PREVALENCE OF VEROTOXIGENIC *Escherichia coli* SEROGROUPS AND VIRULENCE GENES OF RELEVANCE IN PUBLIC HEALTH IN FATTENING CALVES IN SPAIN

Iratxe Perez-Cobo¹, Maria José Ruano Ramos¹, Montserrat Agüero García¹, José Luis Saez Llorente², Soledad Collado Cortés²

¹ Central Veterinary Laboratory (LCV), Ministry of Agriculture Fish and Food (MAPA), Algete (Madrid), Spain

² Subdirectorate General for Animal Health and Hygiene and Traceability, Ministry of Agriculture Fish and Food (MAPA), Spain



Iratxe Perez-Cobo: iperezco@mapa.es

Head of Bacteriology Department -2

Central Veterinary Laboratory-Animal health (LCV)



LABORATORY-ANIMAL HEALTH (LCV)



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INTEGRATED MANAGEMENT SYSTEM





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ORGANIGRAMA LCV

VIROLOGÍA 3







ACTIVITIES OF BACTERIOLOGY DEPARMENT 2

Act as NLR of:

✓ Campylobacteriosis

✓ **VTEC**

- Monitoring and reporting of antimicrobial resistance in zoonotic and commensal bacteria and repealing Implementing Decision 2020/ 1729 /EU (isolation)
 - ✓ Campylobacter coli, Campylobacter jejuni
 - ✓ Indicator commensal *Escherichia coli* (E. coli)
 - \checkmark *E. coli* producing the following enzymes:
 - ✓ Extended Spectrum β -Lactamases (ESBL)
 - ✓ AmpC β -Lactamases (AmpC);
 - ✓ Carbapenemases (CP)
- \checkmark Leptospirosis
- ✓ Botulism
- ✓ Tularemia
- ✓ Contagious Equine Metritis
- Bacteriological honeybees diseases (American Foulrood, European Foulbrood)





INTRODUCTION

Verotoxigenic *Escherichia coli* (VTEC) is a foodborne pathogen of relevance in Public Health in European Union (EU), being in **2020** the fourth most frequent bacterial agent detected in foodborne outbreaks with an increase between 2015 to 2019.

 hospitalisation for 35.8% of all confirmed VTEC cases in the EU in 2020 (EFSA Journal 2021;19(12):6971)

The most common serogroups among **HUS cases** were:

%	O 26	O157	080	O145	ONT
2019	38.7	23.0	9.0	9.8	4.7
2020	41.8	11.9	13.2	8	24.1

- Directive 2003/99/EC established that verotoxigenic Escherichia coli (VTEC) is a zoonotic agent to be included in monitoring (list A). In Spain it was transposed by RD 1940/2004 on the surveillance of zoonoses and zoonotic agents
- In the EU, animal monitoring is not widely implemented or carried out with a harmonized sampling strategy.







 The objetive of this surveillance was determine VTEC prevalence in Spain in calves less than one year of age and strains characterization.

 Assist the investigation of foodborne outbreaks in humans, providing support to the competent authorities under a One Health approach.





SAMPLING

- ✓ Based on the technical specifications for the monitoring and reporting of VTEC on animals and food drafted by EFSA (EFSA Journal 2009; 7(11):1366).
- A nationwide monitoring in the most representative national production slaughterhouses throughout the Spanish geography (Sampling of AMR surveillance was taken advantage of)
- $\checkmark\,$ For years 2019 and 2021.
- ✓ Targeted population: calves under one year of age
- Epidemiological unit: herd. One sample from each herd avoiding repetitions
- A total of 498 hide swabs samples (breast) were taken throughout the year avoiding seasonal bias.





EFSA Journal 2009; 7(11):1366

SCIENTIFIC REPORT OF EFSA

Technical specifications for the monitoring and reporting of verotoxigenic *Escherichia coli* (VTEC) on animals and food (VTEC surveys on animals and food)¹

European Food Safety Authority^{2, 3}

European Food Safety Authority (EFSA), Parma, Italy





LABORATORY ANALYTICAL METHODS



2019: INTERNAL METHOD ACCREDITED BASED ON ISO 13136:2012 FOR THE PRESUMPTIVE DETECTION, ISOLATION AND IDENTIFICATION.

STEP 1: Microbiological enrichment of the sample

1 sponge of hide calve: 90 ml BPW 18-24 h (37°C ±1°C)

STEP 2: Presumptive detection in BPW: real-time PCR

- vtx1, vtx2 genes
- \circ eae gene
- if positive to eae: determine 6 of the more prevalent serogroups of VTEC in human health (Top 5 (0157, 026,0111, 0103,0145) + 0104)

STEP 3: Isolation, confirmation and characterization of strains

- Inoculation in specific solid media for *E. coli: TBX, MCA, CT-SMAC, CT-RMAC*
- Pick up to **50 colonies** with *E. coli* morphology
- Inoculate, dot or streak each colony in TSA
- Test of pools of 10 colonies by real-time PCR for the detection of virulence genes or markers associated with STEC (stx1/stx2).
- Subculture of isolated colonies for subsequent confirmation and characterization by real-time PCR





2019 SURVEILLANCE RESULTS

• STEP1: Microbiological enrichment of the sample

236 samples

• STEP2: Presumptive detection in BPW: real-time PCR

Results in 95,3% positive samples.

