



LE GOUVERNEMENT
DU GRAND-DUCHÉ DE LUXEMBOURG
Ministère de l'Agriculture, de la Viticulture
et du Développement rural

Administration luxembourgeoise vétérinaire
et alimentaire



Salmonella in foodstuff and veterinary diagnostics

Dr. Alexandra SCHOOS

Veterinary inspector
Technical manager

Luxembourg Veterinary and Food Administration (ALVA)
State Laboratory of Veterinary Medicine

EURL Salmonella Workshop
22.-23.05.2023

Our missions



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- **Animal health:** Detection of notifiable diseases, zoonoses and pathogens



- **Public health:** Bacteriological quality of food of animal origin (Safety criteria and hygiene criteria)



- **“One Health” approach:** surveillance of emerging diseases



Our lab in numbers



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4 Veterinarians	12 Technicians	1 Quality manager	1 Secretary accountant	2 Receptionist	1 IT	1 Metrologist	2 Technical and logistics assistant



Year	Requests	Samples	Analyses	Salmonella
2020	10.184	222.374	245.227	2.769
2021	11.388	258.198	289.706	2.950
2022	10.083	270.282	305.572	2.690



Year	Requests	Samples	Analyses	Salmonella
2020	240	1.167	5.923	655
2021	179	870	5.272	544
2022	251	1214	7.534	841



Samples and methods



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- Serology (Swine Salmonella Ab Test IDEXX) in **meat juice** samples of slaughter pigs
- Culture (ISO 6579-1*, MSRV, XLD, RSal) and identification (Kauffmann-White*) on **organs, feces and boot socks** of different species
 - EU 517/2011 : laying hens
 - EU 200/2012 : broilers
 - EU 1729/2020 : AMR-monitoring
- Culture (BRD 07/11-12/05*, RSal, ISO 6579-1* on request) and identification (Kauffmann-White*) in all **foodstuff of animal origin** from retail, processing plants, dairy and border control point



*accredited

In 2021...



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Species	Sample type	Detected	Not detected
Porcine	Meat juice	404	1.644
	Organs	0	11
	Feces/boot cover	44	179
Bovine	Organs	1	52
	Feces	8	106
Poultry	Boot cover/feces/dust	8	435
	Organs	0	24
Goat/sheep	Organs	1	21
	Feces	0	3

S. Goldcast, S. Rissen, S. Typhimurium, monophasic S. Typhimurium, S. Brandenburg, S. Bredeney, S. Derby, S. Enteritidis, S. Infantis, S. Kottbus

S. Typhimurium, S. Havana, S. Enteritidis

S. Typhimurium, S. Diarizonae, S. Salamae, S. Paratyphi B

S. Dublin

Veterinary diagnostics: antisusceptibility testing



Species	Sample type	Detected	Not detected
Animal origin	foodstuff	4	540

S. Typhimurium, monophasic S. Typhimurium, S. Brandenburg



In 2022...



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Species	Sample type	Detected	Not detected
Porcine	Meat juice	444	1524
	Organs	0	1
	Feces/boot cover	9	15
Bovine	Organs	3	54
	Feces	8	163
Poultry	Boot cover/feces/dust	6	400
	Organs	0	30
Goat/sheep	Organs	0	8
	Feces	0	2
Wild animals	Organs	0	6

S. Goldcast, S. Rissen, S. Typhimurium, monophasic S. Typhimurium

S. Typhimurium, S. Muenster, S. Indiana, S. Dublin

S. Enteritidis, S. Dublin, S. Brandenburg

Veterinary diagnostics: antisusceptibility testing



Species	Sample type	Detected	Not detected
Animal origin	foodstuff	1	840

monophasic S. Typhimurium



Data processing and reporting



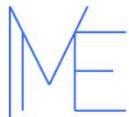
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- Whole genome sequencing (Illumina®) + collection in a database (SeqSphere®) and comparison with human cases



- **efsa** reporting: prevalence + AMR
EUROPEAN FOOD SAFETY AUTHORITY

- National antibiotic action plan

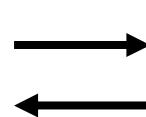
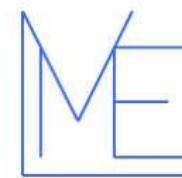


National reference laboratory



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- *Salmonella*
- *Listeria monocytogenes*
- VTEC
- CPS – Coagulase positive *Staphylococcus*
- *Campylobacter*
- AMR



2 labs = 1 team



Thank you for your attention!



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EURL Salmonella Workshop 2023



Catarina Martins

Technical Lab Manager for Food
Microbiology and Molecular
Biology

Laboratoire National de Santé
Food Monitoring Service / Surveillance Alimentaire
Food Microbiology Lab

Service of Surveillance Alimentaire



- 1 – the biggest official control laboratory for food and feed in Luxembourg
- 2 – Accreditation according to ISO 17025 since 2003
- 3 – National Reference Laboratory in 14 domains

Expertise:

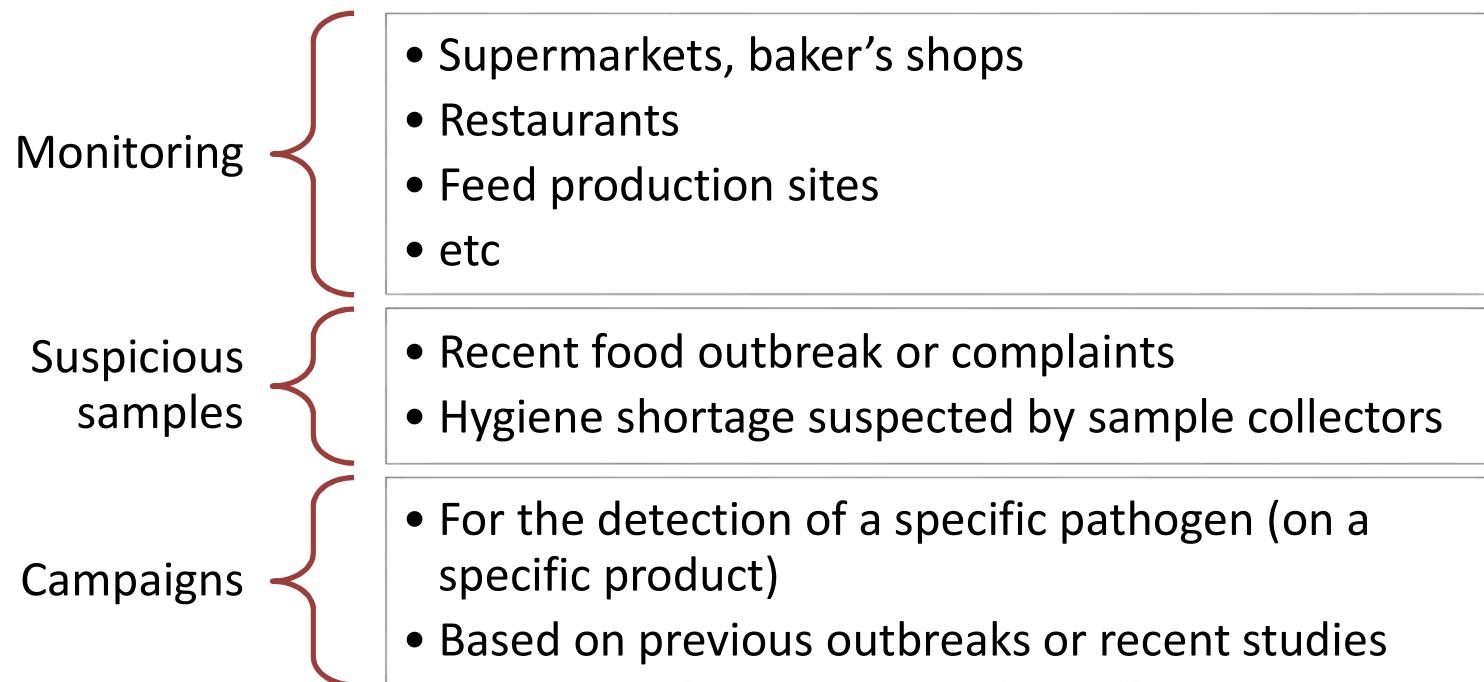
- Microbiological analysis (*Salmonella*, *L. monocytogenes*, STEC, *S. aureus*, *Foodborne Virus*)
- GMO: detection and quantification of genetically modified organisms
- allergens using ELISA and PCR technology
- pesticides in fruits & vegetables, cereals and feed
- mycotoxins and plant toxins in food and feed
- additives (food dyes, sweeteners, preservatives)
- process contaminants: furan, acrylamide, MCPD, PAHs
- food contact materials (FCM)



Service of Surveillance Alimentaire – Food microbiology



- +/- 2500 analyses for *Salmonella* every year
- Diversity of samples : all kind of samples except for a part of animal origin products
- 1-2 positif samples per year
- Samples are brought to us by our competent authority (ALVA)



