

Application of whole genome sequencing for public health interventions around foodborne pathogens

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Public Health Estimates of foodborne infectious disease



- Estimated more than 500,000 cases of foodborne illness linked to known pathogens in the UK annually
- High economic burden £1.8 billion annually in medical, financial and welfare costs

Estimate of number of cases per year of illness due to common foodborne pathogens

Campylobacter	280,000	Cryptosporidium 2770		
C. perfringens	80,000	Giardia	8000	
E. coli 0157	10,000	Norovirus	74,100	
Listeria	183			
Salmonella	33,100		O'Brien et al, BMJ Open 2016 doi: 10.1136/bmjopen-2016-011119.	





Gastrointestinal Bacteria Reference Unit

- National Reference Laboratory
- Salmonella Reference Service
- *E. coli*, Shigella, Vibrio and Yersinia Reference Service
- Foodborne Pathogen Reference Service Listeria monocytogenes, Bacillus cereus, Clostridium botulinum and Clostridium perfringens, Staphylococcus aureus
- Campylobacter and Helicobacter Reference Service

Receive bacterial isolates from clinical microbiology laboratories and food and water laboratories ie from clinical cases and from foods, across England and Wales for identification and typing for national surveillance and for outbreak detection and investigation



Value of Typing

- Provides information to assist the understanding and management of local outbreaks/incidents
- Identifies clusters and outbreaks occurring at local, regional, national level

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- Provides microbiological evidence to identify and track source of contamination so effective control and prevention measures can be implemented
- National surveillance and trend monitoring over time
- Monitoring effectiveness of control and preventative measures







Outbreak investigation questions

- Is there an increase in a particular type a cluster of cases/outbreak?
- Is the type from clinical cases same as the type from food – if available?
- Have we seen this type before in food?
- Is this a known type in the UK or elsewhere?



Move to WGS

- WGS provides a single one step method for identification and typing and provides a wealth of additional information
- Faster TATs, lower costs
- High quality data, standardisation, rapid global comparability
- Safety



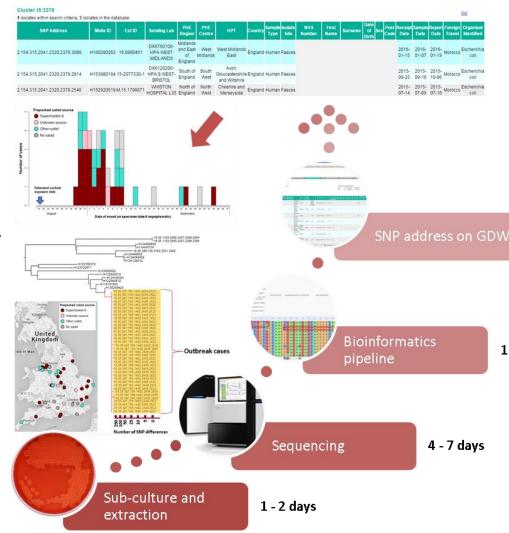


- Highest degree of resolution for strain discrimination
- Provides phylogenetic information showing unequivocal strain relatedness
- Improved cluster detection
- Provides information on potential sources and geographical location
- Ability to rapidly screen large number of isolates for virulence genes, AMR genes
- Leads to improved control measures



Overview

WGS of: Salmonella E. coli STEC Campylobacter Listeria



- \rightarrow Identification
- \rightarrow Serotype
- \rightarrow MLST ST, CC
- ^{1-5 days} → Virulence and AMR gene detection
 - ightarrow SNP typing address
 - \rightarrow Cluster detection
 - → Patient and food isolate matching

Assessing similarity between genomes by SNP typing

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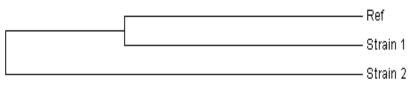
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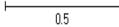
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Single nucleotide polymorphism (SNP) i.e. a different nucleotide to the reference genome

