

**Prevalence of the 7 major serogroups of enterohemorrhagic *Escherichia coli* (EHEC) in fresh minced beef in France:
A novel real-time PCR strategy for their early detection in food**

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Rome, 8th -9th November 2012**



VetAgro Sup

1- Lyon University, VetAgro Sup, LMAP, French National Reference Laboratory for *E. coli*, FRANCE

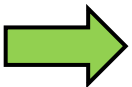


2- French Food Safety Agency (ANSES)-Laboratoire de Sécurité des aliments, FRANCE

Introduction

Precise virulent profiles of EHEC strains are still unknown.

2 approaches to defining EHEC strains outside a clinical context in humans:



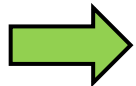
Epidemiological criteria

Identify serotypes more frequent and more associated with outbreaks and severe disease (Karmali *et al.*, 2003)

O157:H7, O26:H11, O103:H2, O111:H8, O145:H28, O121:H19, O45:H2
and non motile derivates

Top 5

Top 7



Molecular criteria

On the basis of molecular markers associated with pathogenicity

Association of *eae* variants with specific EHEC serotypes

Association of genomic O islands with major EHEC strains (notably O122)



Bugarel *et al.*, 2010, *Appl. Environ. Microbiol.* (76) 1:203-11; Coombes *et al.*, 2008, *Appl. Env. Microbiol.* (74) 2153-60; Karmali *et al.*, 2003, *J. Clin. Microbiol.* (41) 4930-40; Immamovic *et al.*, 2010, *Infect. Immun.* (78) 11: 4697-704.

Identification of the major typical EHEC strains

Major EHEC	Serogroups maker	stx genes	LEE marker	eae variants	O1 122 marker
EHEC O157:H7/H-	<i>rfbE</i> -O157	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	γ	<i>nleB</i>
EHEC O26:H11/H-	<i>wzx</i> -O26	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	β	<i>nleB</i>
EHEC O103:H2/H-	<i>wzx</i> -O103	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	ϵ	<i>nleB</i>
EHEC O111:H8/H-	<i>wbdI</i> -O111	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	θ	<i>nleB</i>
EHEC O145:H28/H-	<i>ihp1</i> -O145	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	γ	<i>nleB</i>
EHEC O121:H19/H-	<i>wzx</i> -O121	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	ϵ	<i>nleB</i>
EHEC O45:H2/H-	<i>wzx</i> -O45	<i>stx1</i> or <i>stx2</i>	<i>eae</i>	ϵ or β	<i>nleB</i>

Top 5

Top 7

ISO TS 13136 and MLG 5B

?

 **New tools for EHEC screening in meat?**

Bugarel *et al.*, 2010, Int. J. Food. Microbiol. (142) 318- 29 ; Madic *et al.*, 2011, Appl. Environ. Microbiol. (77) 6:2035-41; Nielsen *et al.*, 2003, J. Clin. Microbiol. (41) 7 :2884-93; Perelle *et al.*, 2004, Mol. Cell Probes (18) 185-92

Objectives



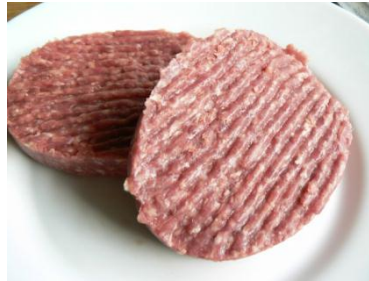
with permission of Caprioli A.



- ➔ **Evaluate the prevalence of the top 7 EHEC in fresh minced beef samples collected in France**
- ➔ **Propose a novel detection strategy for their early detection in minced meat.**

