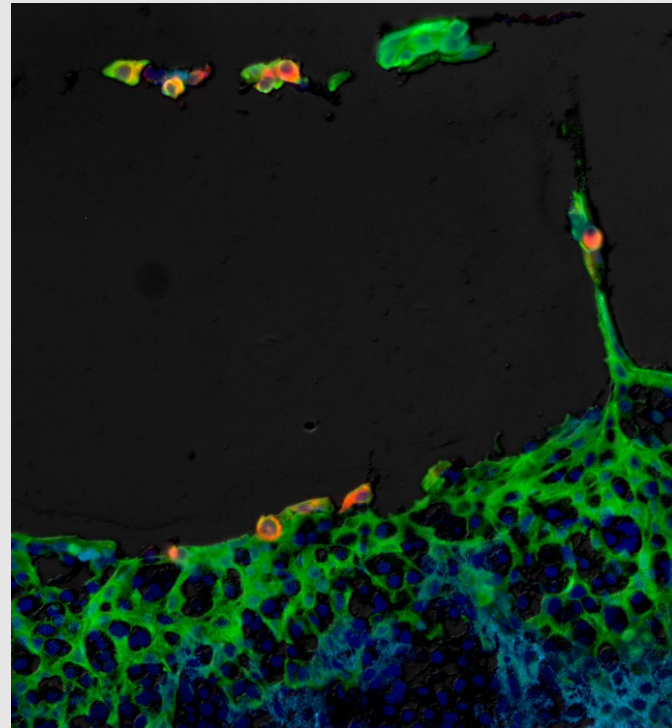


The role of asymptomatic carriers in FMD ecology; unifying knowledge from controlled laboratory experiments and field studies

Jonathan Arzt, Luis Rodriguez, Barbara Brito, Carolina Stenfeldt

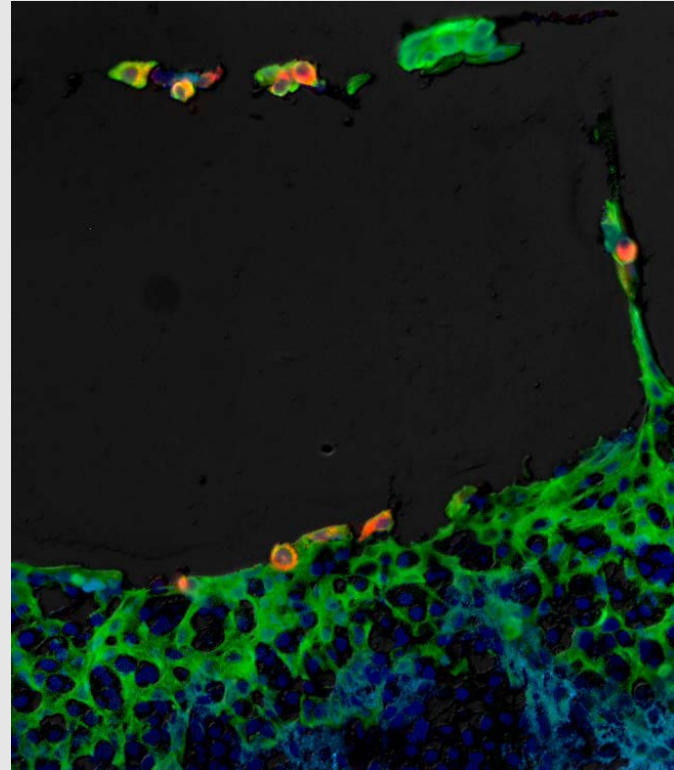


Jonathan Arzt
Veterinary Medical Officer
Foreign Animal Disease Research Unit
ARS, USDA
Plum Island Animal Disease Center





field studies



laboratory experiments

The FMDV Carrier State; early landmark papers

- ***Van Bekkum et al., 1959***

- Recovery of infectious FMDV in OPF (saliva) up to 8 months after recovery from clinical FMD
- Similar subclinical persistence found in vaccinated cattle
- Reports that FMDV carriers do not transmit infection further

- ***Sutmoller and Gaggero, 1965***

- Standardization of probang sampling technique
 - Although, similar approach as used by van Bekkum

- ***Burrows 1966***

- Isolation of infectious FMDV from nasopharyngeal tissues of persistently infected cattle

- ***Sutmoller et al 1968***

- Definition of carriers at 28dpi

First description of carrier state;

Van Bekkum et al., 1959

Observations on the carrier state of cattle exposed to foot-and-mouth disease virus

by

J. G. VAN BEKKUM, H. S. FRENKEL, H. H. J. FREDERIKS
and S. FRENKEL.

(Centraal Diergeneeskundig Instituut, Afd. Amsterdam)

Tijdschrift Voor Diergeneeskunde, Aflevering 20, Deel 84, 15 Okt. 1959.