RefGTStrain 1 GGStrain 2 AG

A A G





 10403S - 1.1.1.1.1.1

 RL15001614 - 1.4.18.20.21.23.29

 H160280558 Human - 1.7.24.28.34.36.44

 H154820912 Human - 1.7.24.28.29.31.38

 RL15001291 - 1.3.15.16.16.18.22

 H154360522 Food - 1.2.11.12.33.35.43

 H110400356 Human - 1.2.11.12.32.34.41

 H131960044 Food - 1.2.11.12.12.13.47

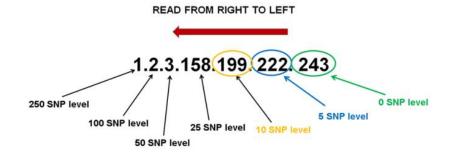
 H132860463 Food - 1.2.11.12.12.13.46

 H132340048 Human - 1.2.11.12.12.13.45

 RL15001372 - 1.2.11.12.13.17

 H140680419 Human - 1.2.11.12.12.13.42

 H140760087 Human - 1.2.11.12.12.13.42



0.2

SNP address



WGS led surveillance

- Identify linked strains with unprecedented sensitivity and specificity
- Provides context for strains
 - Infer strain source
 - Native / Imported
 - Geographical signals within UK
 - Virulence potential
 - Demonstrate relatedness of isolates over years

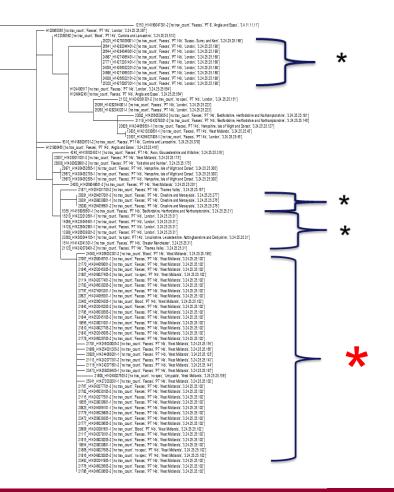
Building global databases – clinical, animal, food

NCBI BioProject PRJNA248042



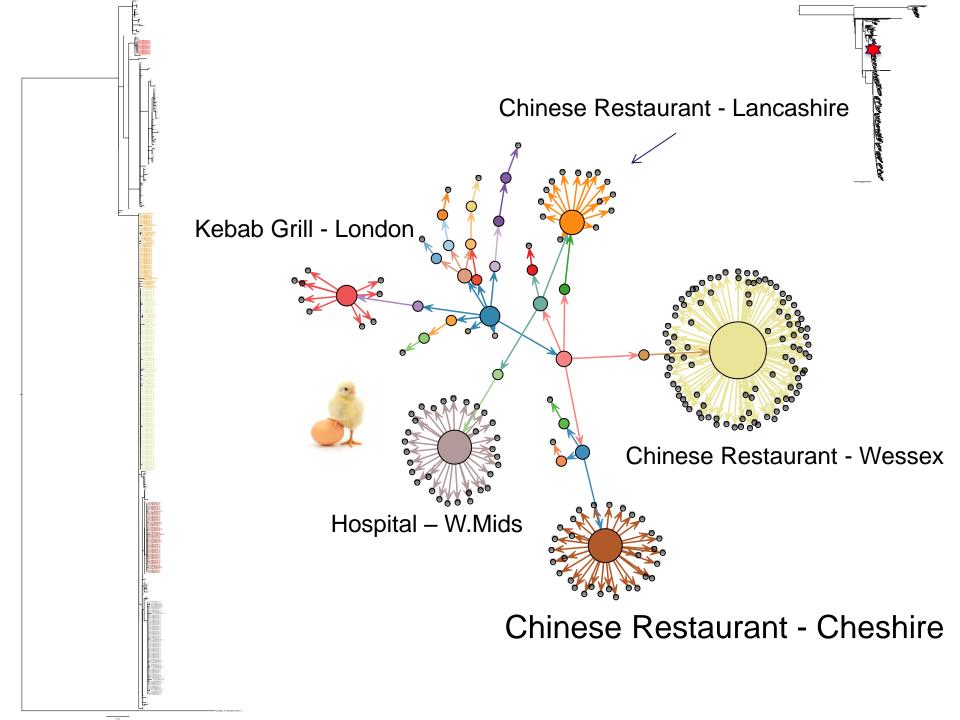
International outbreak of Salmonella Enteritidis PT14b Summer 2014