➔ Food samples

2476 fresh minced beef samples



collected in supermarkets of 92 French departments



from March to December 2010

Materials and methods

Food PCR detection

Test portion 25g

Enrichment
in 225 ml BPW 37C 18/24h

Test portion (1ml)
DNA extraction purification

ISO TS 13136
MLG 5B

RT-PCR detection
of *stx* and *eae* genes

GD1

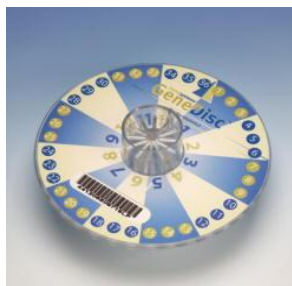
stx/eae +

RT-PCR detection
of top 7 associated genes

GD2

stx/eae/top7 +

Isolation



New
approach

RT-PCR detection
of *stx* and *eae* genes

New
GD1

stx/eae +

RT-PCR detection
of *nleB* genes

stx/eae/nleB +

RT-PCR detection
of top 7 associated genes

New
GD2

stx/eae/nleB/top7 +

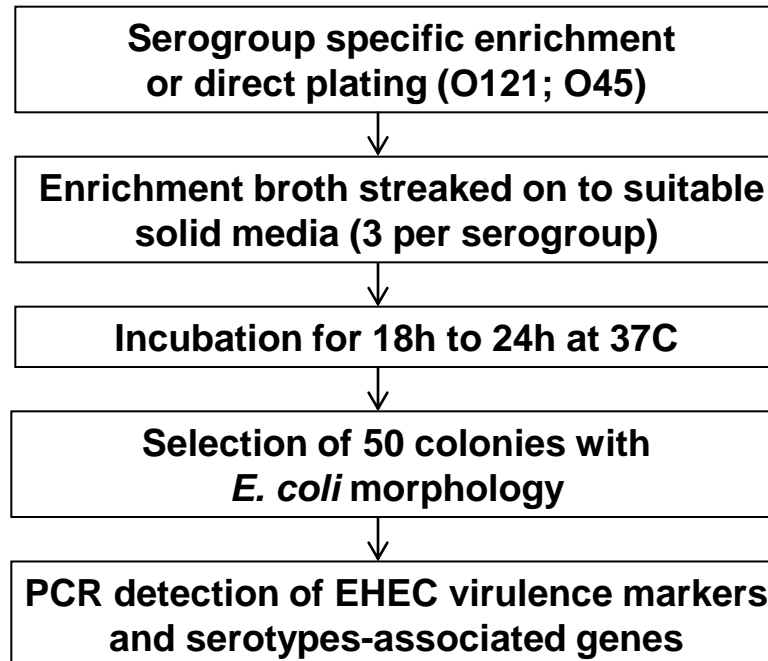
RT-PCR detection
of top 7 associated *eae* variants

stx/eae/nleB/top7
/eae variants +

Isolation

➔ Isolation and characterization procedure

ISO TS 13136



Results and discussion

→ **Top 5 prevalence: 6/2476 (0,2 % CI₉₅ [0,09%- 0,5%])**



Results and discussion

➔ **Top 5 prevalence: 6/2476 (0,2 % CI₉₅ [0,09%- 0,5%])**

- Low prevalence
- In accordance with previous results observed in France



Year	Type of minced beef	Top 5 prevalence
2007	Frozen (production)	0,3% (11/3605) CI ₉₅ [0,2 – 0,5%]
2008	Frozen trims (production)	0,4% (15/4000) CI ₉₅ [0,2 – 0,6%]
2009	Fresh (at retail)	0,1% (2/1527) CI ₉₅ [0,04 – 0,5%]

➔ **Top 7 prevalence: 7/2476 (0,3 % CI₉₅ [0,1%- 0,6%])**

- First data in French minced beef
- Low prevalence (comparison with other areas? USA: 0,2% CI₉₅ [0,05 – 0,3%] (6/4133) (Bosilevac and Koohmaraie, 2011))

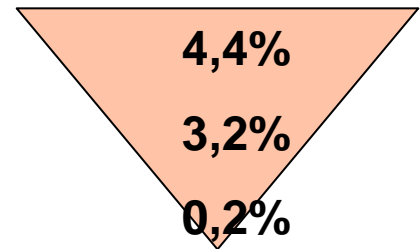
→ Top 5 screening

N = 2476 samples (25g)



Markers/strains detected	Nb of positive samples
<i>stx+</i> <i>eae+</i>	109
<i>stx+</i> <i>eae+</i> top 5+	79
EHEC strains isolated (top 5)	6

ISO TS 13136



Results and discussion



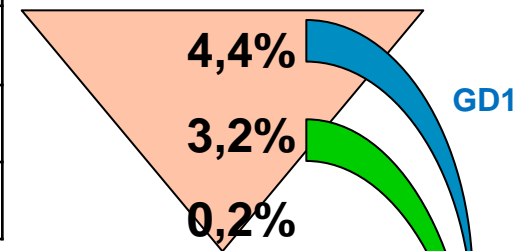
➔ Top 5 screening

N = 2476 samples (25g)

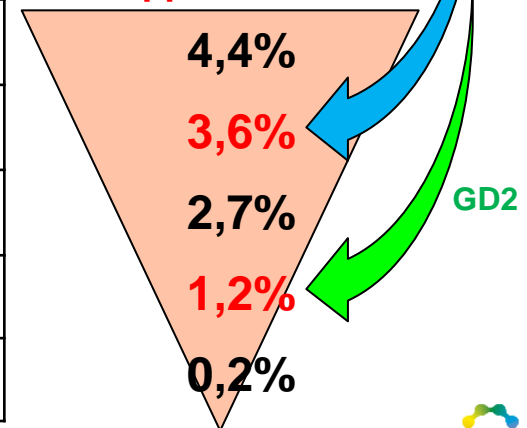
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Markers/strains detected	Nb of positive samples
<i>stx+</i> <i>eae+</i>	109
<i>stx+</i> <i>eae+</i> <i>nleB+</i>	88
<i>stx+</i> <i>eae+</i> <i>nleB+</i> top 5+	67
<i>stx+</i> <i>eae+</i> <i>nleB+</i> top 5/ <i>eae</i> assoc.+	29
EHEC strains isolated (top 5)	6

