



Clusterdetection and interpretation using cgMLST and SNP analysis

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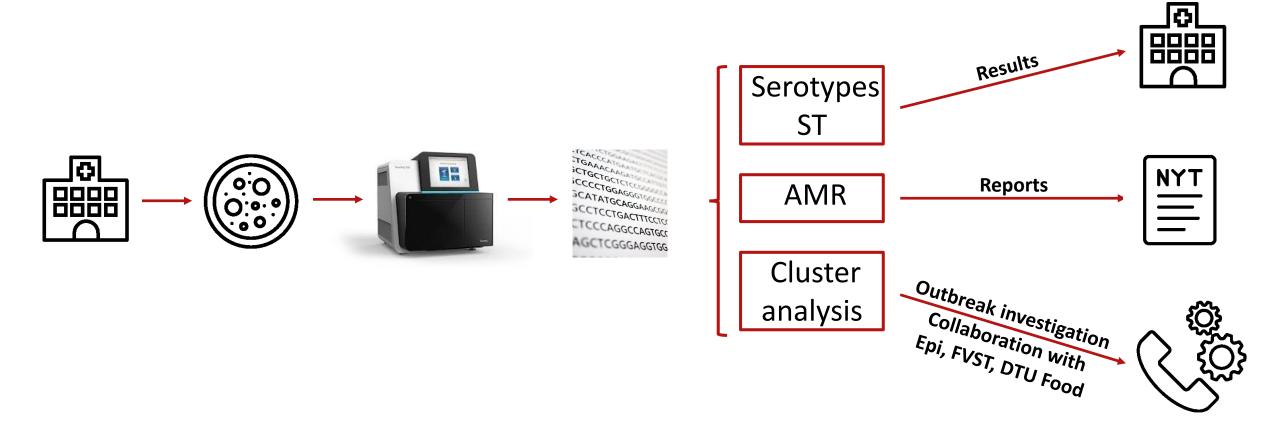


Content

- Surveillance of *Salmonella* in Denmark
 - Outbreak definition and communication
 - Cluster analysis and detection
 - Examples on cluster detection
- Surveillance of *Campylobacter* in Denmark

WGS-based surveillance of Salmonella in Denmark

- Real time surveillance since 2018
- Flow from Hospital to SSI



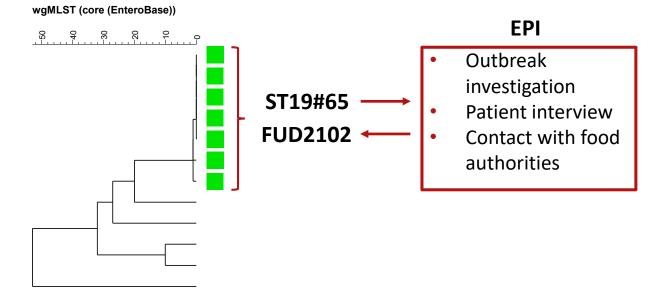


Outbreak definitions and communication



Outbreak definition

- Genetic cluster of \geq 4 isolates within 3 months
- Communication
 - Genetic cluster ID
 - Outbreak ID
- Retrospective analysis
 - Cases to existing cluster new event
 - Same cluster ID but different outbreak ID



Cluster analysis and detection



- Cluster detection
 - cgMLST and single linkage clustering
 - Applied Maths/Enterobase scheme Bionumerics
 - Cluster cut-off
 - ≤ 3 AD
 - ≤ 1 AD for clonal types (Enteritidis ST11 and monophasic Typhimurium ST34)
- Inclusion criteria not one fits all
 - Type, time, place
 - Typing method (SNP, cg/wgMLST)
 - Clustering methods

