

# Three epidemics of multidrug resistant invasive Salmonella disease in Blantyre, Malawi

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### Malawi Liverpool Wellcome Trust Clinical Research Programme at Queen Elizabeth Central Hospital













# LSTM

### Longitudinal bacteraemia surveillance since 1998

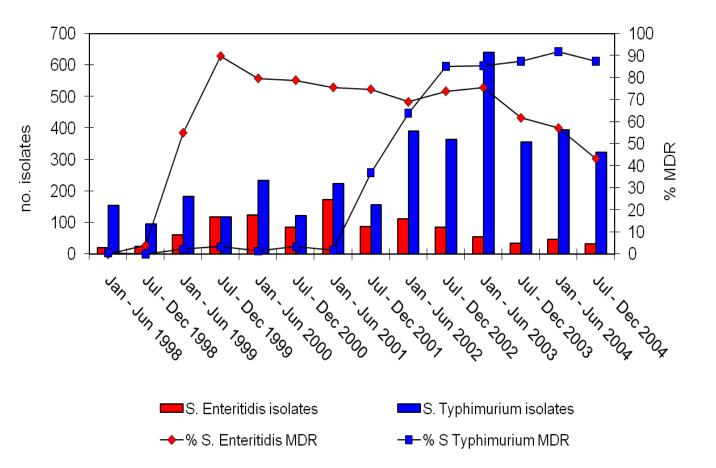




# Relationship between MDR and incidence of NTS LSTM

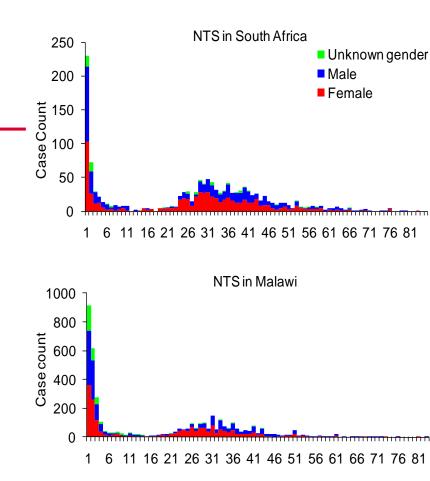






## **Risk factors for iNTS disease**

- Bimodal age distribution
- Adults with severe HIV
- Children with
  - HIV
  - Malaria
  - Malnutrition