At the XVth Veterinary Congress held in Madrid in 1959, Schang gave as his opinion, that virus carriers are of no importance in the Argentine. According to Fogedby a comparable situation exists in Scandinavia. The available evidence at most seems to warrant the conclusion, that only a small proportion of recovered individuals may excrete the agent (Olitsky c.s., 1928).

In 1955 Dijkstra, basing his opinion on an epizootiologic study of the disease in the Dutch province of Friesland, reached the conclusion, that in cattle either virus excretion is not continuous, or the amount of virus liberated is so small.

Experimental studies

Advantages

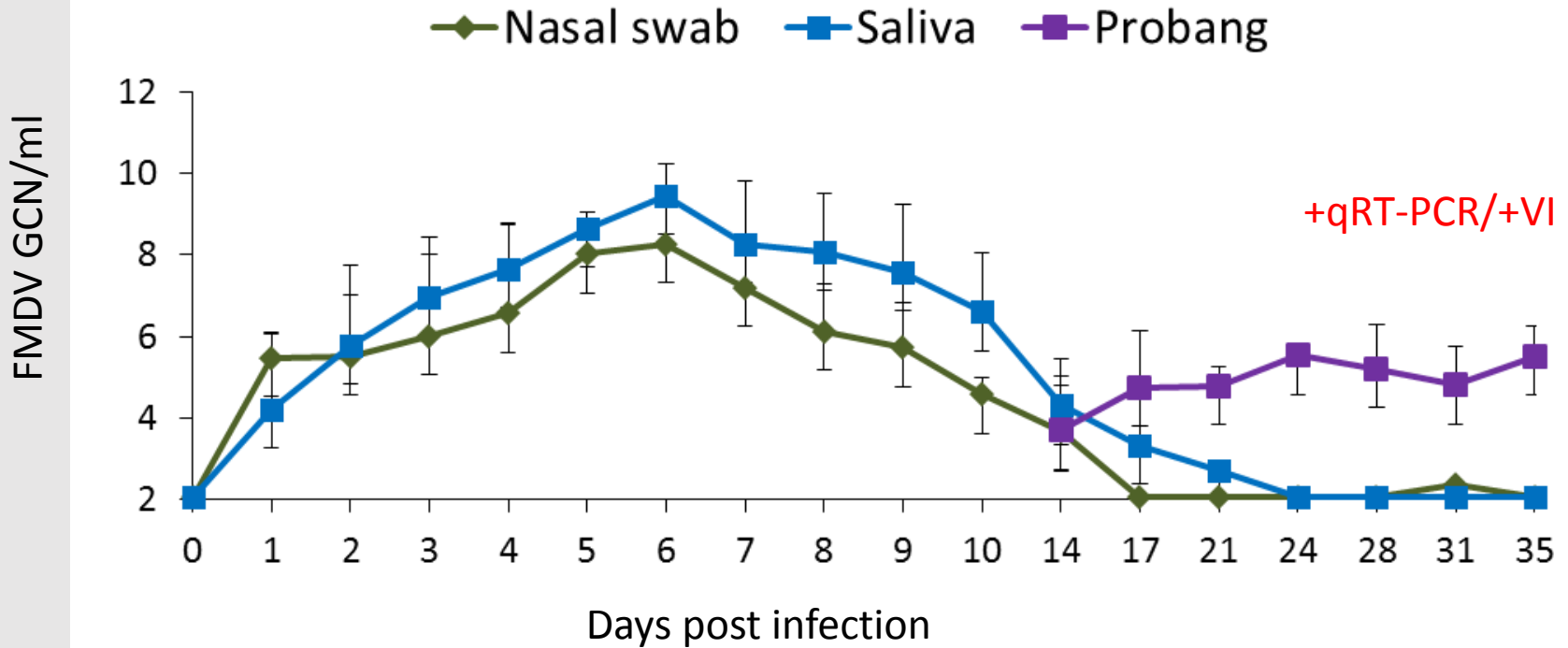
- Prospective study design
- Controlled environment
 - Known virus and exposure conditions
 - Strategic & intensive sampling
 - Possible to reduce/control confounding variables
 - No delays in sample processing
 - Reduced operator variability
- Provides detailed knowledge of FMDV pathogenesis