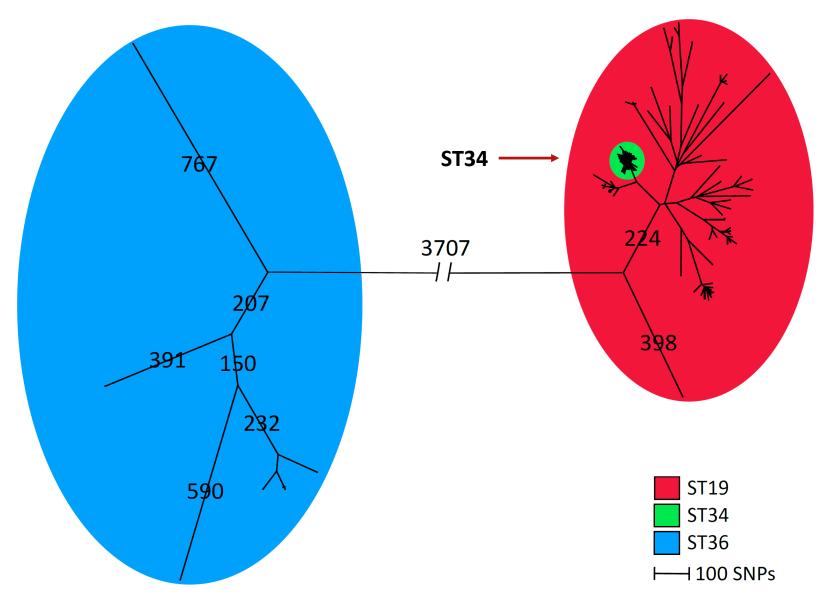
Clonal types

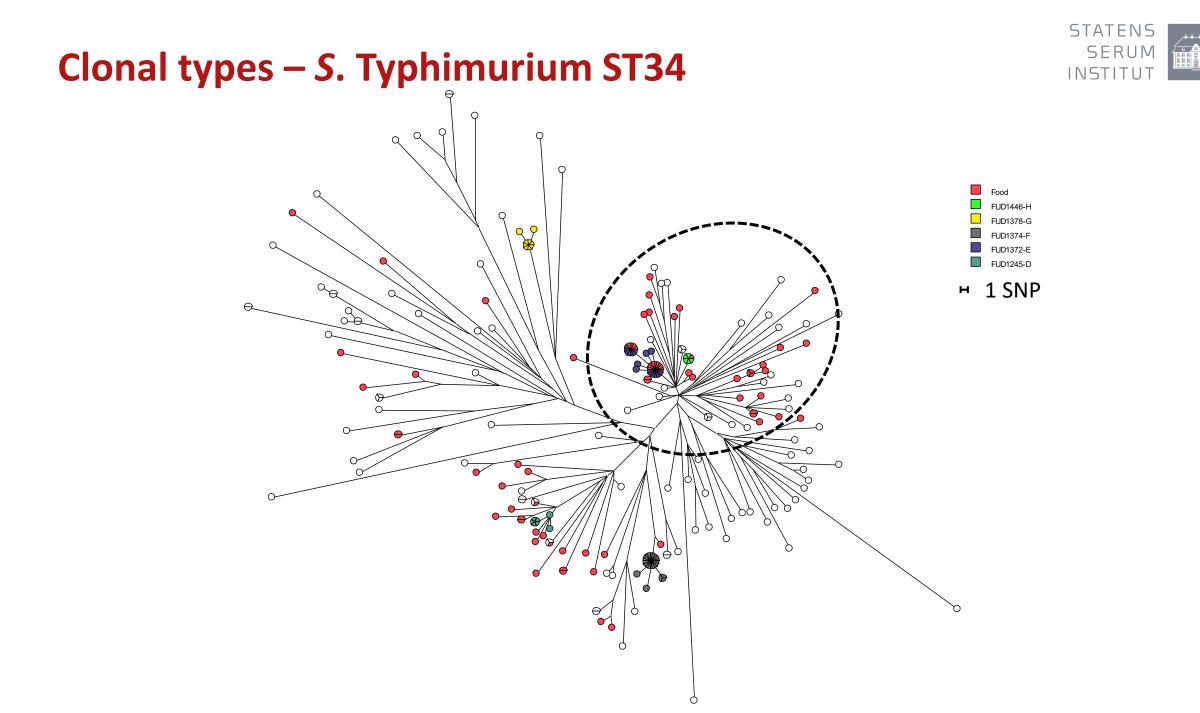


- Enteritidis ST11 and monophasic Typhimurium ST34
 - New clones
 - Little genetic diversity
 - Challenging cluster analysis
 - Supporting analysis SNP or wgMLST
 - Additional analysis AMR, plasmids, prophages
 - Other clustering methods
 - More strict cluster cut-off