Salmonella surprise

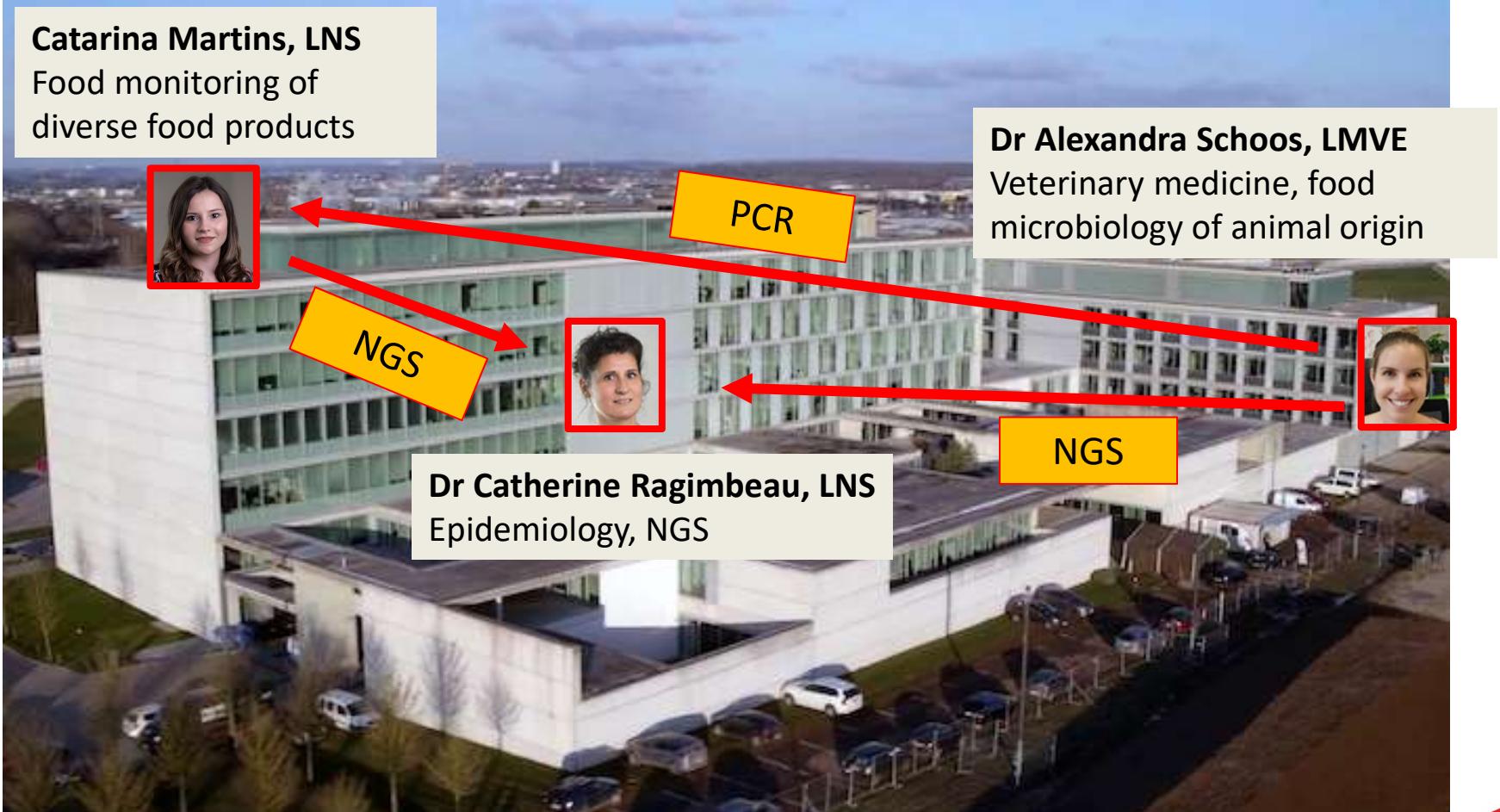


- Approximately 30 samples of chocolate products reached our lab in march/april 2022
- Most of the suspicious lot numbers of the products weren't available anymore
- Analyses with 2 methods :
 - Real-time PCR (iQ-Check Salmonella Kit on CFX 96)
 - ISO 6579-1 method using RVS and MKTTn
→ isolation on Compass *Salmonella* and XLD + confirmation on mini Vidas
- None of the samples was positif
- In case of positif sample : isolation of colonies to give them to Dr Ragimbeau's service for sequencing and matching with human cases



<https://www.foodsafetynews.com/2022/04/kinder-chocolate-eggs-other-products-recalled-in-canada-because-of-link-to-multi-country-salmonella-outbreak/>

Collaboration of the 2 luxembourgish reference labs



Thanks for your attention!





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Epidemiology and Microbial Genomics (NGS Platform)