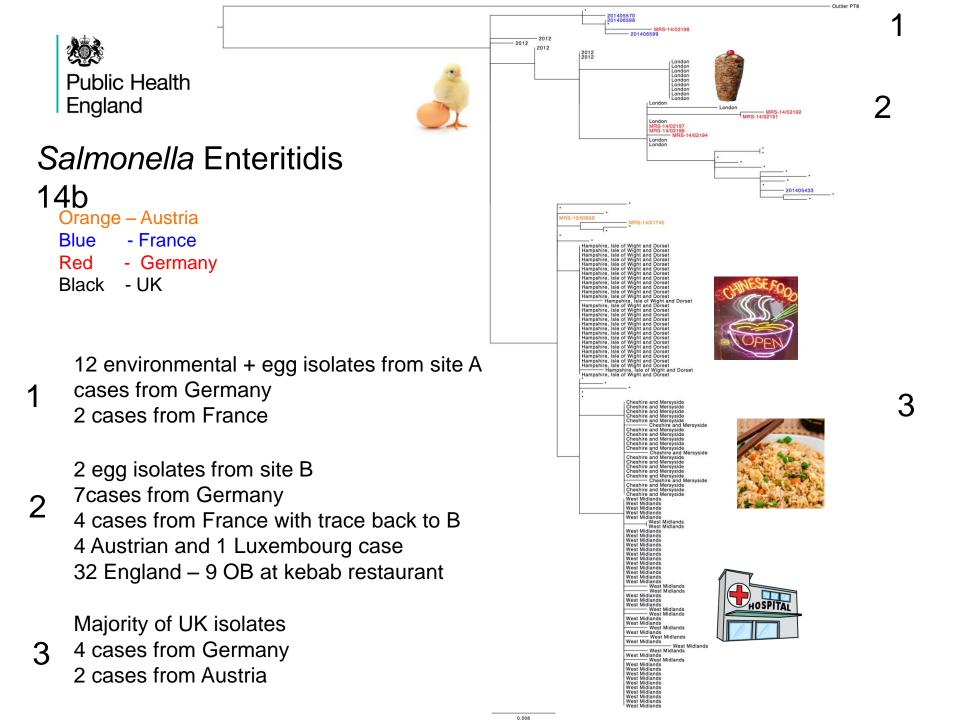
- In England an outbreak of Salmonella Enteritidis PT 14b in a hospital in Birmingham with WGS profile *
- Also noticed small pockets of increase in PT14b around England with WGS profile – individual point source outbreaks *



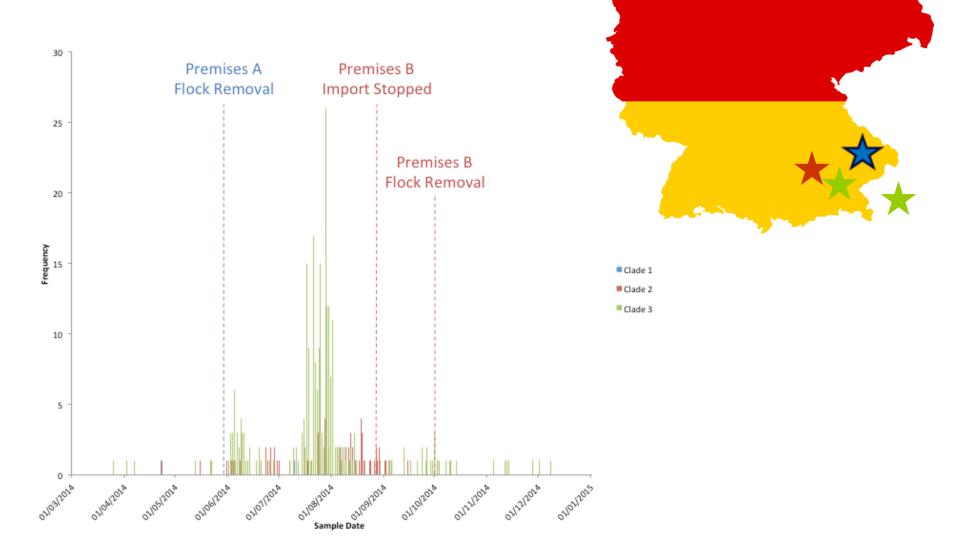
Public Health International Outbreak

- Cases in Austria, Germany, France and Luxemburg plus French egg isolate with German egg mark – same MLVA or SLV as UK
- Trace back in UK All outbreaks involved eggs from same German supplier
- Trace back investigations lead to sampling of egg production premises in Germany
- Company has 4 separate egg production plants supplied with layer flocks from single supplier
- European isolates sequenced cases / eggs