- 98,7% from these were eae positive.
 - 97,8% of them were positive at least to one or more of the 6 investigated serogroups "Top 5" + O104)
- STEP3: Isolation, confirmation and characterization of strains
 - prevalence of 21,2% (79 isolates)
 - 32,9 % VTEC (vtx1 and/or vtx2, eae +, O-T (O157, O111, O26, O103)
 - 19,0 %: VTEC (vtx1 and/or vtx2, eae +, O-NT)
 - 48,1 % VTEC (vtx1 and/or vtx2, eae -)

 Serogroups
 2019

 O157 VTEC
 19.0%

 O26 VTEC
 7.6%

 O103 VTEC
 5.1%

 O145 VTEC
 0%

 O111 VTEC
 1.3%

 O104 VTEC
 0%

finantial and laboratory work cost

2019	vtx1,eae	vtx2,eae	vtx1,vtx2,eae	vtx1	vtx2	vtx1,vtx2
0157	1,3%	6,3%	11,4%			
O26	6,3%		1,3%			
O103	5,1%					
0145						
0104						
0111	1,3%					17
O-NT	5.1%	7.6%	6.3%	11.4%	27.8%	8.9%





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Nº de animales en cebaderos, julio 2018

2019 Municipalties analyzed

Resultado

Detectado No detectado





2021: INTERNAL METHOD WITH REVIEW ACCREDITED FOR THE ISOLATION AND IDENTIFICATION OF SHIGATOXIN-PRODUCING *Escherichia coli* (VTEC) AND DETERMINATION OF SEROGROUPS 0157, 0111, 026, 0103, 0104 AND 0145.

STEP 1: Microbiological enrichment of the sample

1 sponge of hide calve: 90 ml BPW 18-24 h (**41,5°C** ±1°C)

STEP 2: Isolation of **100%** of the samples, confirmation and characterization.





2021 SURVEILLANCE RESULTS

• STEP 1: Microbiological enrichment of the sample in BWP

262 samples

- STEP 2: Isolation, confirmation and characterization
 - prevalence of 36,6 % (112 isolates)
 - 27,7% VTEC (vtx1 and/or vtx2, eae +, O-T (O157, O26, O103,O145)
 - 14,3 %: VTEC (vtx1 and/or vtx2, eae +, O-NT)
 - 58,0 % VTEC (vtx1 and/or vtx2, eae -)

Serogroups	2021
O157 VTEC	15.2 %
O26 VTEC	8.9 %
O103 VTEC	1.8 %
O145 VTEC	1.8 %
O111 VTEC	0 %
O104 VTEC	0%

2021	vtx1,eae	vtx2,eae	vtx1,vtx2,ea	vtx1	vtx2	vtx1,vtx2
0157	0,9%	10,7%	3,6%			
O26	5,4%	0,9%	2,7%			
0103		1,8%				
0145		1,8%		0,9%	0,9%	
0104						
0111						
O-NT	7,1%	3,6%	3,6%	17,9%	35,7%	2,7%
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EFSA Journal

CONCLUSIONS 1

- ✓ Wide effort in sampling and laboratory analysis: a total of 498 samples (cattle sampling units were analysed in 2019 and 2021)
- \checkmark The detection increase from 2019 to 2021 from **21.2%** to **36.6%**.
 - May be due to the optimisation of the temperature of enrichment in BPW (changed from 37°C to 41,5°C).

SCIENTIFIC REPORT

- EU 2019: 17,1% positive herds (9 MS reported data, 1493 cattle sampling units)
- EU 2020: 5,2% positive herds (3 MS reported data, 678 cattle sampling units)
- ✓ Sampling in primary production: it could be used at EU level taking advance of the sampling of AMR surveillance every two years (calves).





CONCLUSIONS 2

- The more prevalent serogroups in calves (<1 year old) in Spain were 0157, 026, 0103 and 0145, representing the 17.1%, 8.3%, 3.4% and 0.9% of the isolates respectively. Not detection for 0104.
 - $\checkmark~$ In line with human HUS cases
- ✓ **Future**: subtyping
- ✓ The periodic surveillance in targeted animal populations could assist the investigation of foodborne outbreaks in humans, providing support to the competent authorities under a One Health approach.







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Without all of them...

it would not be possible.