ISO TS 13136



New approach



Results and discussion



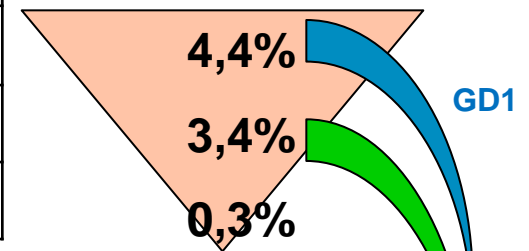
➔ Top 7 screening

N = 2476 samples (25g)

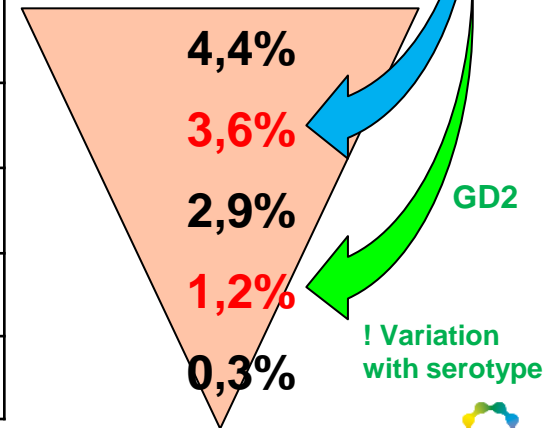
Markers/strains detected	Nb of positive samples
<i>stx+</i> <i>eae+</i>	109
<i>stx+</i> <i>eae+</i> top 7+	84
EHEC strains isolated (top 7)	7

Markers/strains detected	Nb of positive samples
<i>stx+</i> <i>eae+</i>	109
<i>stx+</i> <i>eae+</i> <i>nleB+</i>	88
<i>stx+</i> <i>eae+</i> <i>nleB+</i> top 7+	71
<i>stx+</i> <i>eae+</i> <i>nleB+</i> top 7/ <i>eae</i> assoc.+	29
EHEC strains isolated (top 7)	7

ISO TS 13136
MLG 5B



New approach





Characteristics of the 7 EHEC strains isolated

Serotype	<i>stx</i> genes	<i>eae</i> variants	O1 122 marker	Nb of strains isolated
EHEC O157:[H7]	<i>stx2</i>	<i>eae</i> γ	<i>nleB</i>	<u>1</u>
EHEC O26:[H11]	<i>stx1</i>	<i>eae</i> β	<i>nleB</i>	4
EHEC O103:H2/H ⁻	<i>stx1</i> or <i>stx2</i>	<i>eae</i> ϵ	<i>nleB</i>	0
EHEC O111:H8/H ⁻	<i>stx1</i> or <i>stx2</i>	<i>eae</i> θ	<i>nleB</i>	0
EHEC O145:[H28]	<i>stx1</i>	<i>eae</i> γ	<i>nleB</i>	1
EHEC O121:H19/H ⁻	<i>stx1</i> or <i>stx2</i>	<i>eae</i> ϵ	<i>nleB</i>	0
EHEC O45:[H2]	<i>stx1</i>	<i>eae</i> ϵ	<i>nleB</i>	1

Top 5

Top 7

No EHEC O103, O111 nor O121

Only 1 EHEC O157:H7 and a majority of EHEC O26:H11 (4/7)





9 particular AEEC strains isolated

Isolated in samples in which *stx* genes were detected by PCR
Resemble the major EHEC strains (top 5 / top 7)

Serotype	<i>eae</i> variants	O1 122 marker	Nb of strains isolated
AEEC O157:nonH7	<i>no eae</i> γ	<i>no nleB</i>	2
AEEC O26:[H11]	<i>eae</i> β	<i>nleB</i>	7
AEEC O103:[H2]	<i>eae</i> ϵ	<i>nleB</i>	2
AEEC O103:nonH2	<i>no eae</i> ϵ	<i>no nleB</i>	2
AEEC O111:H8/H ⁻	<i>eae</i> θ	<i>nleB</i>	0
AEEC O145:H28/H ⁻	<i>eae</i> γ	<i>nleB</i>	0
AEEC O121:H19/H ⁻	<i>eae</i> ϵ	<i>nleB</i>	0
AEEC O45:H2/H ⁻	<i>eae</i> ϵ	<i>nleB</i>	0

Public health significance of these particular strains?

➔ Prevalence of major EHEC in fresh minced meat in France

The 5 and the 7 major EHEC serogroups (top 5 and top 7) were detected at a low prevalence (0,2% and 0,3% respectively)

O26 was the main serogroup isolated in meat
None O111 nor O121 strains were isolated



➔ A novel real-time PCR strategy for EHEC detection in food

using O122 marker and association of serotype with specific eae variants

A reliable strategy (specific and sensitive)

Based upon complementary tools to those described in the ISO TS 13136

A cost effective screening (reduces the number of PCR suspicions without increasing time analysis)



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anses

French agency for food, environmental
and occupational health safety



Thanks !

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