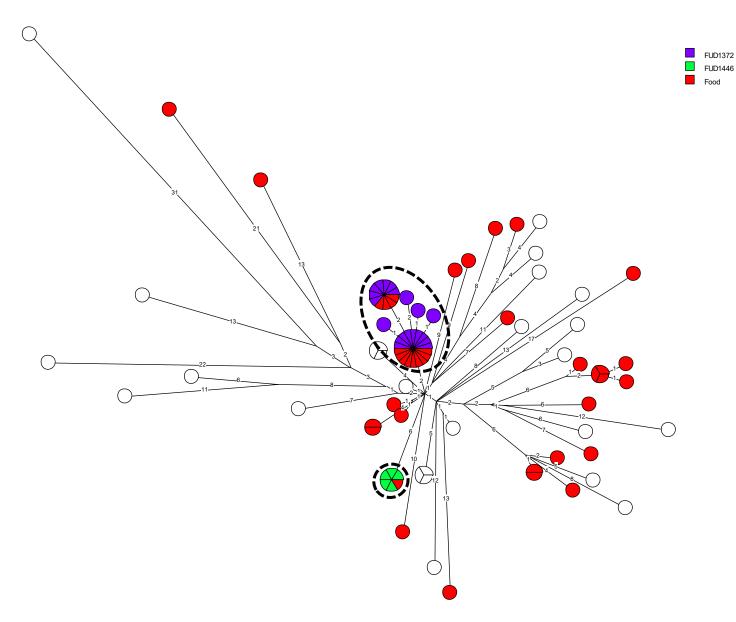
Clonal types – S. Typhimurium ST34







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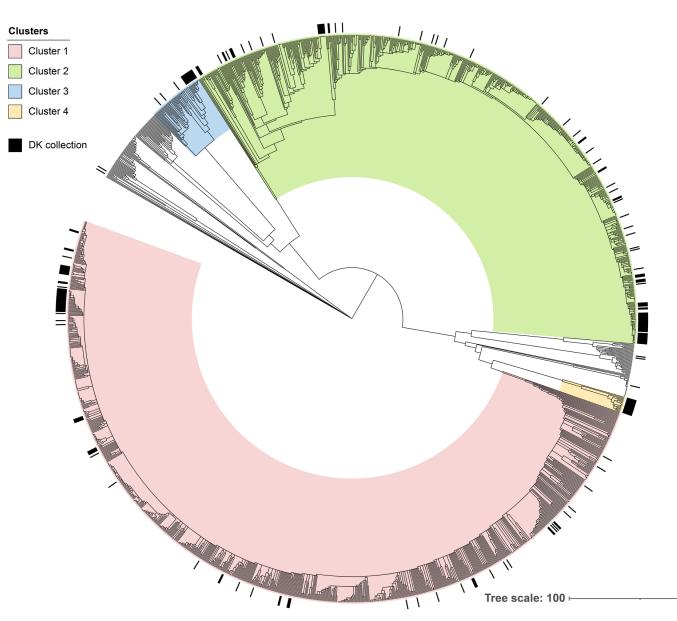




