









The 2017-2018 listeriosis outbreak, South Africa: situation update



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Epidemiology of listeriosis

- Sporadic cases occur worldwide, mostly in persons at high risk
 - Estimated 2 4 cases per million population per year in developed countries
- Outbreaks associated with contaminated food products increasingly common
- Nosocomial outbreaks also reported

Surveillance for listeriosis cases

- Requirements for surveillance
 - Diagnostic algorithm: <u>case definition</u>, <u>diagnostic tests</u>
 - Staff members
 - Sampling frame
 - Competent laboratory
- Surveillance system should allow
 - Timely data processing
 - Results and action
 - Interpretation
 - Facilitating appropriate action based on surveillance results

Case definition

 Only those cases that meet the 'confirmed' case definition are included; there are no probable, possible, or suspected cases.

Confirmed case:

- A person who has clinical features in keeping with listeriosis (symptoms and signs depend on site and severity of disease*)
 and
- laboratory confirmation of Listeria monocytogenes (by culture or PCR).

^{*}Types of disease include: febrile gastroenteritis; pregnancy-associated (maternal listeriosis, fetal listeriosis, neonatal listeriosis); bacteraemia; central nervous system disease (including meningitis, meningoencephalitis, rhombencephalitis, cerebral abscesses); other focal infections (e.g. mycotic aneurysms, septic arthritis, endocarditis, endophthalmitis, and osteomyelitis).

Passive laboratory-based surveillance

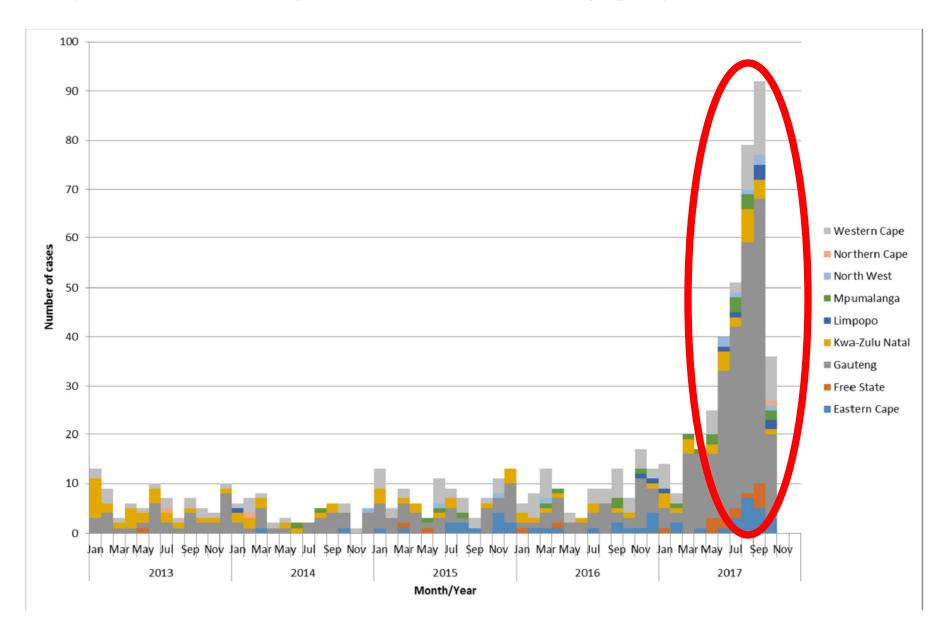
- <u>Laboratory-confirmed cases</u> are reported, then documented
- Epidemiologic analysis of lab-confirmed cases performed
 - Establish trends (time, person, place)
 - Seasonality?
 - Detect unusual signals
 - Time
 - Person
 - Place

Case investigation

- Relies on timely reporting of laboratory-confirmed cases to health department
- Triggers case investigation
 - Patient interview: epidemiologic information and food consumption history
 - Standardised CIF developed by CED, NICD
 - Additional clinical data
 - Visit to patient's home: sample food from refrigerator, freezer
 - Awareness for other cases (?cluster)

- Depending on case investigation may trigger further traceback investigation
 - Case part of a cluster
 - Food consumption history provides signal

Epidemic curve of laboratory-confirmed cases of Listeria monocytogenes, South Africa, 2013-2017



Situation update on listeriosis outbreak, issued 23 April 2018

- As of 17 April 2018, a total of 1 019 laboratoryconfirmed listeriosis cases has been reported to NICD since 01 January 2017
- Where age was reported (n=993), ages range from birth to 93 years (median 18 years) and 42% are neonates aged ≤28 days
- Most cases have been reported from Gauteng Province (59%,579/1 019), followed by Western Cape (13%, 127/1 019) and KwaZulu-Natal (7%, 73/1 019) provinces
- Final outcome data is available for 69% (698/1019) of cases, of which 28% (199/698) died.