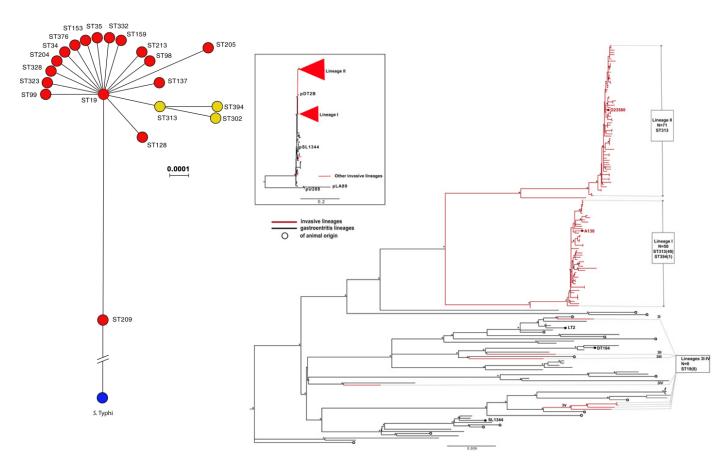


## Susceptible host or novel pathogen?





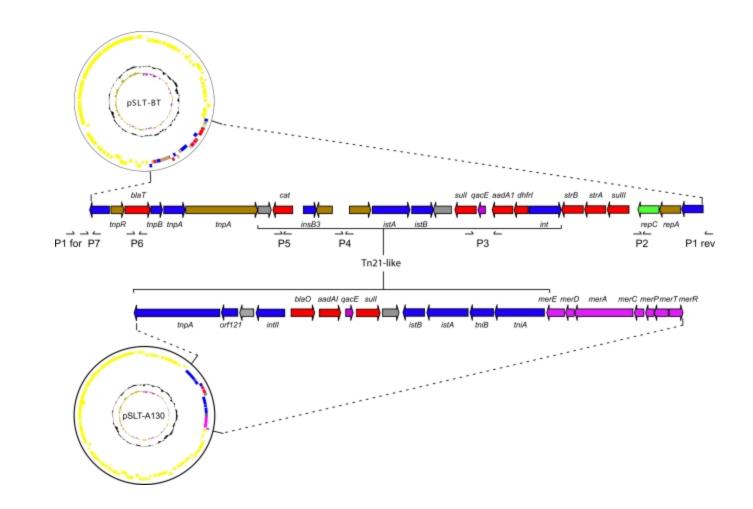




Msefula, Kingsley Gen Res 2009 Okoro et al Nat Gen 2012

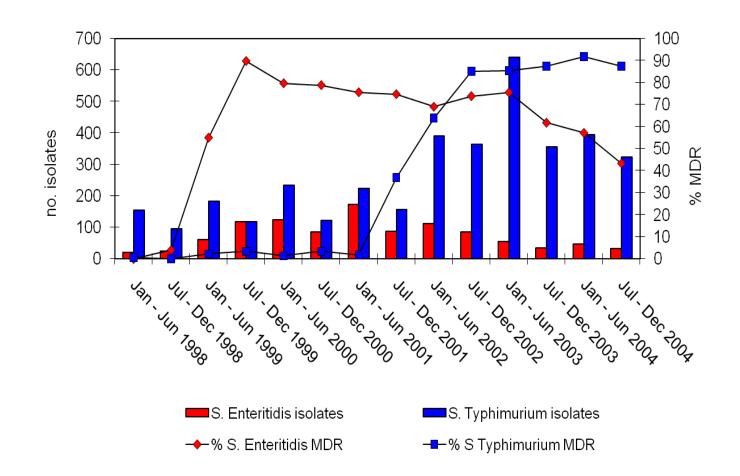


#### **Novel MDR plasmid associated with ST313**



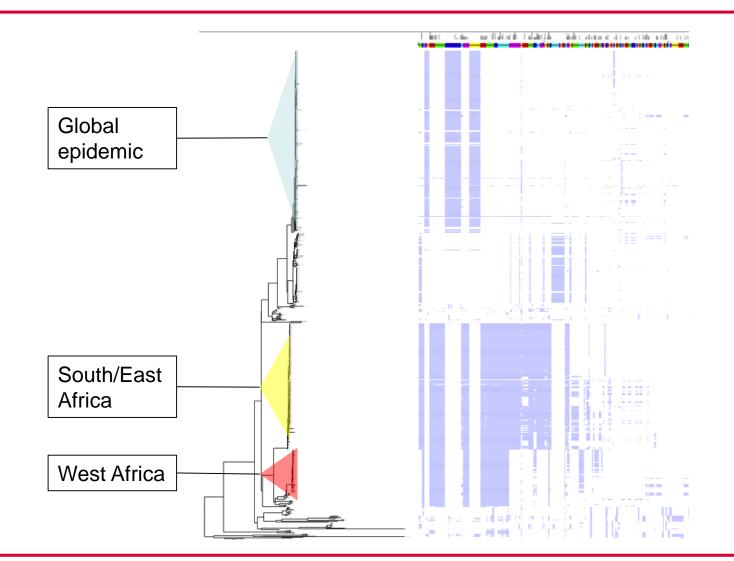
# **Prior to the epidemic of S. Typhimurium ST313, there was an epidemic of S. Enteritidis**