- Outbreak FUD1372
 - Defined by small plasmid

- Outbreak FUD1446
 - Defined by specific epi-data

Clonal types – S. Enteritidis ST11

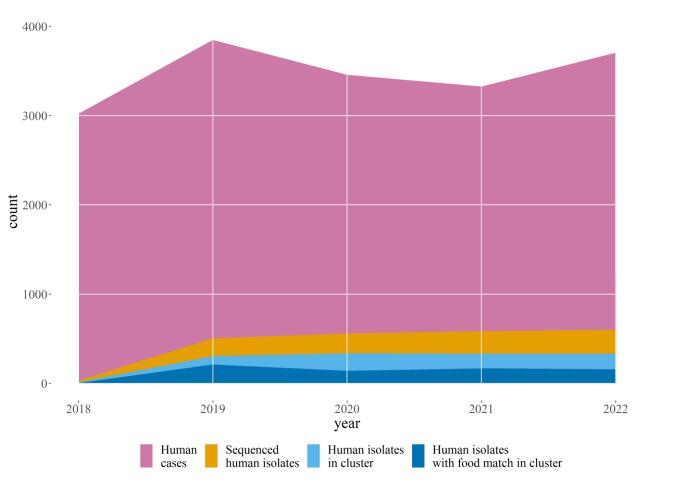




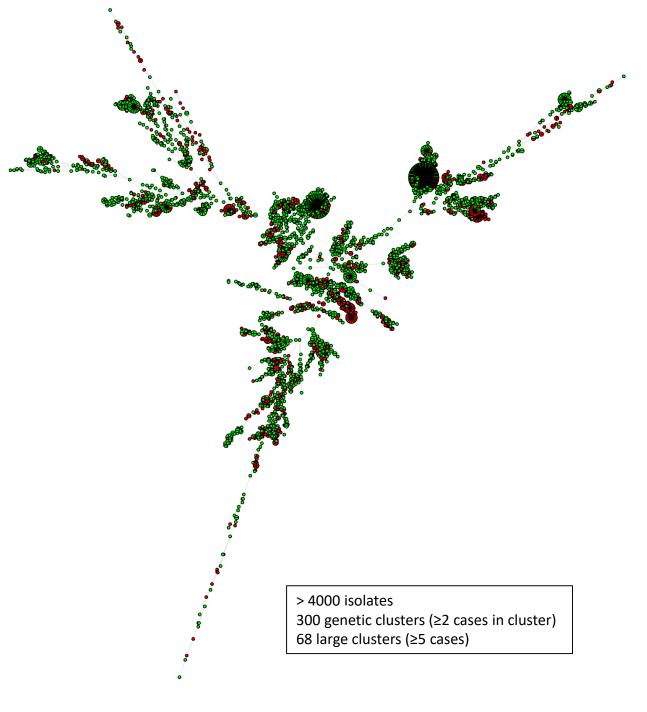
- 2 major clones in Europe
- Extremely little diversity
 - Cluster definition
 - Strict cut-off
 - UPGMA instead of single linkage
 - Additional wgMLST analysis

WGS-based surveillance of Campylobacter jejuni/coli

- Performed routinely since 2019
- Isolates from 10-15% of human cases
- 4/5 regions of Denmark represented
- Food data analyzed routinely at SSI
 - To find food matches









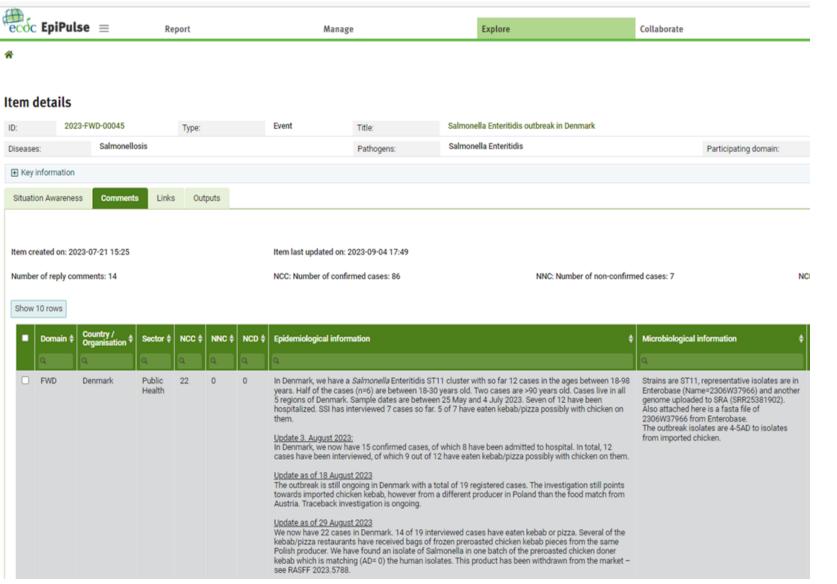
- *Campylobacter* overall very diverse
- cgMLST and single linkage
 - \leq 4 AD cut-off
- ~50% of human isolates form clusters
 mostly small, some big
- 25-30% of human isolates match to food isolates (mostly chicken)