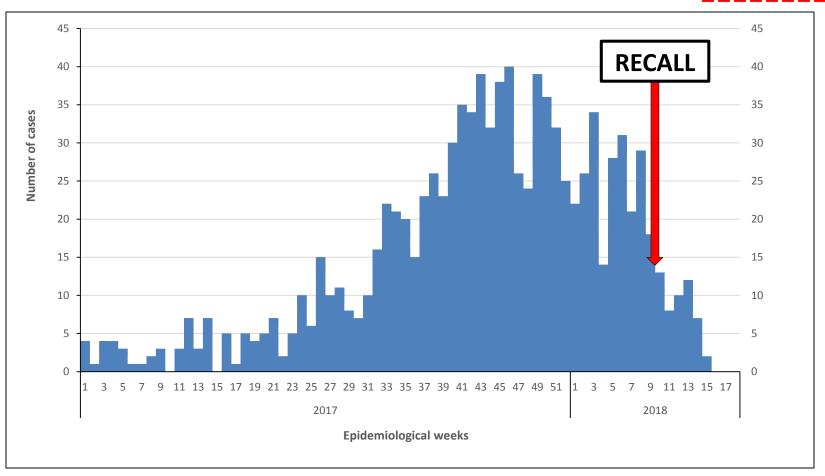


Figure 1: Epidemic curve of laboratory-confirmed listeriosis cases by epidemiological week and date of sample collection, South Africa, 01 January 2017 to 17 April 2018 (n=1 019)

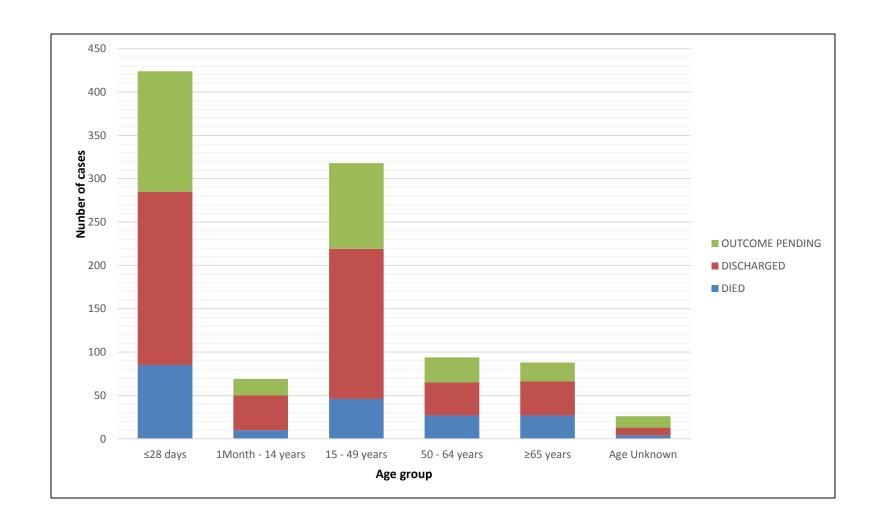
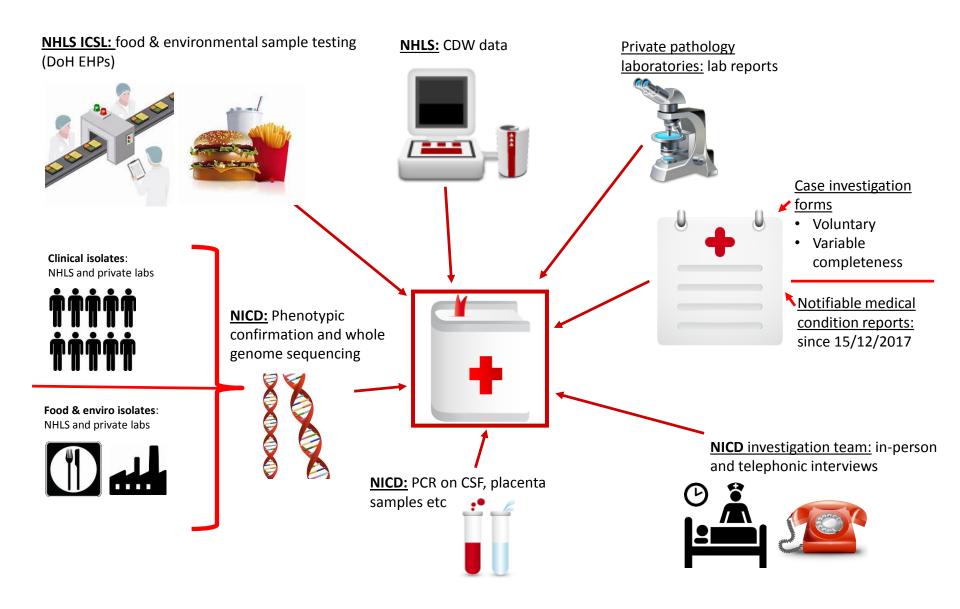