Clade 1 – 12 egg isolates + traceback (site A) Clade 2 – 2 egg isolates + traceback (site B) Clade 3 – 0 egg isolates, 0 traceback (?site C or D)





Shiga toxin-producing (STEC) *E. coli* O157

 Severe bloody diarrhoea and Haemolytic Uraemic Syndrome (HUS).
 Long-term renal, cardiac and neurological sequelae

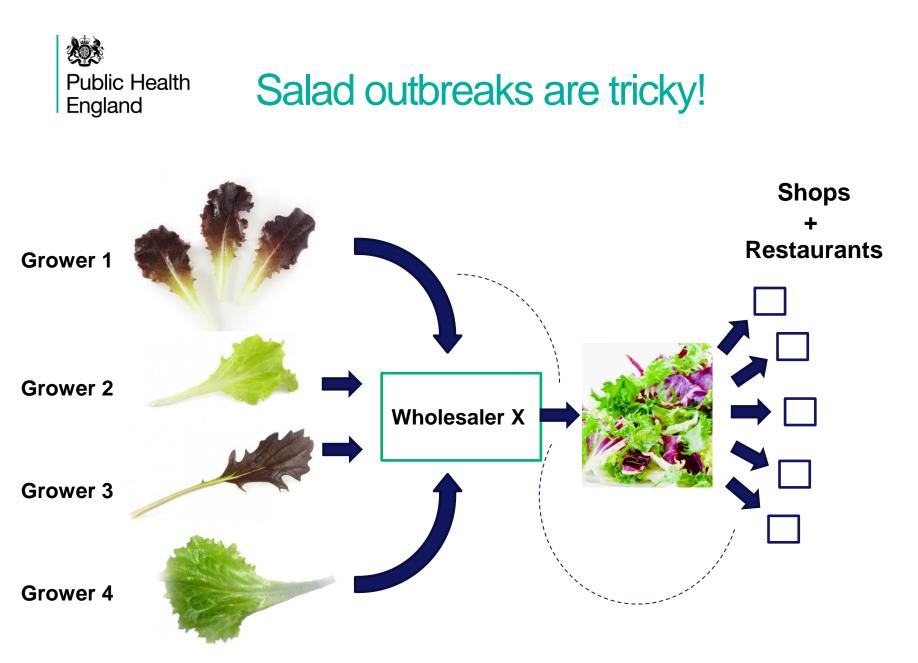
• Limited treatment options – palliative care only as antibiotics contraindicated

• Zoonotic reservoir in the UK mainly cattle, sheep and goats, but almost all animals can act as secondary transmission vectors

• Transmission can be food or waterborne, via direct contact with animals or their environment, or person to person spread





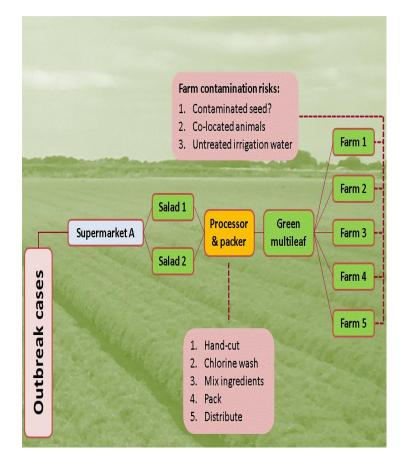


different salad leaf mixes on different days



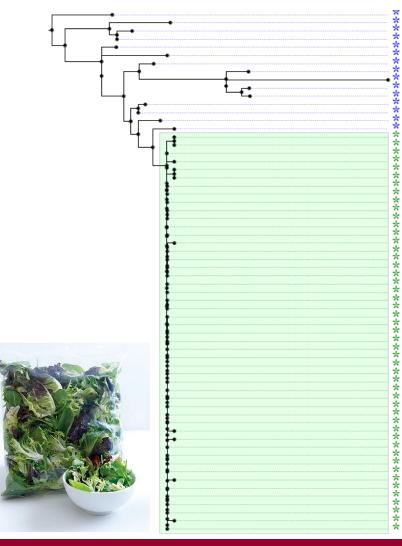
Deeper phylogenetic relationships : National context