Concluding remarks

- Not one fits all
 - Interpretation of results from cluster to cluster
 - Find the method that fits the data and types seen in your country
 - Do validation using well defined outbreaks
- Clear definitions and communication is important
 - Clear information on tools and methods
 - Clear information on cluster and outbreak definition (type, time, place)
 - Better understanding between lab and epi
 - Better understanding between countries
- Epi data is extremely important
- It's a working progress ongoing evaluation of data is needed
- Questions?
 - Is it better to include to many or to few?

Epipulse event – Outbreak of Enteritidis ST11



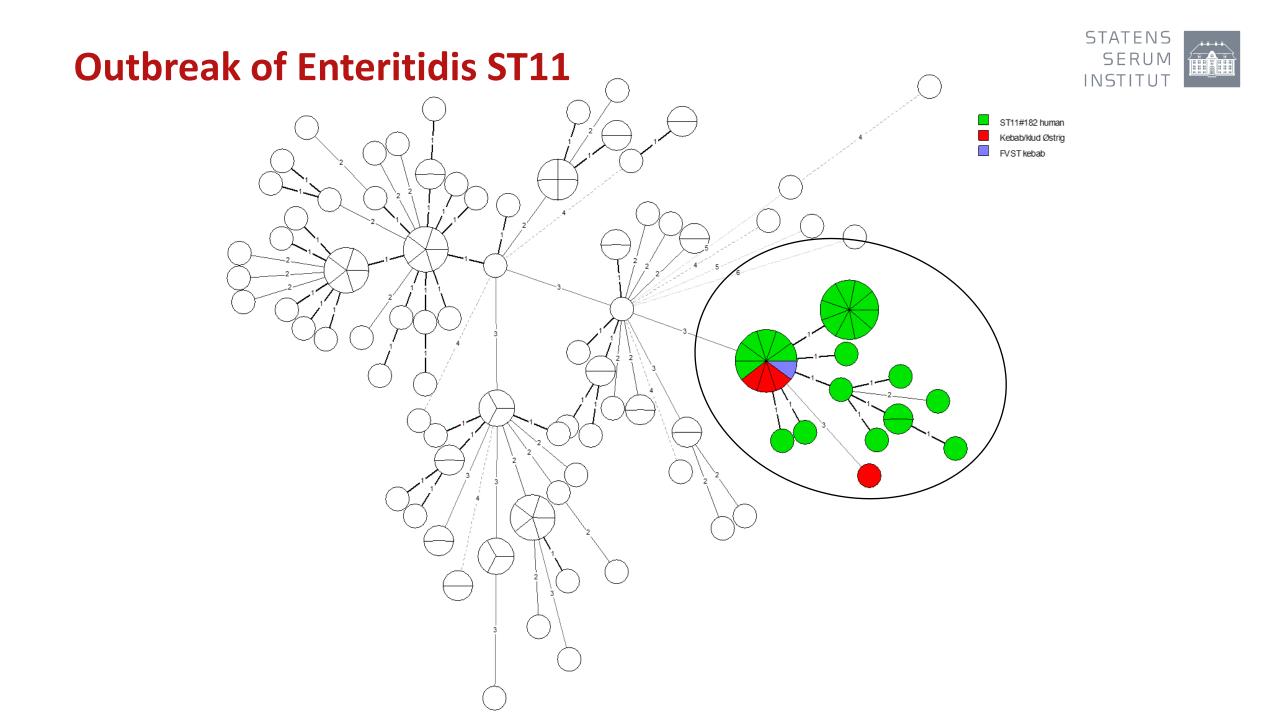


National outbreak of ST11

related to chicken kebab

- Epipulse event
 - Epidata
 - Mikrobiological data
 - Attached sequence

Turned into a international outbreak





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Aktuelt og presse Sygdomme, beredskab og forskning Vaccination Produkter og ydelser Om SSI

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Internationalt udbrud af Salmonella Enteritidis ST11 relateret til Polsk kyllingekebab

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