Survey of potentially zoonotic E. coli of serogroup 026, 0103, 0111, 0145 and 0157 in Norwegian cattle

10th Annual Workshop of the NRLs for *E. coli* Rome, 6. November 2015

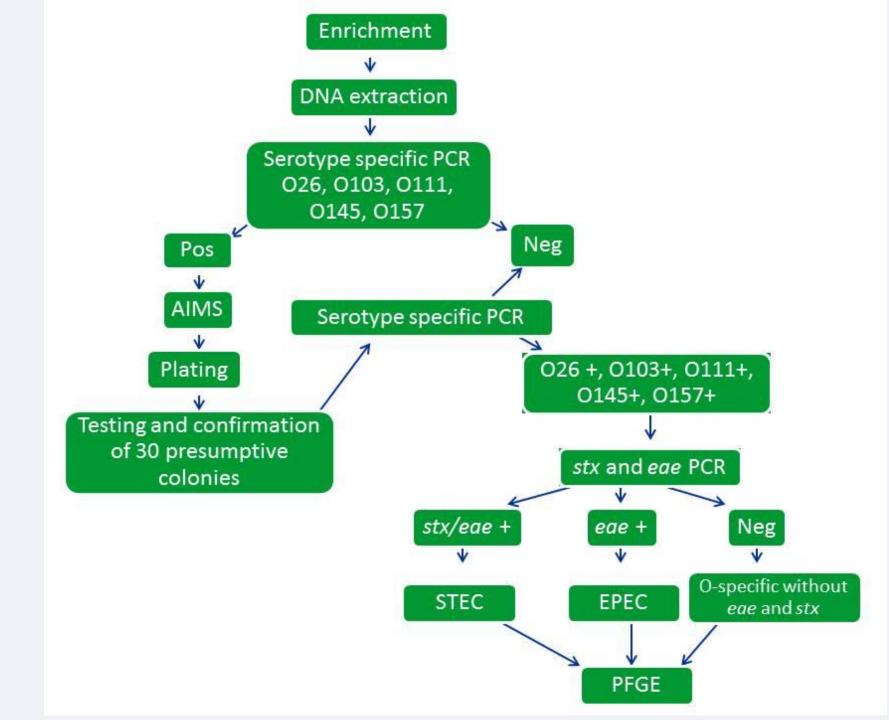
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Introduction

- Nationwide survey in cattle 2014
- Investigating the occurrence of STEC serogroups: 026, 0103, 0111, 0145, 0157
- 179 dairy herds with >50 cows







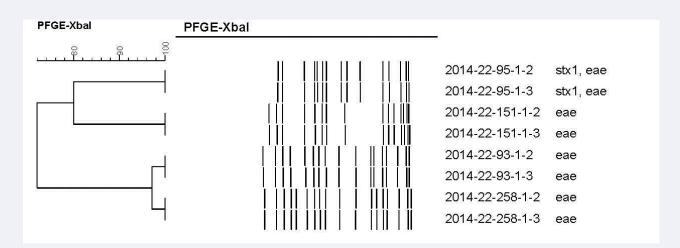
Results 0111 in cattle

- 3 PCR positive samples (Ct values 39.7-41.6)
 - 30 colonies per herd (n=90)
 - 3 aggl+/indol+ from 1 herd
 - 3 colonies for MALDI
- Presumptive O111 were identified as Morganella morganii with MALDI



Results 0145 in cattle

- PCR positive herds: 4 (Ct values 38.5-42.4)
 - 30 colonies per herd (n=120)
 - 47 aggl+/indol+
 - 5 colonies per herd for MALDI and PCR characterization
- Culture confirmed positive herds: 4 (2.2%)
- 0145, stx_1 , eae 1 herd
- 0145, *eae* 3 herds





Results 0157 in cattle

- PCR positive herds: 31 (Ct values 31.9-43.7)
 - 19-30 colonies per herd (n=909)
 - 143 aggl+/indol+
 - 1-5 colonies per herd for MALDI and PCR characterization (n=74)
- Culture confirmed positive herds: 18 (10.2%)
 - STEC 0157

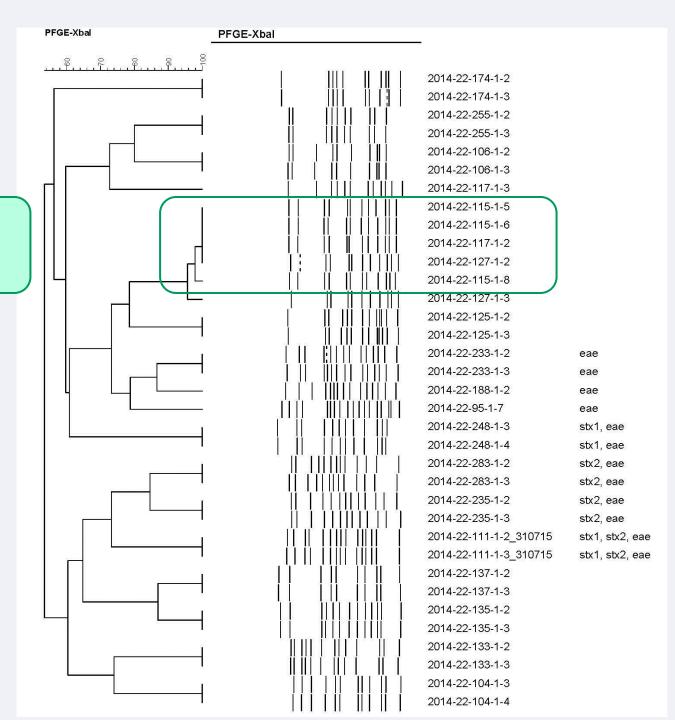
4 herds (2,2%)