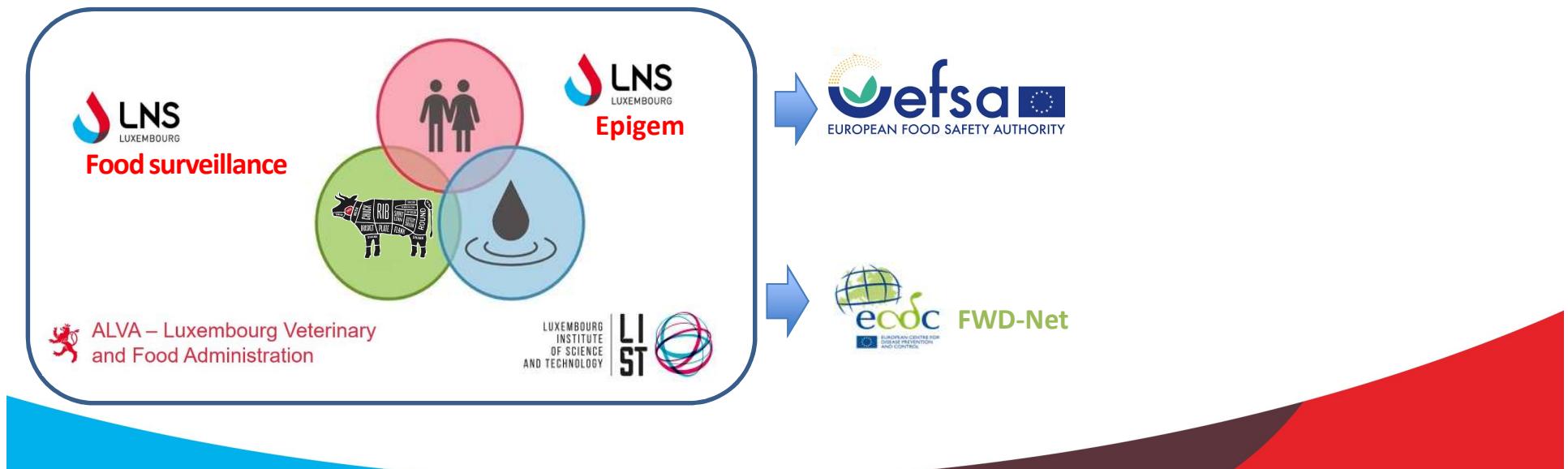
Microbiology Department



EPIGEM EPIDEMIOLOGY AND MICROBIAL GENOMICS



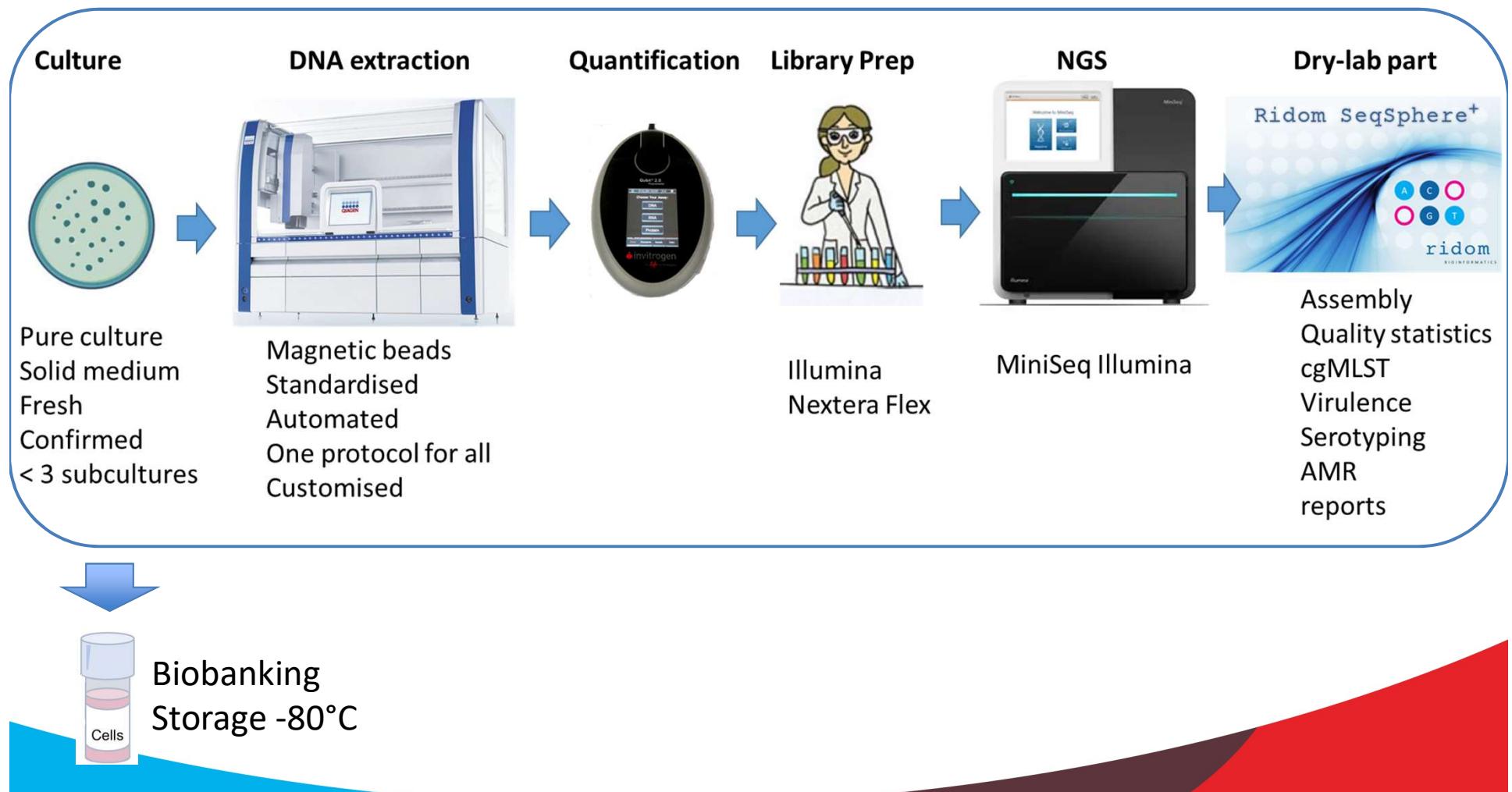
Foodborne pathogens: integrated surveillance – One Health concept
≈1500 isolates/year



Foodborne Surveillance – *Salmonella* – *Listeria* mono.– STEC – *Campylobacter* – *Yersinia enterocolitica* – One Health approach

....+ Staph aureus

Overview of the sequencing workflow





Pilot study

The complete resistance profile of the outbreak monophasic *Salmonella* Typhimurium strain linked to chocolate products.

Genomics tells us the story of the outbreak monophasic *Salmonella* Typhimurium strain linked to chocolate products.



Case definition

The agreed European Union (EU) case definition for confirmed cases was laboratory-confirmed monophasic *S. Typhimurium* with symptom onset on or after 1 October 2021 and belonging to the same five SNP single linkage cluster by SNP typing or cases who clustered within five allelic differences of another confirmed outbreak strain by core genome multilocus sequence typing (cgMLST) analysis or shared the same HC5_296366 by the Enterobase HierCC scheme [1]. This definition therefore depended on the whole genome sequencing (WGS) methodology used at the national level in each country (i.e SNP typing or cgMLST analysis). Probable cases were those with laboratory confirmation of monophasic *S. Typhimurium* with symptom onset on or after 1 October 2021 and phenotypic antimicrobial resistance (AMR) results consistent with the outbreak strain or a multilocus variable number tandem repeat analysis (MLVA) profile 3-11-14-NA-0211.

Source: Euro Surveill. 2022 Apr;27(15):2200314.



In silico definition of the AMR profile (cluster 1)

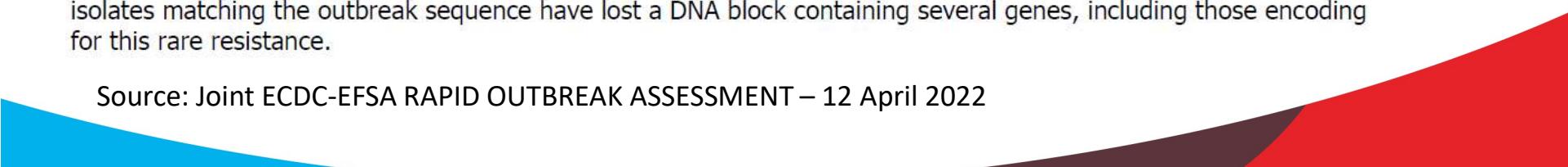
The outbreak strain is **resistant** to six families of antibiotics:

- penicillins (*blaT_{EM-1}*);
- aminoglycosides (streptomycin, spectinomycin, kanamycin, and gentamycin (gene combination varies); strA-strB, aac(6')-Ia, aac(3)-IIId, aph(6)-Id, aadA-2, aadA-8b, aadA-12, aadA-15 and aadA-17;
- phenicols (*cmlA1*, *floR*);
- sulfonamides (sul2 with some strains having an additional sul3 gene);
- trimethoprim (*dfrA12*); and
- tetracyclines (*tetA* and *tetM*).

In addition, some strains contained the *Inu(F)* gene encoding resistance to **lincosamide**, but this could not be confirmed phenotypically.

The resistance to aminoglycosides, phenicols, and trimethoprim is rare in monophasic *S. Typhimurium* and could therefore be used for screening of probable cases. However, France has noted that a minor proportion of their isolates matching the outbreak sequence have lost a DNA block containing several genes, including those encoding for this rare resistance.