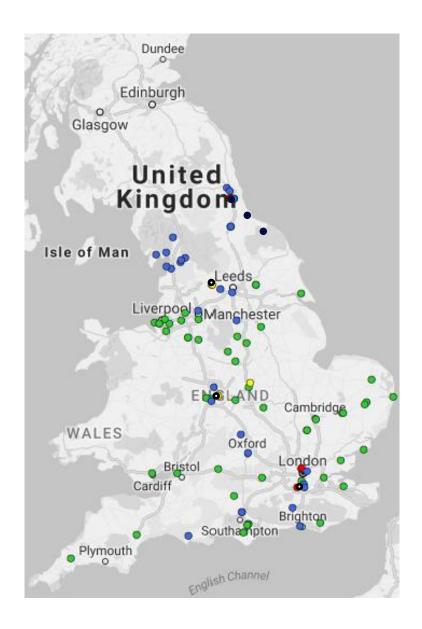
- August 2015 40 cases of STEC 0157 PT 8 detected following routine analysis of the WGS data. Not travel associated, but genetically close to UK cattle and environmental isolates in DB
- Cases exposed to prepackaged salads subsequently found to have particular type of salad leaf in common.
- Green multi-leaf lettuce, supplied to Supermarket A, was grown in England on five different farms, all samples -ve
- EHOs and local HPTs reported poor biosecurity on all farms, eg. animal ingress (wildlife) and use of untreated irrigation water from rivers and ponds