• EPEC 0157

- 3 herds (1.7%)
- Other *E. coli* O157 11 herds



Isolates from 3 herds





Results 026 in cattle

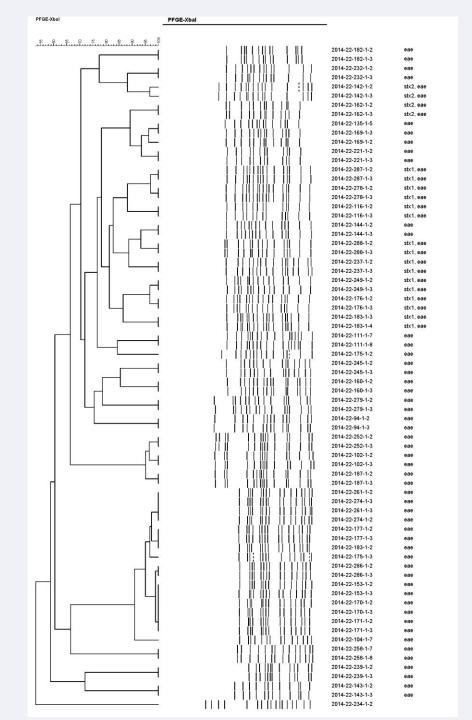
- PCR positive herds: 70 (Ct values 27.7-43.4)
 - 25-30 colonies per herd (n=2083)
 - 381 aggl+/indol+
 - 1-5 colonies per herd for MALDI and PCR characterization (n=160)
 - 5 colonies were not *E. coli*
- Culture confirmed positive herds: 37 (20.7%)
 - STEC 026

10 herds (5,6%)

• EPEC 026

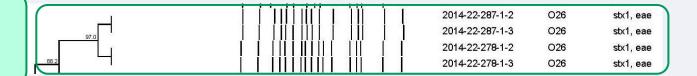
27 herds (15.1%)

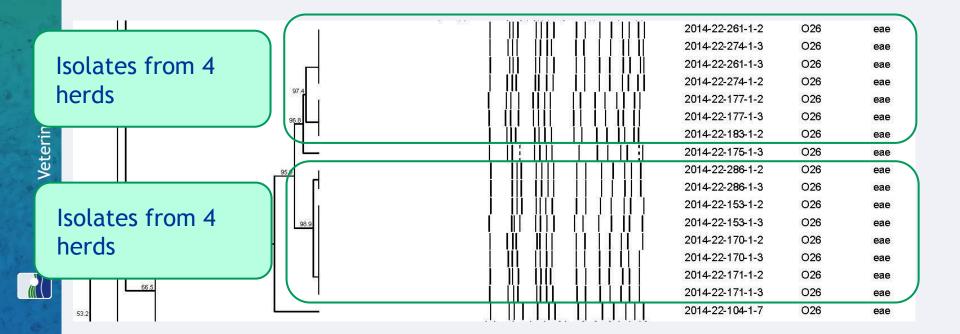






Isolates from 2 herds





Results 0103 in cattle

- PCR positive herds: 83 (Ct values 28.5-44.5)
 - 30 colonies per herd (n=2490)
 - 324 aggl+/indol+
 - 1-5 colonies per herd for MALDI and PCR characterization (n=177)
 - 3 isolates were Citrobacter or M. morganii
 - 45 isolates were not O103
- Culture confirmed positive herds: 46 (25.7%)
 - STEC 0103

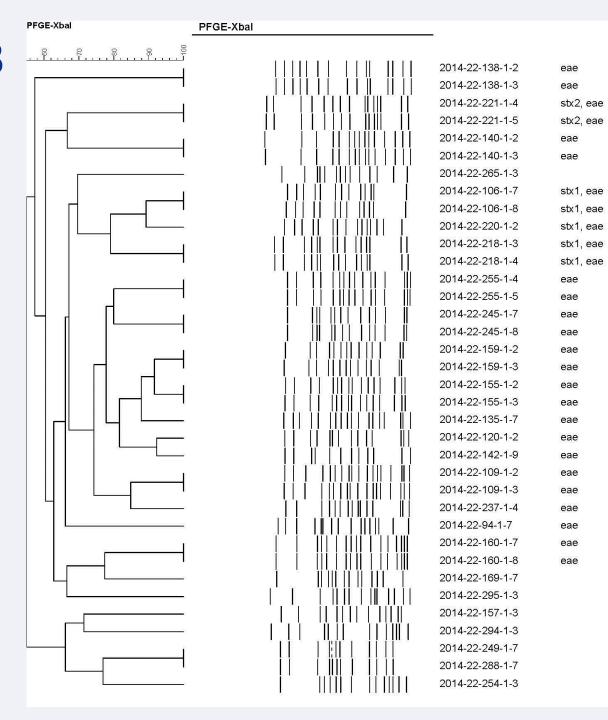
4 herds (2.2%)

• EPEC 0103

- 13 herds (7.3%)
- Other O103 *E. coli* 30 herds



Selected 8 eae-0103 for PFGE





Further work

- Whole genome sequencing for a selection of isolates
 - One isolate per herd different PFGE two selected
 - For eae- O103 only 8 isolates are selected
- Use WGS for:
 - H-typing
 - Virulence characterization (stx subtype, eae, ehxA + others)
 - AB resistence genes
 - MLST and SNP typing
- Screen for serogroup O121, O91 and a specific stx2a PCR



Summary

- Large difference between PCR pos. samples and culture confirmed samples
 - Pick colonies when positive real-time PCR regardless Ct value
- Relatively high prevalence of E. coli O26 and O103 in Norwegian cattle herds
 - Low prevalence of STEC
- More in-depth information about the isolates by WGS



Norwegian Veterinary Institute

Acknowledgement











