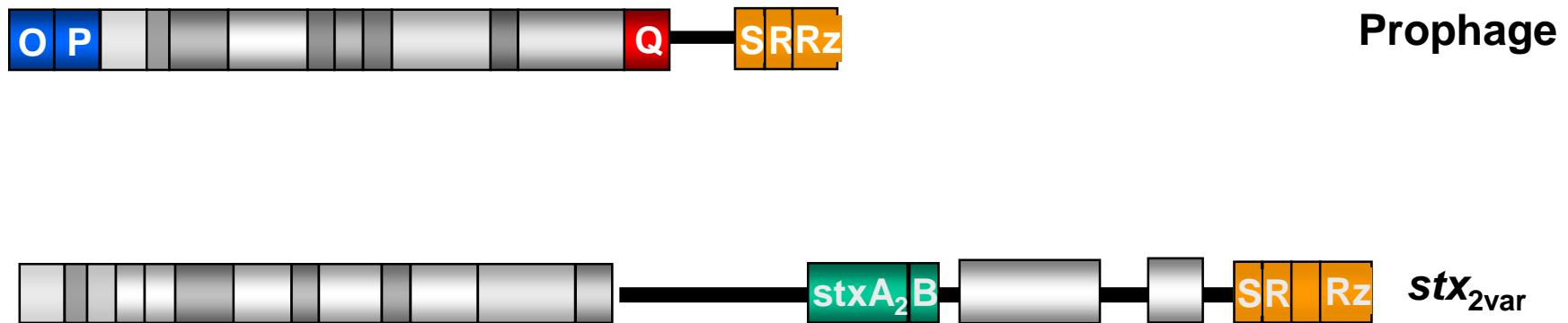
<i>stx1/stx2/eae</i>	763	100%	486	100%
after Pre-enrichment:	718	94.1%	457	94.0%

1249 100%

1175 94.1%

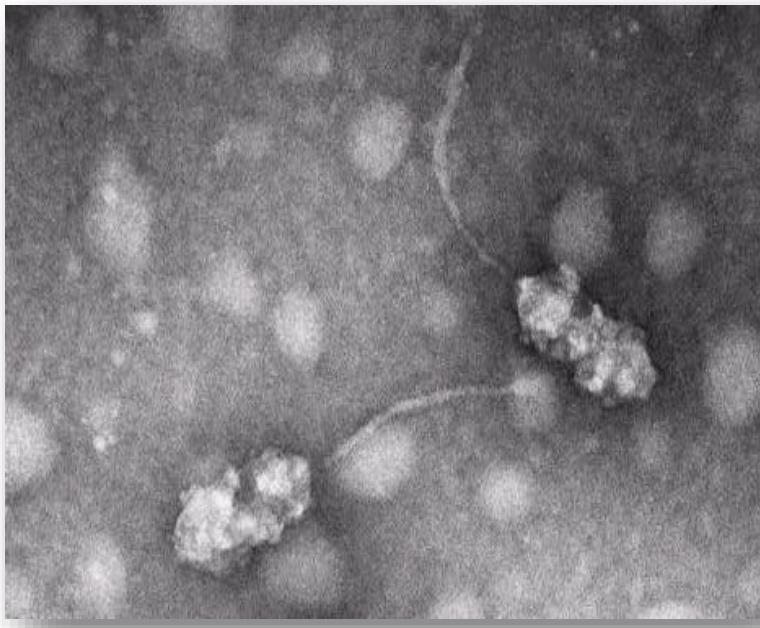
?

# Structure analysis of *stx*-Gene



Antibiotics in pre-HUS-phase increase risk for HUS  
2011 outbreak: Antibiotics after HUS improved clinical outcome

# *stx*-genes: common in nature



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*Appl Environ Microbiol*. 2009 Mar;75(6):1764-8. Epub 2009 Jan 23.  
**Phage-mediated Shiga toxin 2 gene transfer in food and water.**  
Imamovic L, Jofre J, Schmidt H, Serra-Moreno R, Muniesa M.  
Department of Microbiology, Faculty of Biology, University of Barcelona, Diagonal 645, E-08028 Barcelona, Spain.

**Abstract**  
Shiga toxin (*stx*) transduction in various food matrices has been evaluated with lysogens of *Stx* phages. *stx* transduction events were observed for many phages under appropriate conditions. Transduction did not occur at low pH and low temperatures. A total of 10(3) to 10(4) CFU ml(-1) was the minimal amount of donor and recipient strains necessary to generate transductants.

PMID: 19168651 [PubMed - indexed for MEDLINE] PMCID: PMC2655461 Free PMC Article

 Maite Muniesa 35.90  
University of Barcelona

## Article

### Persistence of infectious *Stx* bacteriophages after disinfection treatments.

Anna Allué-Guardia, Alexandre Martínez-Castillo, Maite Muniesa

*Applied and environmental microbiology* (Impact Factor: 3.69). 01/2014;  
DOI:10.1128/AEM.04006-13  
Source: PubMed

**ABSTRACT** In Shiga toxin-producing *Escherichia coli* (STEC), induction of *Stx* phages causes the release of free phages that can later be found in the environment. The ability of *Stx* phages to survive different inactivation conditions will determine their prevalence in the environment, the risk of *stx* transduction and the generation of new STEC. We evaluated the infectivity and genomes of two *Stx* phages ( $\phi$ 534 and  $\phi$ 557) in different conditions. Infectious *Stx* phages were stable at 4, 22 and 37°C and at pH 7 and 9 after one month of storage, but were completely inactivated at pH 3. Infective *Stx* phages decreased moderately when treated with UV (2.2 log<sub>10</sub> reduction for an estimated UV dose of 178.2 mJ/cm<sup>2</sup>) or after treatment at 60 and 68°C for 60 min (2.2 and 2.5 log<sub>10</sub> reduction respectively) and were highly inactivated (3 log<sub>10</sub>) by 10 ppm of chlorine in one min. Assays in a mesocosm showed lower inactivation of all microorganisms in winter than in summer. The number of *Stx* phage genomes did not decrease significantly in most cases, and STEC inactivation was higher than phage inactivation in all conditions. Moreover, *Stx* phages retained the ability to lysogenize *E. coli* after some of the treatments [less]



# Are we prepared?

- STEC of important zoonotic impact also via vegetables
- Stx-genes not enough to determine STEC/EHEC
- Risk analysis of STEC/EHEC needed for public health, infection control and food safety
- Need for national research network on STEC/EHEC for outbreak preparation

INVITATION

# ONE HEALTH SYMPOSIUM FOCUS ON STEC - HUSEC



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