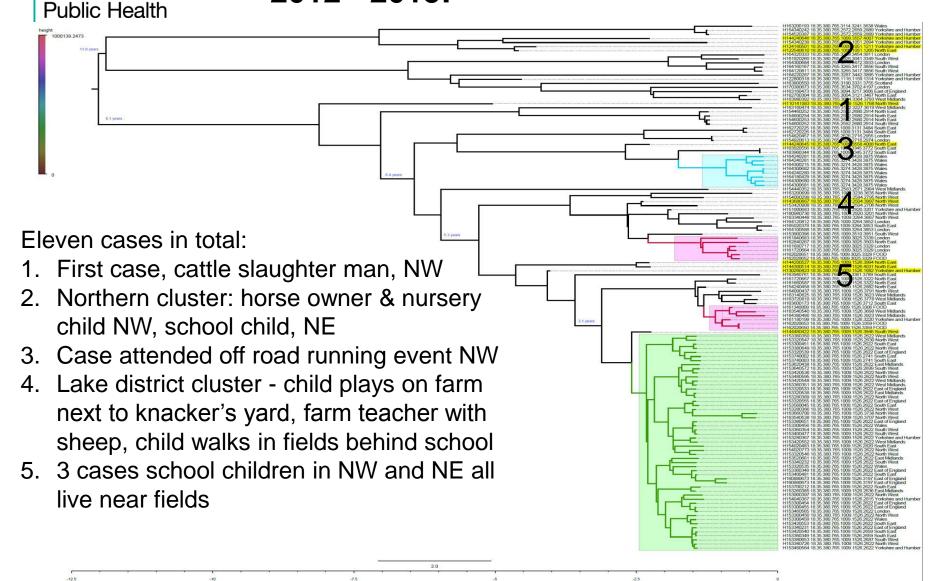


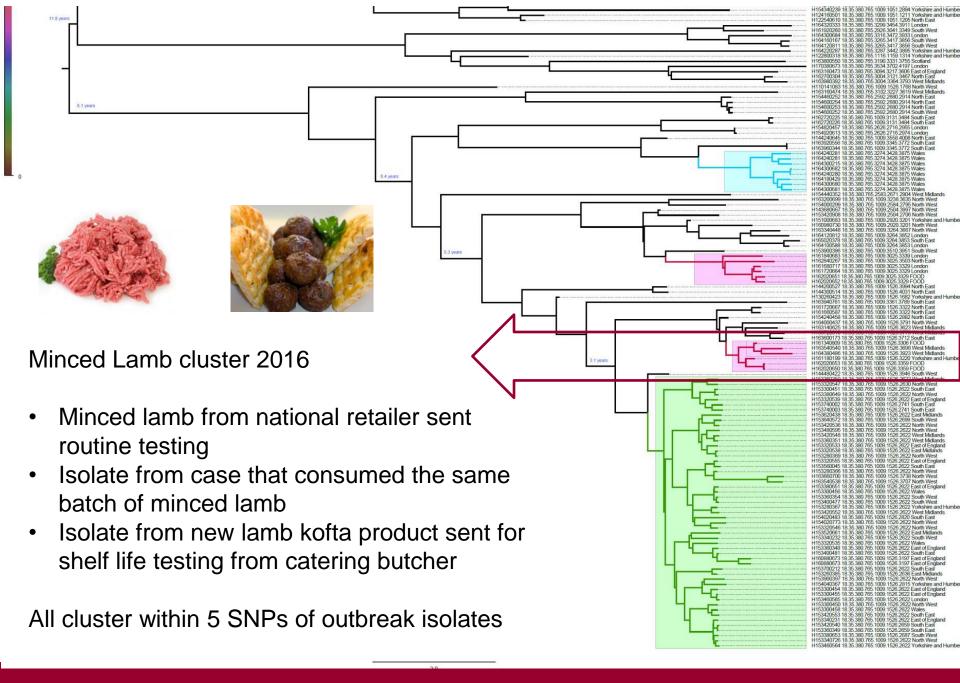


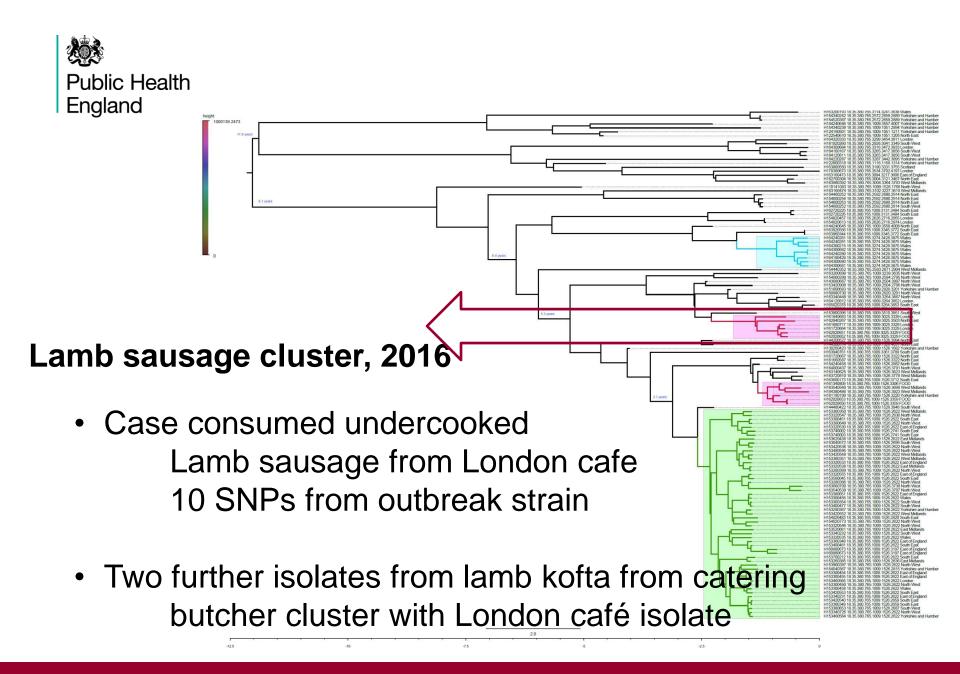




Pre-outbreak cases, 2012 - 2015:









O157 OB summary

- Strain initially detected during the prepacked salad outbreak August -October 2015.
- No new cases in this cluster until March 2016; marked difference in exposures and case demographics; minced lamb or live ovine exposures
- Four independent isolates from minced lamb products in the 10-SNP cluster;
- Increased and continuous activity of this strain to date; although more variation than seen previously, new cases within 5 - 50 SNPs of the original outbreak cluster continue to occur weekly.
- Origin of the prepacked salad outbreak likely to be ovine
- Questions remain around this strain amongst sheep in UK; how widespread is it?