Limitations

Detecting and defining carriers

FMDV “shedding” in oral/nasal swabs and probangs

Non-Vaccinated cattle



← Clinical FMD →

← FMDV persistence →

Stenfeldt et al 2016

AMERICAN SOCIETY FOR MICROBIOLOGY | Journal of Virology | CrossMark

The Foot-and-Mouth Disease Carrier State Divergence in Cattle

Carolina Stenfeldt,^{a,b} Michael Eschbaumer,^{a,b} Steven I. Rekant,^{a,b} Juan M. Pacheco,^a George R. Smoliga,^a Ethan J. Hartwig,^a Luis L. Rodriguez,^a Jonathan Arzt^a

Plum Island Animal Disease Center, Foreign Animal Disease Research Unit, Agricultural Research Service, United States Department of Agriculture, Greenport, New York, USA; Oak Ridge Institute for Science and Education, PIAOC Research Participation Program, Oak Ridge, Tennessee, USA^a

Sutmoller et al 1968

Plum Island Animal Disease Laboratory, Animal Disease and Parasite Research Division, Agricultural Research Service, U.S. Department of Agriculture, Greenport, Long Island, N.Y., U.S.A.

The Epizootiological Importance of Foot-and-Mouth Disease Carriers

I. Experimentally Produced Foot-and-Mouth Disease Carriers in Susceptible and Immune Cattle

By

Paul Sutmoller, John W. McVicar, and George E. Cottral

With 1 Figure

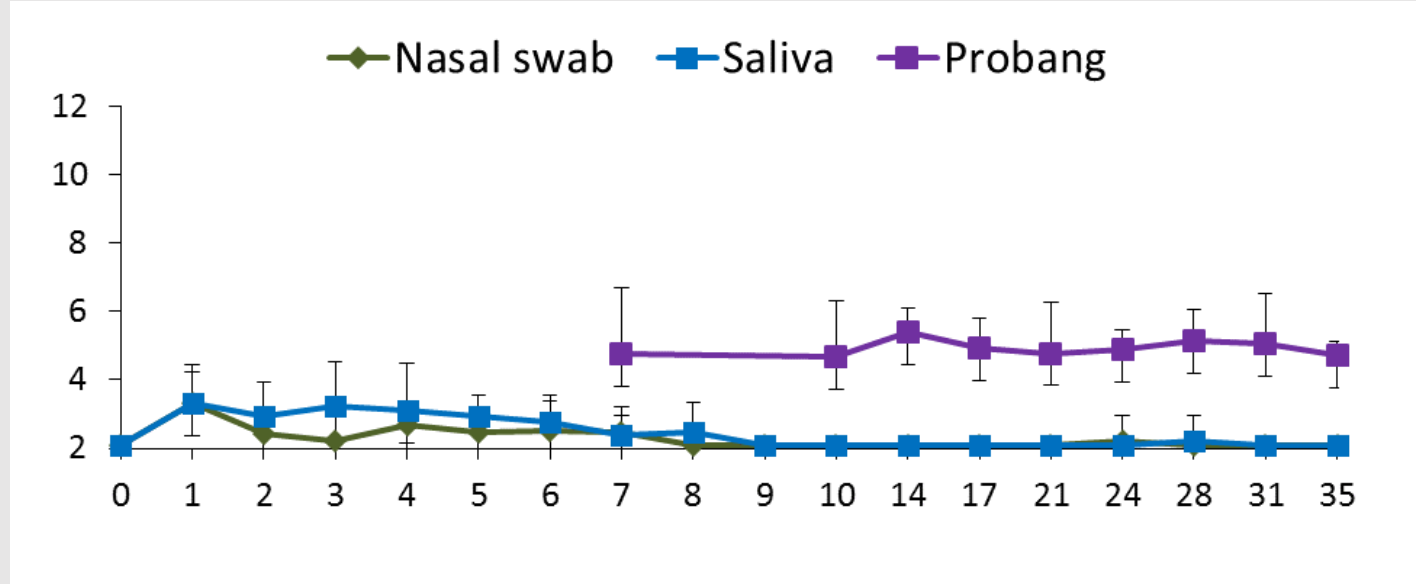
(Received September 9, 1967)

Cattle examination. The data reported are limited to cattle observed for 4 weeks following inoculation. The term virus "carrier" is applied to cattle in which virus was detectable to the end of that period. For a period of 14 DPI, the cattle were examined daily for clinical signs of FMD. Most OP fluid specimens were collected and tested (16) at weekly intervals. However, from each of groups III and IV, 8 cattle were tested only at 28 DPI.

