

Some statistics on E. coli O157

8 O157 eae + vtx1a

NO HUS

85 O157 eae + vtx1a + vtx2c

NO HUS

7 O157 eae + vtx1a + vtx2a

2/6 (33%) with HUS

37 O157 eae + vtx2a

11/37 (30%) with HUS

45 O157 eae + vtx2a + vtx2c

11/45 (24%) with HUS

31 O157 eae + vtx2c

1/31 (3%) with HUS

HUSEC: vtx2a and eae

Current outbreak:

0157 eae + vtx1a + vtx2a

7/12 (58%) with HUS





4th external quality assurance (EQA-4) scheme & PT10 EU-RL for typing of VTEC in 2012-2013

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ECDC contracts for 4 years

ECDC framework contracts with the overall purpose of

- Performing surveillance of foodborne infections based on isolate characteristics (subtyping) → outbreak detection
- Ensure that comparable methods are used in all countries
- Pathogens covered:
 - Salmonella
 - Listeria
 - Vero cytotoxin-producing E. coli

CONTRACT 1: EQA PROGRAMMES

MICROBIOLOGICAL CHARACTERISATION SERVICES



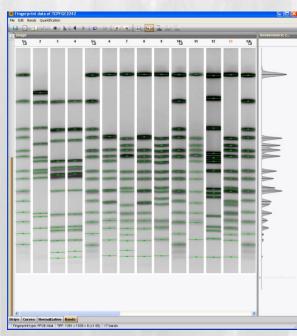
- >30 laboratories; 10-60 strains in each EQA
- Methods:
 - Pulsed-field gel electroforesis (PFGE) for all 3 pathogens
 - MLVA for Salmonella Typhimurium
 - Phage typing of Salmonella (subcontractor: DTU-Food)
 - E. coli serotyping and virulence characterisation
 - Listeria serotypning molecular and conventional (subcontractor: ANSES)
- Trouble-shooting regarding PFGE and MLVA
- The MLVA method: maintain and extend the reference material
- Perform molecular typing for other countries, including data analysis and storage of strains
- Protocols, certificates, individual reports, report, scientific paper
- Since start Sept 1: protocols, invitation letter, selection of strains...



CONTRACT 2: EXPERT MANAGEMENT (curator)

- **Expert management of molecular typing** data (curator)
- **BioNumerics databases with molecular** typing data maintained at ECDC, but as lab-experts we shall:

 - Data validation (quality, normalisation, band assignment)
 - Assign nomenclature (reference types)
 - **Analyse for clusters**
 - Give feedback to laboratories
 - Give expert advice to ECDC regarding interpretation of data, clusters/outbreaks, SOPs, etc
- **Expected amount of typing data for** curation
 - Salmonella: 5000 isolates per year
 - Listeria: 2000 isolates first year; 500 per following years
 - VTEC: 500 isolates per year
- September-October: kick-off meeting pilot project, our input to SOPs...





4th external quality assurance (EQA-4) scheme & PT10 EU-RL for typing of VTEC in 2012-2013

Strains and methods

Strains selected for the EQA-4 programme is based on five criteria:

- they represent commonly reported strains in Europe
- they remain stable during the preliminary testing period at the organising laboratory
- they are easy to type
- they represent the three different subtypes of vtx1 and the seven different subtypes of vtx2 in positive and stable strains
- the PFGE profile is stable and represent the diversity of the occurring VTEC strains in Europe



EQA-4 & PT10 includes the following methods

- molecular typing: Pulsed Field Gel Electrophoresis (PFGE)
- serotyping O:H
- virulence determination:
 - genotyping of VTEC virulence genes: vtx1, vtx2, eae, aaiC and aggR
 - subtyping of vtx1 (vtx1a, vtx1c and vtx1d) and vtx2 (vtx2a to vtx2g)
- phenotypic testing
 - production of Verocytotoxin/Shiga toxin
 - production of Extended Spectrum Beta Lactamases (ESBL)
 - production of β-glucuronidase
 - production of enterohaemolysin
 - fermentation of sorbitol



Test panel

 In total, 15 blinded *E. coli test* strains, including 11 strains that will be used for PFGE

If requested

- 9 reference control strains covering all the established variants of vtx1 and vtx2 genes
- the global standard strain S. Braenderup (if participation in EQA PFGE)



Deadlines

20th of Nov. 2012

Response to the invitation to ecoli.eqa@ssi.dk

26th of Nov. 2012 -3rd of Dec. 2012

25th of Feb. 2013

Shipment of test and control strains

Deadline for submission of the results



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Eva Møller Nielsen Head



Mia Torpdahl Salmonella



Jonas Larsson Listeria



Søren Persson Curator