Multicountry European Public Health **Outbreak of listeriosis**



Listeriosis is rare but severe disease with high fatality rate and causes the highest number of deaths due to a foodborne pathogen in Europe

England

- *L. monocytogenes* isolates from 5 European countries have same WGS profile Nov 2017
- Environmental isolate from French frozen food producer from surface used to prepare frozen vegetables including sweetcorn – matches by WGS Nov 2017

- 23 cases with food consumption data:
- 9 reported consumption of sweetcorn and 1 reported possible consumption
- 6 reported consumption of frozen sweetcorn
- 2 reported frozen vegetables, one non frozen both with sweetcorn
- Sweetcorn consumption not part of food questionnaire for all MS ----UK cases re-interviewed 3/4 sweetcorn consumption



On going multicountry European outbreak of listeriosis identified by WGS June 2018

	Number of cases (No deaths)				Total number	Total number
Country					of cases	of deaths
	2015	2016	2017	2018		
Austria	0	2 (1)	0	0	2	1
Denmark	0	0	2	2 (1)	4	1
Finland	0	4	10 (2)	10	23	2
Sweden	0	3 (1)	3	1	7	2
The United Kingdom	1	2	2	6	11	2
Total	1	11 (2)	17 (3)	19 (1)	47	8



Jan and Feb 2018



L. monocytogenes detected in batches of frozen sweetcorn from Finnish trader (50-140cfu/g) with same WGS as OB strain

Sweetcorn produced in Hungary and packed in Poland

Recent Finnish case reported eating frozen mixed veg containing sweetcorn – same brand as above

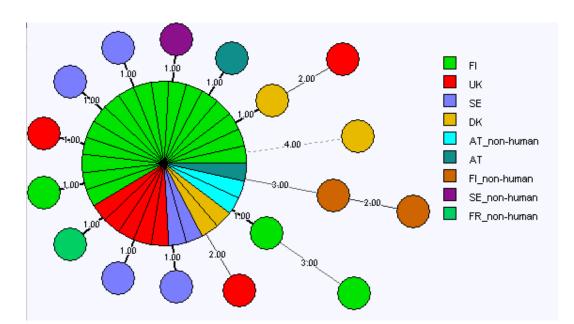
Swedish case reports eating frozen sweetcorn - brand unknown

Isolate with same WGS detected unacceptable levels in sweetcorn from home of a consumer in Sweden – traced back shows packed in Poland and produced in Hungary

Austria reports isolates from 2 different types of frozen mixed vegetables including sweetcorn that originates same company in Hungary