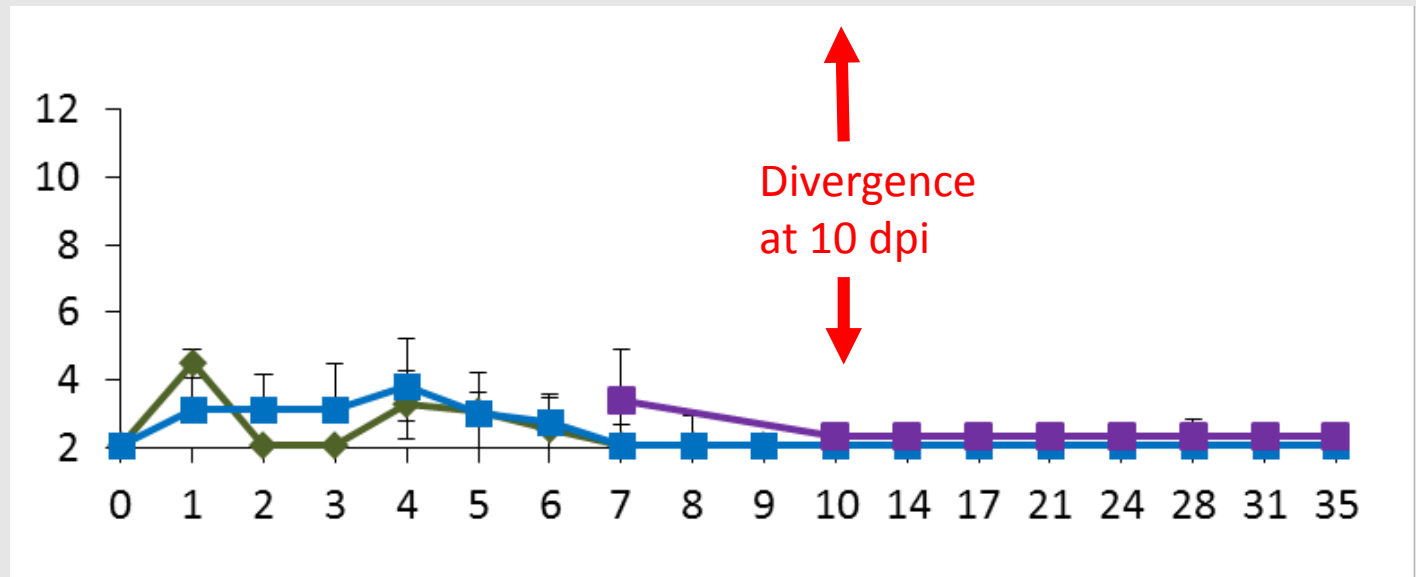
FMDV persistence definition; from laboratory data