Source: Joint ECDC-EFSA RAPID OUTBREAK ASSESSMENT – 12 April 2022



Overview of the number of cases

Update on June 3 th 2022 – ECDC website « Threats and outbreaks of salmonellosis »

Table 1. Number of confirmed and probable cases of monophasic S. Typhimurium in the EU/EEA, the United Kingdom, Canada, Switzerland and the United States, as of 3 June 2022

Country	Confirmed cases	Probable cases	Total number of cases
Austria	14	0	14
Belgium	52	14	66
Denmark	4	0	4
France	118	0	118
Germany	30	5	35
Ireland	17	0	17
Italy	1	0	1
Luxembourg	2	0	2
Netherlands	3	0	3
Norway	1	0	1
Spain	2	3	5
Sweden	4	0	4
Total EU/EEA	248	22	270
United Kingdom	122	0	122
Total EU/EEA and UK	370	22	392
Canada	4	0	4
Switzerland	48	0	48
United States	1	0	1
Total	423	22	445



This outbreak is characterised by a high proportion of hospitalised (about 40%) cases and some cases with severe clinical symptoms such as bloody diarrhoea

In silico characterization of strains from cluster 1



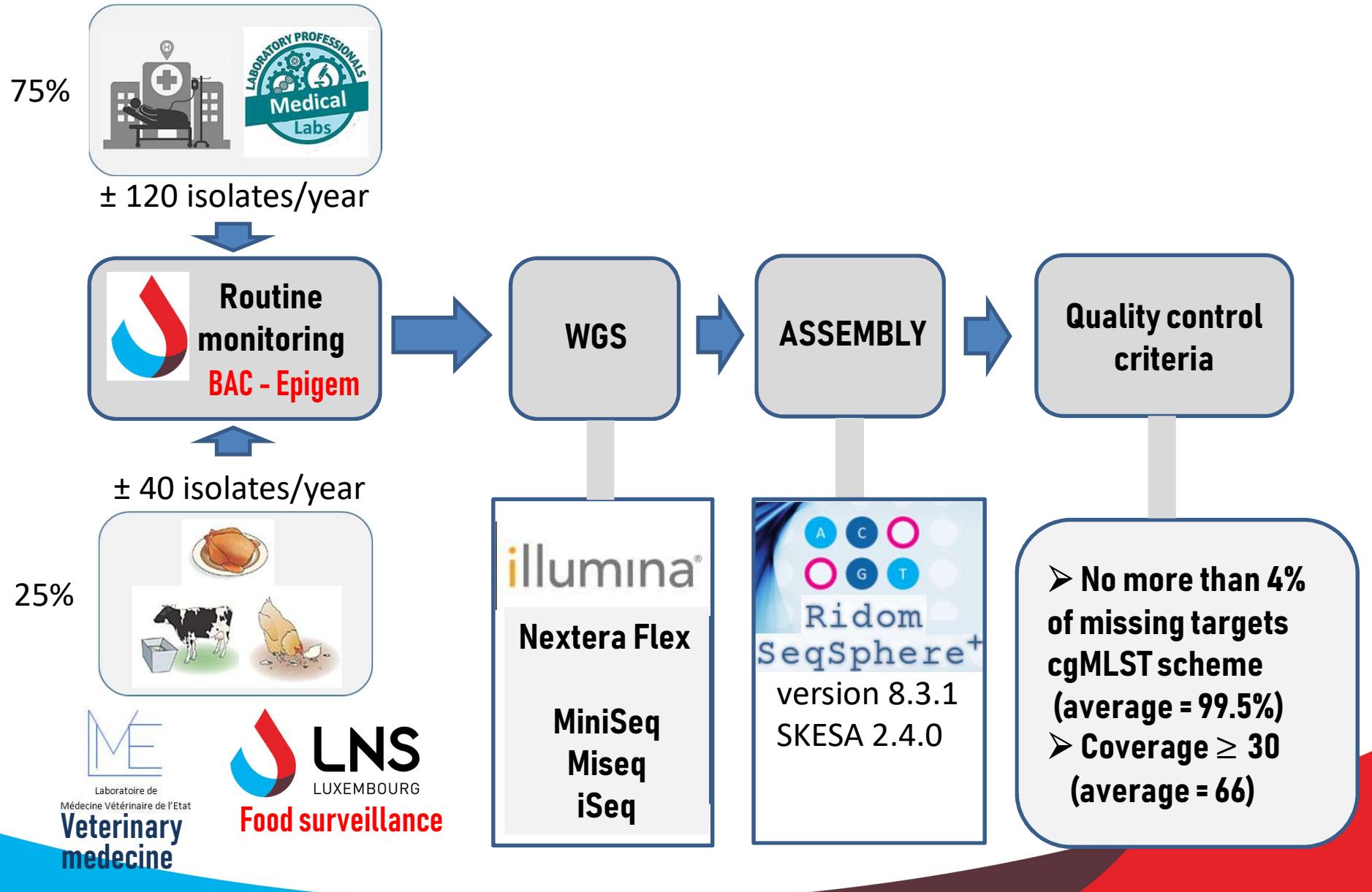
- In Luxembourg:
2 human cases linked to the outbreak = 2 strains from cluster 1
- Identification of ***qacL gene*** encoding reduced susceptibility to a variety of guanidinium including the most notable quaternary ammonium compounds (QACs)



