









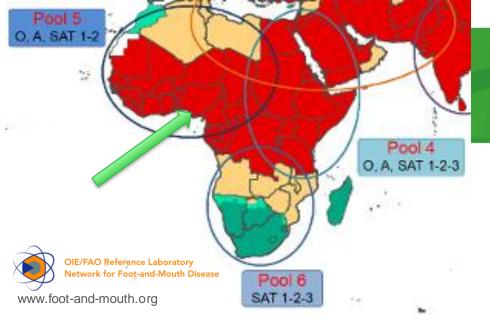
Foot-and-mouth disease in Nigeria

presented by David Lefebvre (Sciensano, Belgium)
in collaboration with
the National Veterinary Research Institute (Nigeria)
and the Plum Island Animal Disease Center (ARS/USDA, USA)

EuFMD OS18

Borgo Egnazia, Italy, October 29th-31st 2018





Nigeria

- 924.000 km²
- 190 million people
- Largest African economy
- Exportation of oil and gas
- Subsistence-oriented agriculture 20 million cattle, 40 million sheep,
 70 million goats, 7 million pigs demand largely exceeds production
- Significant movement of livestock across the borders and inside the country due to importation and nomadic pastoralism
- No systematic surveillance for FMD, only sporadic notifications
- No control program for FMD, vaccines not locally available







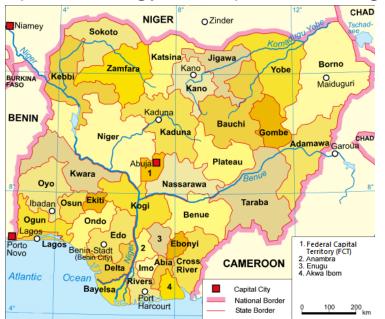




Previous results of the OIE Laboratory Twinning program for capacity building

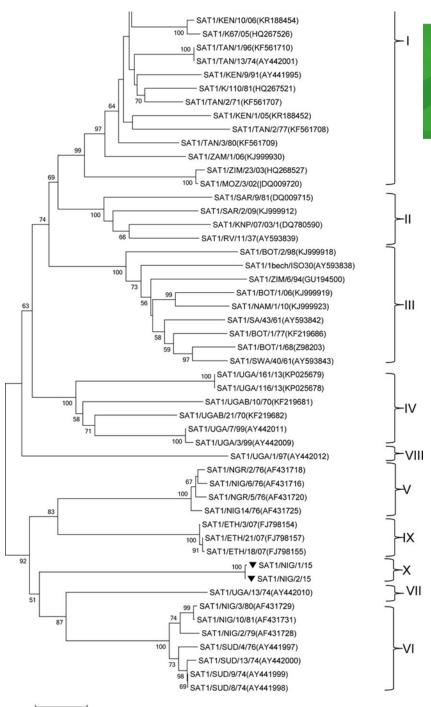
 Detection of FMD virus in 4 provinces of Northern Nigeria during the period 2012-2015: O/WA, O/EA-3, A/Africa/G-IV, SAT1/X, SAT2/VII (Ehizibolo, Haegeman et al., 2017a, Transbound Emerg Dis)

Epidemiology: complex and highly dynamic



- Most prevalent serotype in Plateau State:
- 2013: SAT2/VII
- 2014: O/EA-3
- 2015: A/Africa/G-IV
- From end 2015: SAT1/X





Identification of a new topotype: SAT1/X

| | <u>% NT</u> homology | % AA homolog |
|----------------|-------------------------|-----------------|
| Topotype V | | |
| SAT1/NGR/2/76 | 71 | 79 |
| SAT1/NGR/5/76 | 70 | 79 |
| SAT1/NIG/6/76 | 71 | 79 |
| SAT1/NIG/14/76 | 71 | 79 |
| Topotype VI | | |
| SAT1/NIG/2/79 | 71 | 78 |
| SAT1/NIG/3/80 | 71 | 76 |
| SAT1/NIG/10/81 | 71 | 78 |
| | | |

- Ehizibolo, Haegeman et al., 2017b, Transbound Emerg Dis
- Vandenbussche et al., 2017, Genome Announc



Studies during the period 2016-2018: epithelial samples

- Epithelial samples from suspected FMD-infected cattle were collected in Northern and South-West Nigeria
 - 2016: 45 samples from 3 States and 3 archived samples from 2014 were analyzed in PIADC
 - 2017: 44 samples from 7 States were analyzed in Sciensano
 - 33 archived samples (2012 and 2014) from the Kachia Grazing Reserve (KGR) were analyzed in Sciensano











Studies during the period 2016-2018: serum samples

- Archived serum samples were analyzed in Sciensano
 - Serosurveys:
 - 300 samples from sheep and goat (2014)
 - 38 samples from wildlife (2009 end 2014)
 - 1220 samples from cattle and 160 samples from sheep from KGR (2012)
 - FMD outbreaks:
 - 59 samples from diseased cattle from KGR (2012 2014)











Results from epithelial samples

- Samples from 2014 (3) and 2016 (45):
 O/WA, O/EA-3, A/Africa/G-IV, SAT1/X, SAT2/VII
- Samples from 2017 (44):
 - O/EA-3, A/Africa/G-IV, SAT2/VII
 - In a single State outbreaks with 2 or 3 different serotypes observed in only 2 or 3 months of time
 - A single serotype isolated at the same time in remote States
- Samples from KGR (33):
 - 2012: SAT2/VII
 - 2014: O/EA-3











Results from serum samples

Serosurveys:

- Lower prevalence of antibodies against FMDV in small ruminants compared to cattle and wildlife
 - 16.3 21.7% vs. 30.6 44.7%
- Most small ruminants are seropositive for a single serotype while more wildlife and cattle are seropositive for 2 or more serotypes
 - 72.3% mono-specificity vs. 46.7 58.0%
- Until 2014 serological evidence for circulation of O, A and SAT2
- FMD outbreaks in Kachia Grazing Reserve:
 - Until 2012 serological evidence for circulation of O and A
 - Since 2012 outbreaks also serological evidence for SAT2













Discussion

- No isolation of O/WA and SAT1/X in 2017
 - Outcompeted by O/EA-3, A/Africa/G-IV and SAT2/VII?
 - No access to samples from States in which these topotypes are circulating?
- No evidence for circulation of SAT1/X in Nigeria until at least 2014
 - Assumption of a new introduction into Nigeria in 2015 still stands
 - No data available from neighboring countries, particularly Niger and Chad,
 that act as an access road for the importation of live animals
- Lower prevalence of FMD in small ruminants, even at household level
 - Less susceptible to infection with FMD virus?
 - Lower and thus less detectable antibody levels?
 - Younger age structure of the herd due to larger number of offspring?













Acknowledgements



Laboratory Twinning program for capacity building (Sciensano and NVRI)



FASTA project

Co-authors:



NVRI: Hussaini Ularamu, Yiltawe Wungak, David Lazarus, Y.J. Atuman, A.G. Ardo, Chika Nwosuh, David Ehizibolo



Ahmadu Bello University (Zaria, Nigeria): Jarlath Umoh, Emmanuel Okolocha, Haruna Kazeem



Andy Haegeman, Annebel De Vleeschauwer, Ilse De Leeuw, Frank Vandenbussche, Steven Van Borm, Kris De Clercq



PIADC: Ian Fish, Steven Pauszek, Carolina Stenfeldt, Jonathan Arzt







Any questions or remarks?