Vaccinated cattle

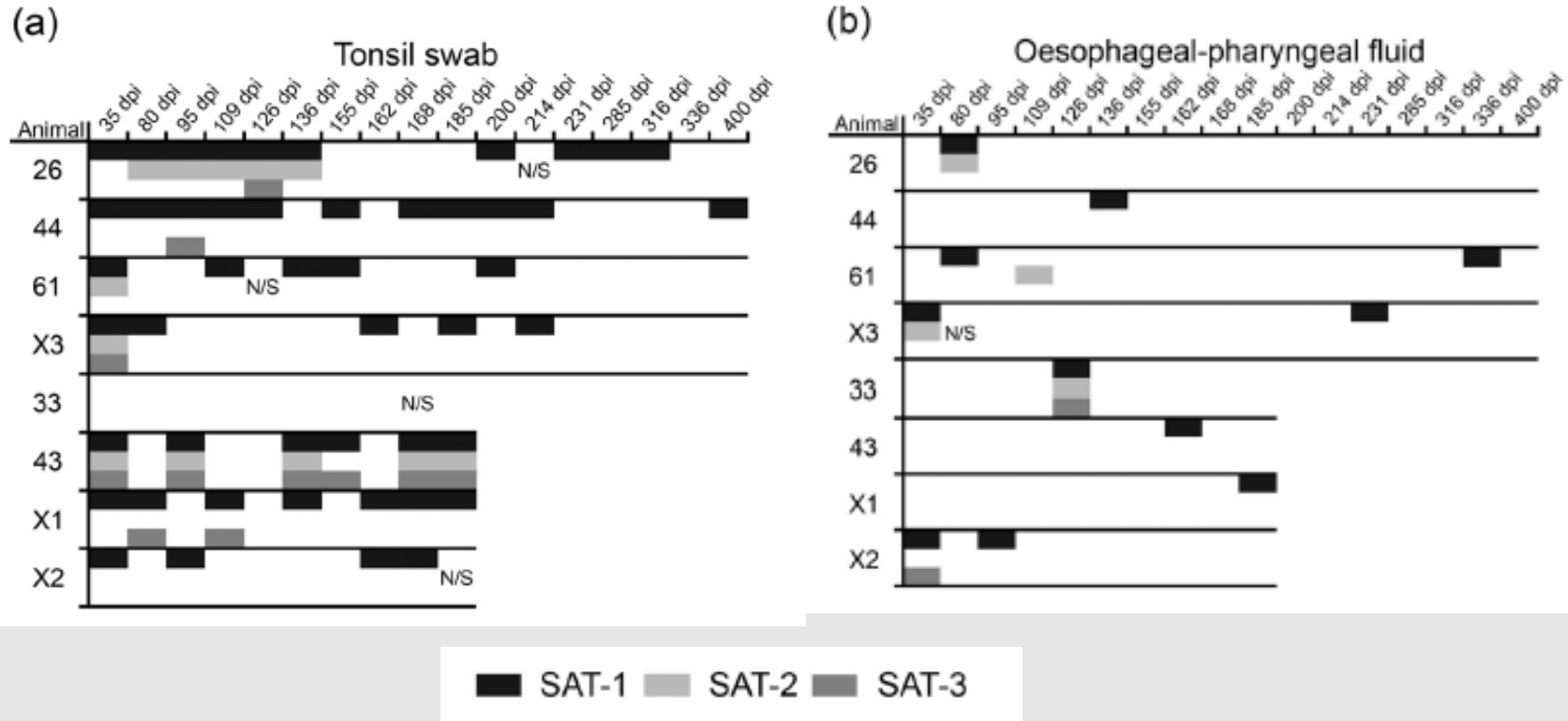
Carriers



Non-Carriers



FMDV shedding in tonsil swabs and probangs



Maree et al '16

Experimental studies

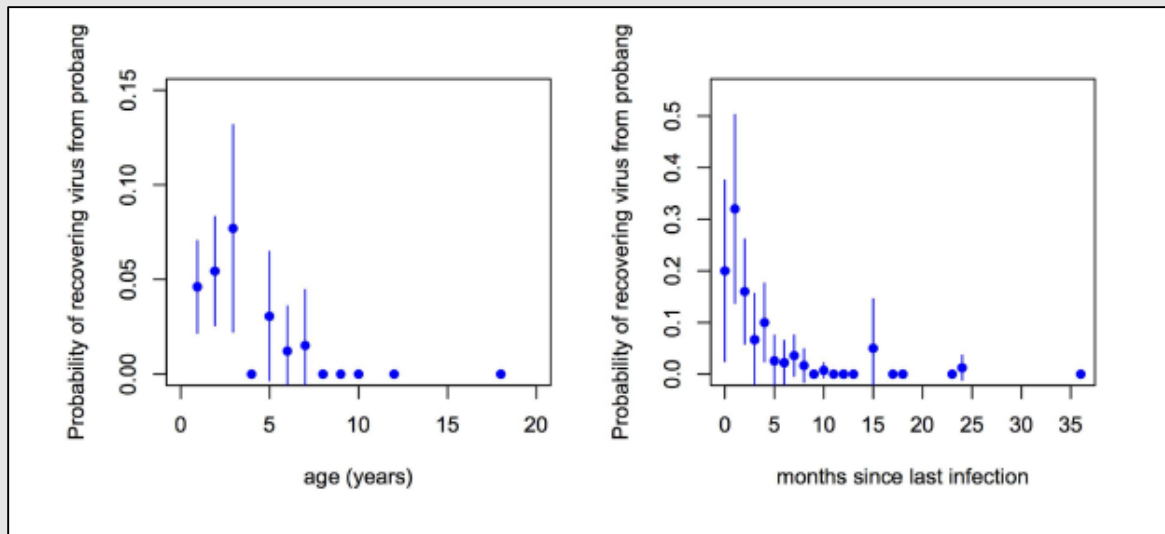
Limitations

- Experimental capacity (practical burden)
 - Restrictions in animal numbers and duration of investigations
 - Logistics and Animal Welfare concerns
- Questionable applicability to field situation
 - Virus used
 - Environmental conditions
 - Findings limited to experimental virus strain/host species
 - Feed
 - husbandry