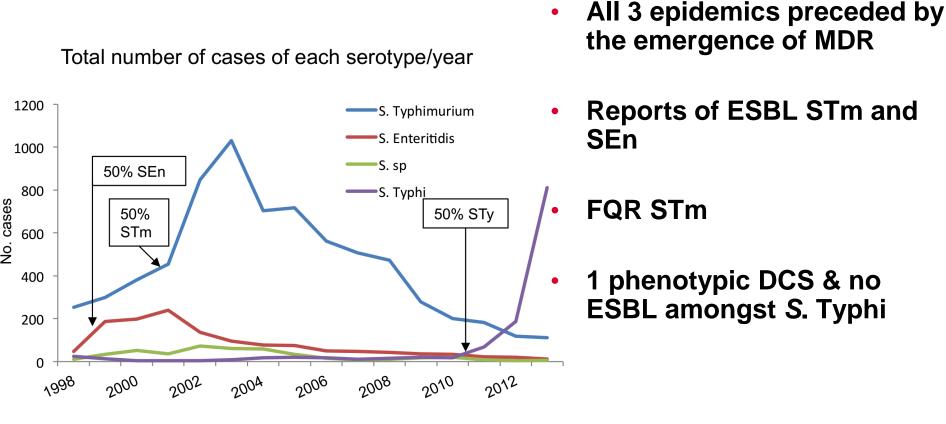
#### Novel clades of *S*. Enteritidis restricted to sub-Saharan Africa with MDR-plasmids





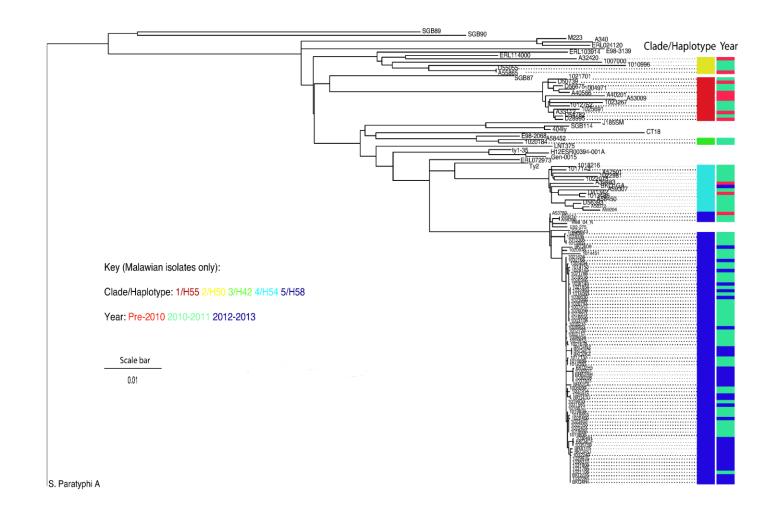
# NTS remains a prominent pathogen, but MDR S. Typhi has now emerged







### H58-S. Typhi, but with chromosomal MDR locus



### **Clinical features and complications of iNTS disease**



	2001	2009-11
Recruited	100	63
Inpatient mortality	47%	11%
1 year mortality	77%	38%
HIV positive	99%	98%
Recurrence	43% (19/44)	5%(3/56)
Multiple recurrences	11% (5/44)	2%(1/56)



# Adults (n= 77)

- 16% HIV infected
- 5% severe/complicated
- 1% perforation
- 4% mortality

# Children (n=330)

- HIV data pending
- 21% complicated
- 2% perforation
- 2% mortality



- Decline has leveled off
- Still a major pathogen
- iNTS disease likely to be responsible for approximately double the number of deaths caused by Typhoid fever in Blantyre, 2014



- 3 epidemics of Salmonella bloodstream infection since 1999
- Epidemics of iNTS disease associated with novel clades
- Recent emergence of H58-S. Typhi
- Risk factors for iNTS disease and Typhoid fever differ
- BUT all 3 epidemics associated with emergence of MDR
- In-turn driving increased use of cephalosporins and fluoroquinolones
- Reservoirs of pathovars associated with iNTS disease unknown

#### Acknowledgements



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