Figure 2: Age distribution and outcome of laboratory-confirmed listeriosis cases, South Africa, 01 January 2017 to 17 April 2018 (n=1 019)

Table 1. Number of laboratory-confirmed listeriosis cases and deaths by province, where outcome data is available:

Province	Outcome available (%)	Number of deaths (%)	Total cases (%)
Gauteng	370 (62.0)	105 (28.4)	597 (58.6)
Western Cape	113 (89.0)	29 (25.7)	127 (12.5)
KwaZulu-Natal	47 (64.4)	18 (38.3)	73 (7.2)
Mpumalanga	45 (95.7)	11 (24.4)	47 (4.6)
Limpopo	33 (63.5)	7 (21.2)	52 (5.1)
Eastern Cape	30 (56.6)	11 (36.7)	53 (5.2)
Free State	30 (85.7)	8 (26.7)	35 (3.4)
North West	25 (86.2)	7 (28.0)	29 (2.8)
Northern Cape	5 (83.3)	3 (60.0)	6 (0.6)
Total	698 (68.5)	199 (28.5)	1 019

Gathering and analysing listeriosis outbreak investigation data

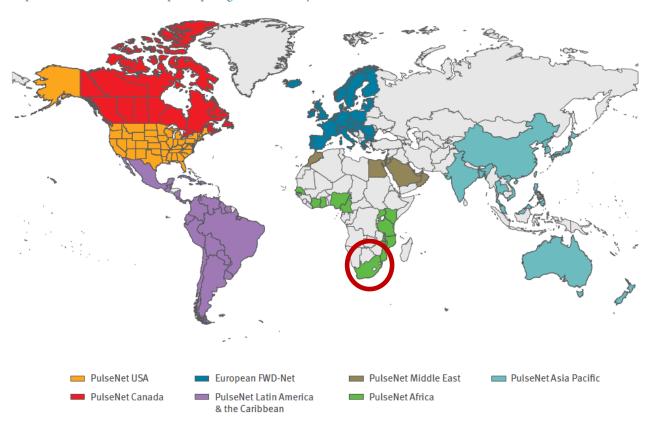


PulseNet International: Vision for the implementation of whole genome sequencing (WGS) for global foodborne disease surveillance