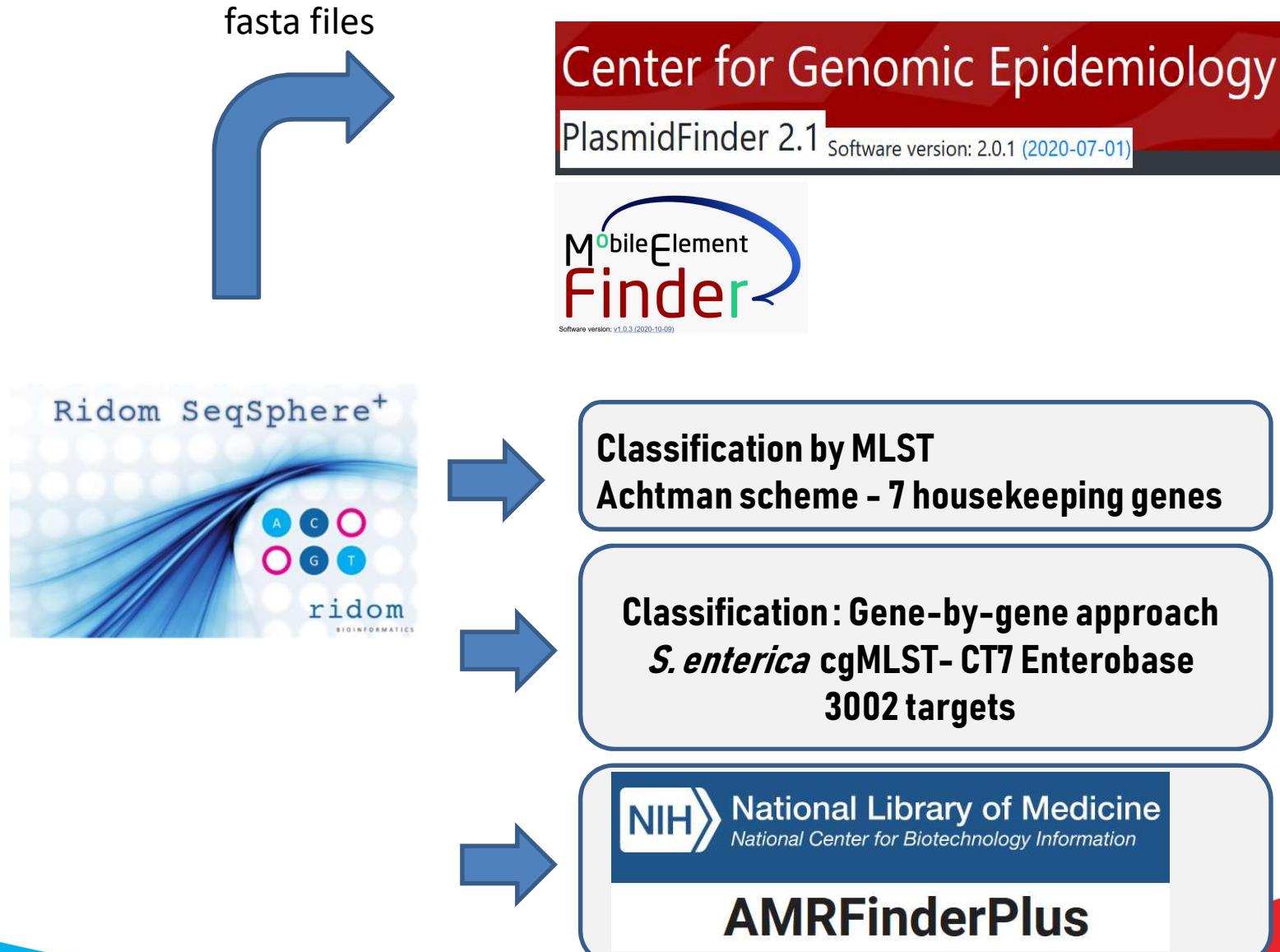
key ingredient in many cleaning and disinfection products

- Frequency of this accessory gene in our collection of S. Typhimurium/monophasic Typhimurium isolates?