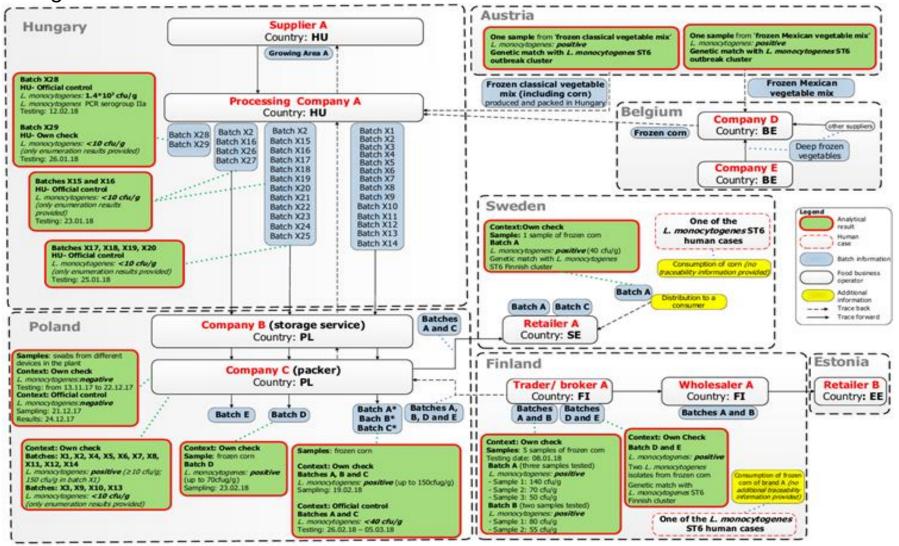
Clonal complex 6



WGS_Allelelds (Core loci)		
°·····	 FI 	
1.0	 FI 	
2.0	DK	2017-05-18
	AT	2016-10-20
	FI	2017-08-31
	UK	2015-12-22
	FI	2018-01-10
	FI	2016-12-02
	FI	2017-06-27
	SE	2016-10-19
	UK	2016-10-19
	SE	2017-09-06
	FI	2017-09-08
	FI	2017-08-02
	FI	2017-01-11
	DK	2017-01-11
	FI	2017-01-08
		2017-12-14
	FI	
	FI	2016-10-28
		2017-02-02
	FI	2017-03-23
	• AT	
1.0	• AT	2047.00.22
1 1 1	FI	2017-06-23
1.0	UK	2016-05-08
1.0	SE	2016-11-25
1.0	★ FI	2017-09-26
1.0	DK	2018-02-02
1.0	SE	2016-06-09
1.0	UK	2018-01-17
1.0	SE	2017-12-06
1.0	 SE 	
1.0	FR	
1.0 1.0	SE	2017-10-26
1.0 1.0	AT	2016-11-03
2.0	UK	2017-11-24
3.0	DK	2018-02-20
3.0	FI	2017-04-23

WGS_Allelelds (Core loci)

Public Health Frozen sweetcorn investigations



Food investigations UK (11cases) Public Health England

- Company supplying sweetcorn is large European-wide company suppling fresh and frozen vegetables + other products to major retailers, food service companies and industry across Europe
- 10 production sites across Europe including 2 in UK that supply a range of major supermarkets and food service providers many of which have been mentioned or have links with one or more cases in the UK
- Frozen Sweetcorn from implicated Company in Hungary is supplied to the 2 production sites in the UK and thence to various supermarket and food service suppliers and is known to be supplied to hospital where one of UK cases was inpatient
- Two recent cases have major supermarket branded frozen sweetcorn originating from Hungary in their home freezers. Supermarket known to be supplied by UK production sites – sampling and testing underway

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- First multicounty European outbreak of listeriosis identified by WGS
- Provides unequivocal evidence that isolates are related and come from a common source and have been present in food chain for 2-3 years
- Isolates from frozen sweetcorn originating in Hungary have same WGS profile isolates from cases
- Need to implement effective control measures to prevent more cases recalling contaminated product
- Improve communication on preventing consumption of uncooked frozen sweetcorn



WGS summary

- WGS is being used in real time for monitoring Salmonella, STEC, Shigella, Campylobacter and Listeria monocytogenes
- Highest degree of resolution for typing plus phylogenetic information unequivical evidence that strains are related and from a common source
- Improved surveillance and outbreak detection and investigation:
 - \circ $\,$ Real time monitoring of clusters, of virulence and AMR of all strains
 - Detecting more outbreaks smaller OBs, geographically spread, over longer time frame, international outbreaks
 - Accurate and robust outbreak definition finds cases and rules out unrelated cases from outbreak – refines OB investigation
 - o Increased case ascertainment and indication of location/source of infection

WGS is improving public health interventions for controlling foodborne pathogen illness and the monitoring of their effectiveness



Acknowledgements

All staff in GBRU and all those in Gastrointestinal Infections Department

PHE Food and Water Laboratories and NHS hospital clinical laboratories and PHE regional labs

PHE Genomic Services Unit and Bioinformatics Unit

PHE Health Protection Teams, Field Epidemiology Services

Food Standards Agency, Animal and Plant Health Agency

Local Authorities and Environmental Health Officers

ECDC, EFSA and all EU member states involved in outbreak investigations