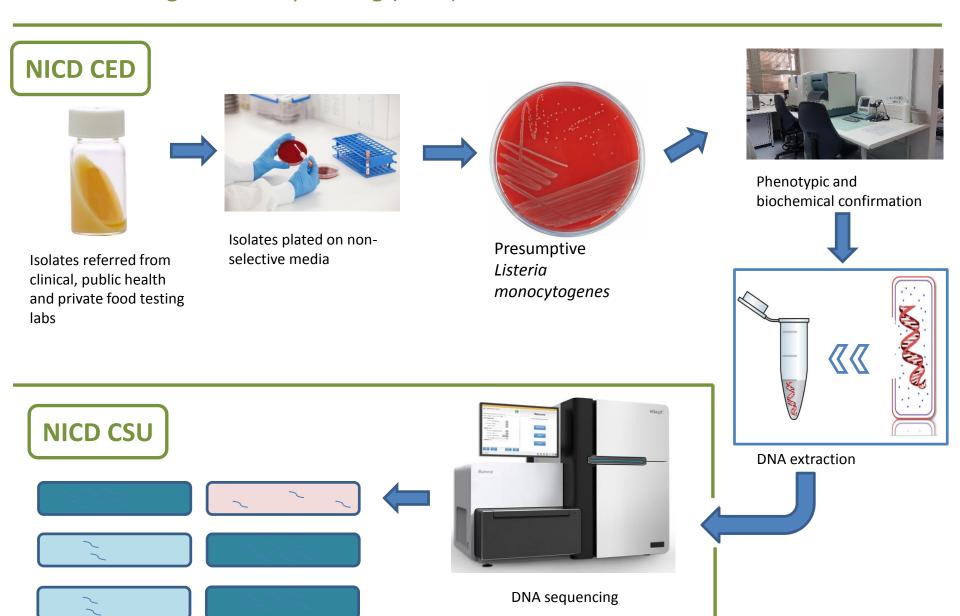
Celine Nadon^{1,2}, Ivo Van Walle^{2,3}, Peter Gerner-Smidt⁴, Josefina Campos⁵, Isabel Chinen⁵, Jeniffer Concepcion-Acevedo⁴, Brent Gilpin⁶, <u>Anthony M. Smith</u>⁷, Kai Man Kam⁸, Enrique Perez⁹, Eija Trees⁴, Kristy Kubota¹⁰, Johanna Takkinen³, Eva Møller Nielsen¹¹, Heather Carleton⁴, FWD-NEXT Expert Panel¹²

Euro Surveill. 2017;22(23):pii=30544.

FIGURE 1
Map of PulseNet International participating countries, May 2017



Whole genome sequencing (WGS): The Listeria detective's ultimate tool

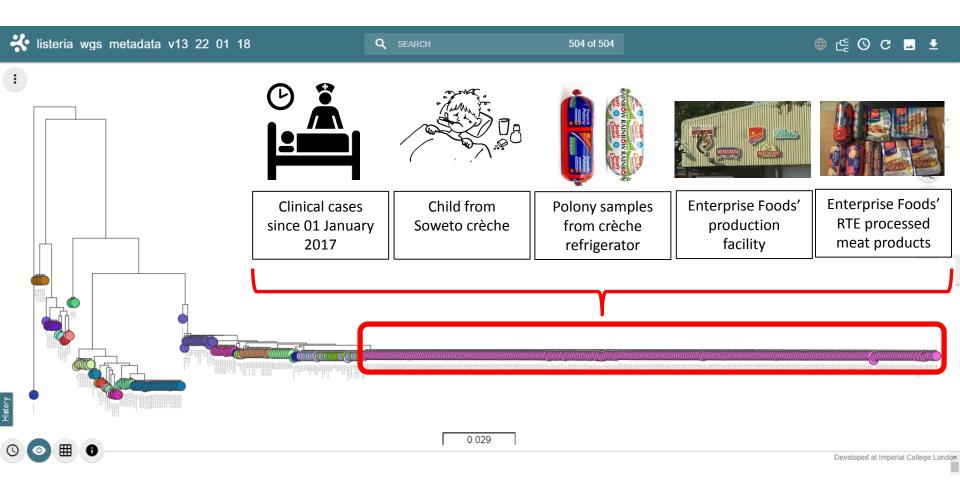


Whole genome sequence analysis

Solving the outbreak

- 12 January 2018: a cluster of febrile gastroenteritis in 9 children attending a crèche in Soweto (Johannesburg), who all consumed polony sandwiches
 - L. monocytogenes ST6 (outbreak strain) confirmed in one case
 - L. monocytogenes ST6 (outbreak strain) detected in food samples (two different polony brands produced by different manufacturers) collected from refrigerator at crèche
- Investigation of polony manufacturer (Enterprise Foods' production facility in Polokwane)
 - L. monocytogenes ST6 (outbreak strain) detected in 28 environmental and product casing swabs
 - L. monocytogenes ST6 (outbreak strain) also detected in several RTE processed meat products manufactured at the Polokwane facility

Phylogenetic tree of WGS



Distance and topology calculated by RaXML; indicates ST6 isolates of 92% clinical cases identical, and Enterprise Foods' production facility environmental and product isolates also identical; i.e. confirms exposure to common contaminated food item/s produced at Enterprise Foods' production facility in Polokwane

Acknowledgements

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- Several private sector food microbiology laboratories
- WHO
- US CDC
- Institut Pasteur