Overview of the *Salmonella* workflow – Genomic surveillance in LU

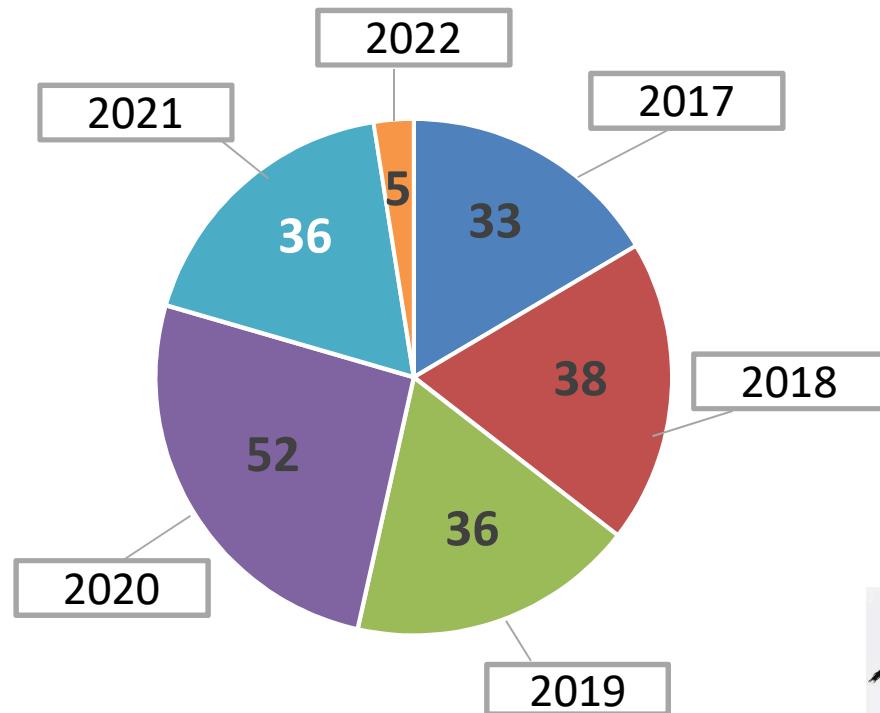


Quick overview of pipelines /bioinformatic tools

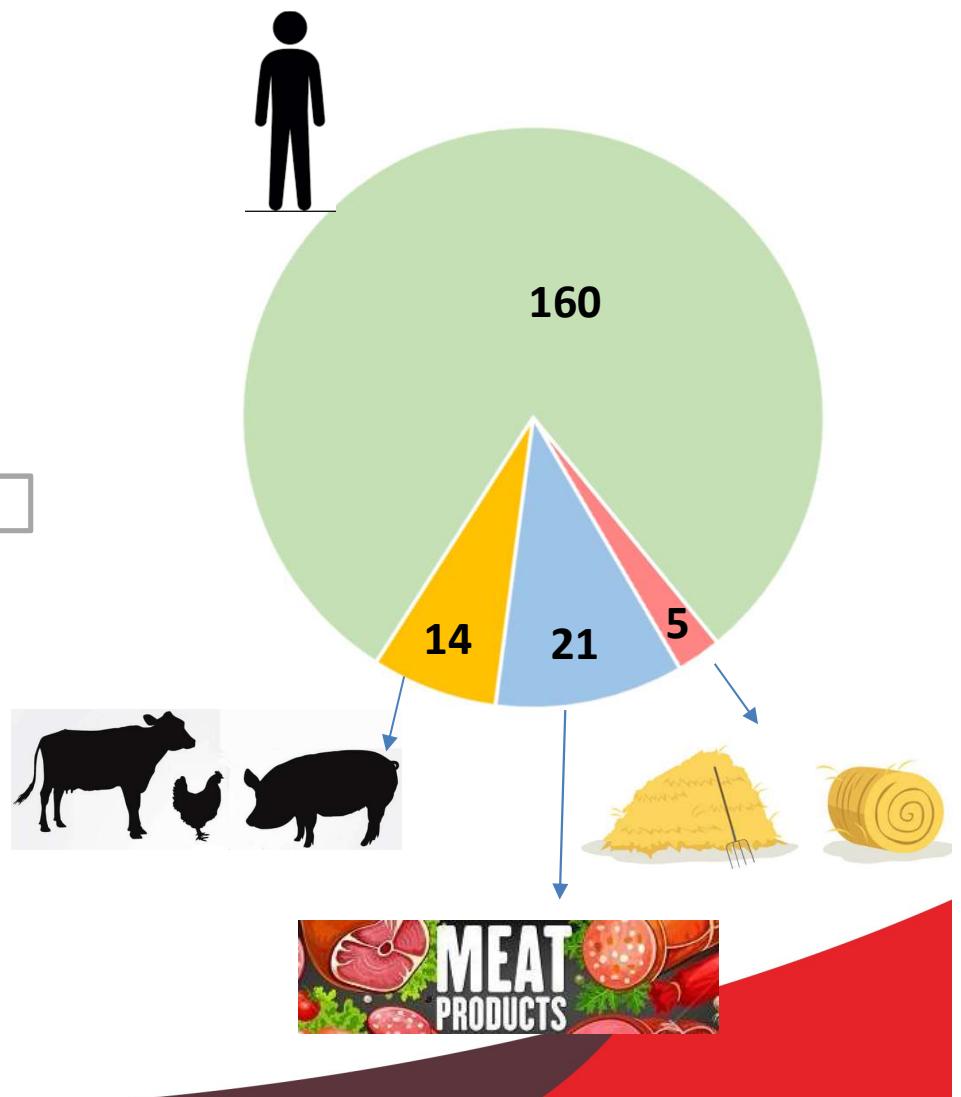


Collection of 200 strains *Salmonella* Typhimurium/ monophasic S. Typhimurium (ST19-ST34)

Isolated between 2017-2022 (5 years +)



One Health



Resistance profiles in silico associated to *qacL*

Sources	Classes of antibiotics							Biocides	Metals				Plasmids
 A								<i>qacL</i>					IncFIB(AP001918)
 F								<i>qacL</i>					IncFIB(AP001918)
 H								<i>qacL</i>					IncFIB(AP001918)
								<i>qacL</i>					IncN -
								<i>qacL</i>					Col8282 - IncHI2 - IncI1-I(Alpha)
								<i>qacL</i>					IncFIB(AP001918)
								<i>qacEΔ1</i>					IncFIB(AP001918) - IncFIB(S)- IncFII(S)
								<i>qacL</i>					IncFIB(S)-IncFII(S)
								<i>qacL</i>					IncFIB(AP001918)
								<i>qacL</i>					IncFIB(AP001918)
Outbreak O	O												

Legend for Classes of antibiotics:

- Penicillin (light green)
- Aminoglycosides (light blue)
- Phenicols (orange)
- Sulfonamides (pink)
- Trimethoprim (yellow)
- Tetracycline (red)
- Lincosamide (light green)
- Macrolide (blue)

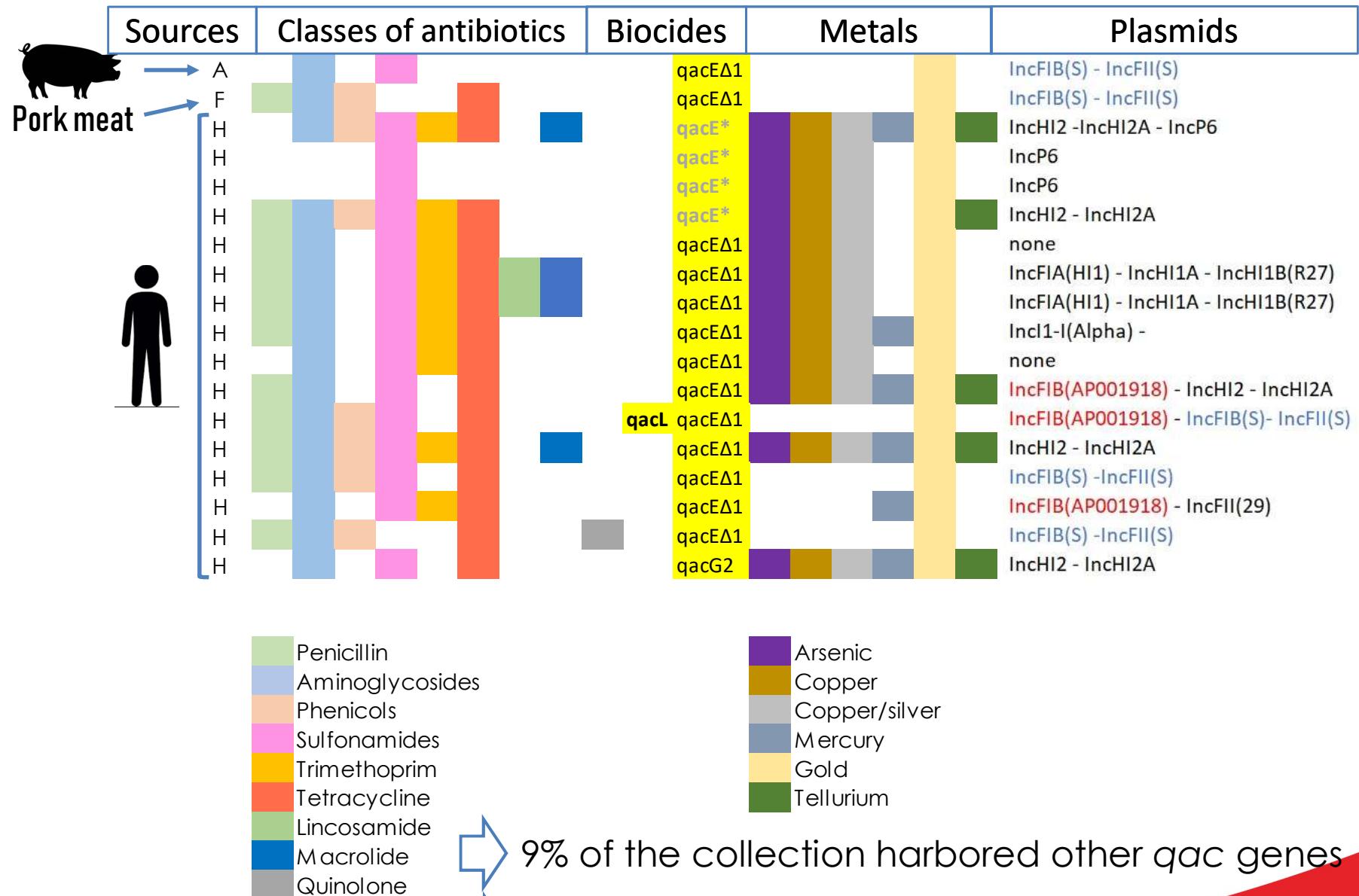
Legend for Metals:

- Arsenic (purple)
- Copper (gold)
- Copper/silver (grey)
- Mercury (dark blue)
- Gold (yellow)
- Tellurium (green)

→ 9/200 strains (unique cgMLST types) over the 5 past years harboured *qacL* => **4,5%**

→ 5 to 7 classes of antibiotics R + reduced susceptibilities for 5 heavy metals

Resistance profiles in silico associated to other Biocides (qacEΔ1)



Conclusions & Perspectives



- In silico characterization of the outbreak strain from cluster 1 highlighted its resistance profile:
 - 6+ classes of antibiotics
 - reduced susceptibilities to quaternary ammonium compounds (*qacL*)
 - reduced susceptibilities to 5 heavy metals
- *qacL* gene quite uncommon in *S. Typhimurium*/monophasic *Typhimurium* \approx 4.5% of strains
=> strong interest to analyse data through integrated surveillance
- Environmental persistence of this strains linked to biocide & heavy metals “resistance” ?
- Role of biofilm ?
- Interest for the surveillance to characterize strains in their whole profiles of resistance: antibiotics genes/mutations & genes conferring an advantage to face environmental conditions
- Correlation phenotype – genotype? Perspective to test the strains with various concentration levels of biocide (quaternary ammonium)



References

Larkin L, Pardos de la Gandara M, Hoban A, Pulford C, Jourdan-Da Silva N, de Valk H, Browning L, Falkenhorst G, Simon S, Lachmann R, Dryselius R, Karamehmedovic N, Börjesson S, van Cauteren D, Laisnez V, Mattheus W, Pijnacker R, van den Beld M, Mossong J, Ragimbeau C, Vergison A, Thorstensen Brandal L, Lange H, Garvey P, Nielsen CS, Herrera León S, Varela C, Chattaway M, Weill FX, Brown D, McKeown P. Investigation of an international outbreak of multidrug-resistant monophasic *Salmonella* Typhimurium associated with chocolate products, EU/EEA and United Kingdom, February to April 2022. Euro Surveill. 2022 Apr;27(15):2200314. doi: 10.2807/1560-7917.ES.2022.27.15.2200314. PMID: 35426359; PMCID: PMC9012091.

Joint ECDC-EFSA RAPID OUTBREAK ASSESSMENT – Multi-country outbreak of monophasic *Salmonella* Typhimurium sequence type (ST) 34 linked to chocolate products- 12 April 2022 .

https://www.ecdc.europa.eu/sites/default/files/documents/ROA_monophasic-S-Typhimurium-ST34-linked-to-chocolate_2022-00014-final_UK.pdf

15 July update: Monophasic *Salmonella* Typhimurium outbreak linked to chocolate products- 15 Jul 2022
<https://www.ecdc.europa.eu/en/news-events/15-july-update-monophasic-salmonella-typhimurium-outbreak-linked-chocolate-products>

Thank you for your attention and special thanks to...

Maria Pardos de la Gandara – Institut Pasteur FR– NRC E. coli – Shigella – Salmonella

Johanna Takkinnen & Therese Westrell - ECDC - FWD

Thank you to all active partners involved in our integrated surveillance in Luxembourg!

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Catarina Martins



Manon Bourg
Alexandra Schoos
Dominique Claude



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