FMDV persistence definition; from modeling field data

Predictive statistical models for identifying persistently infected cattle:

$$\text{Prob}(\text{probang}+'ve) = \text{invlogit}(-2.91 - 0.32 \times \text{monlast} + 0.22 \times \text{age} - 0.14 \times \text{age}^2 + 2.07 \times \text{VNTany} + 0.04 \times \text{age} \times \text{monlast})$$



Field studies

Advantages

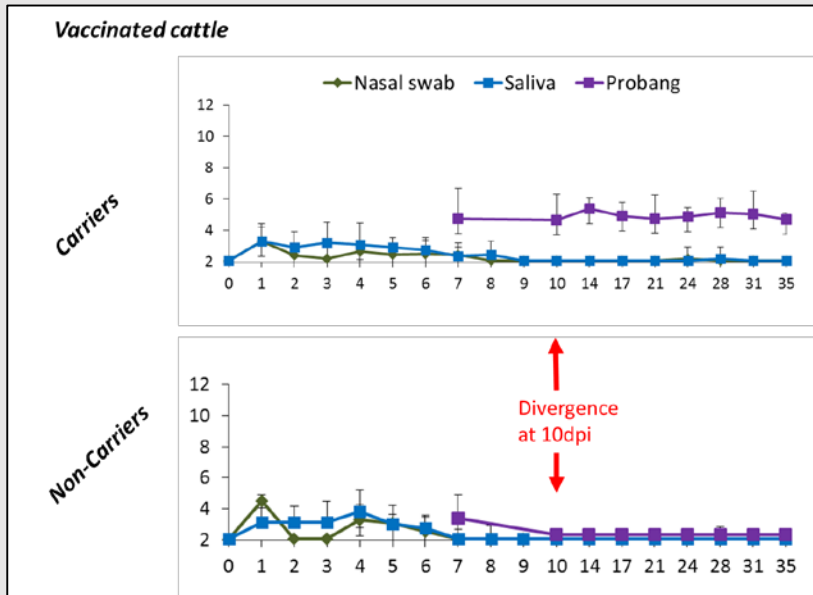
- Scale, scope, and duration unlimited
- Undeniable link to reality
- Enables association of specific findings and known events
 - Risk analysis based on actual
- Spatiotemporal relationships of viruses and outbreaks
 - Retrospective estimation of transmission pathways
- Multi-factorial analysis → FMD Epidemiology & Ecology

Limitations

- Unknown (uncontrolled) background context
- Limited information available
- Outcome and conclusions specific to geographic region investigated
- Intrinsic technical challenges(cold chain, transport)
- Limited capability (no daily sampling, operator variability)
- Breadth of relevance difficult to ascertain

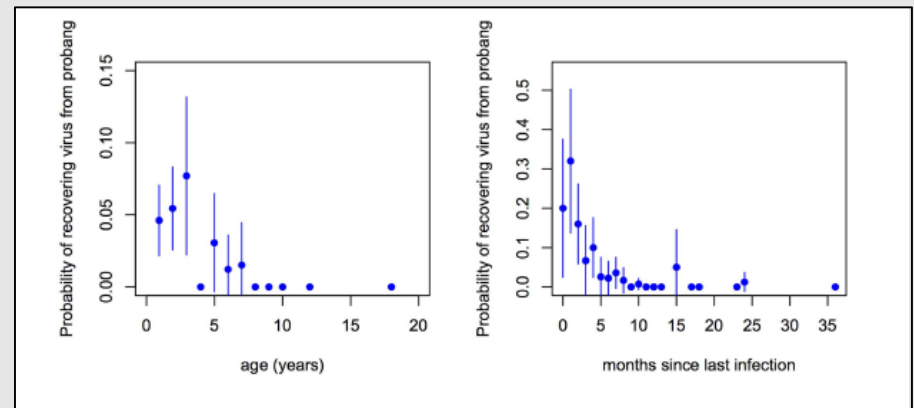
2 new criteria to characterize the carrier state

Unified



Stenfeldt et al 2016

$$\text{Prob}(\text{probang}+'ve) = \text{invlogit}(-2.91 - 0.32 \times \text{monlast} + 0.22 \times \text{age} - 0.14 \times \text{age}^2 + 2.07 \times \text{VNTany} + 0.04 \times \text{age} \times \text{monlast})$$

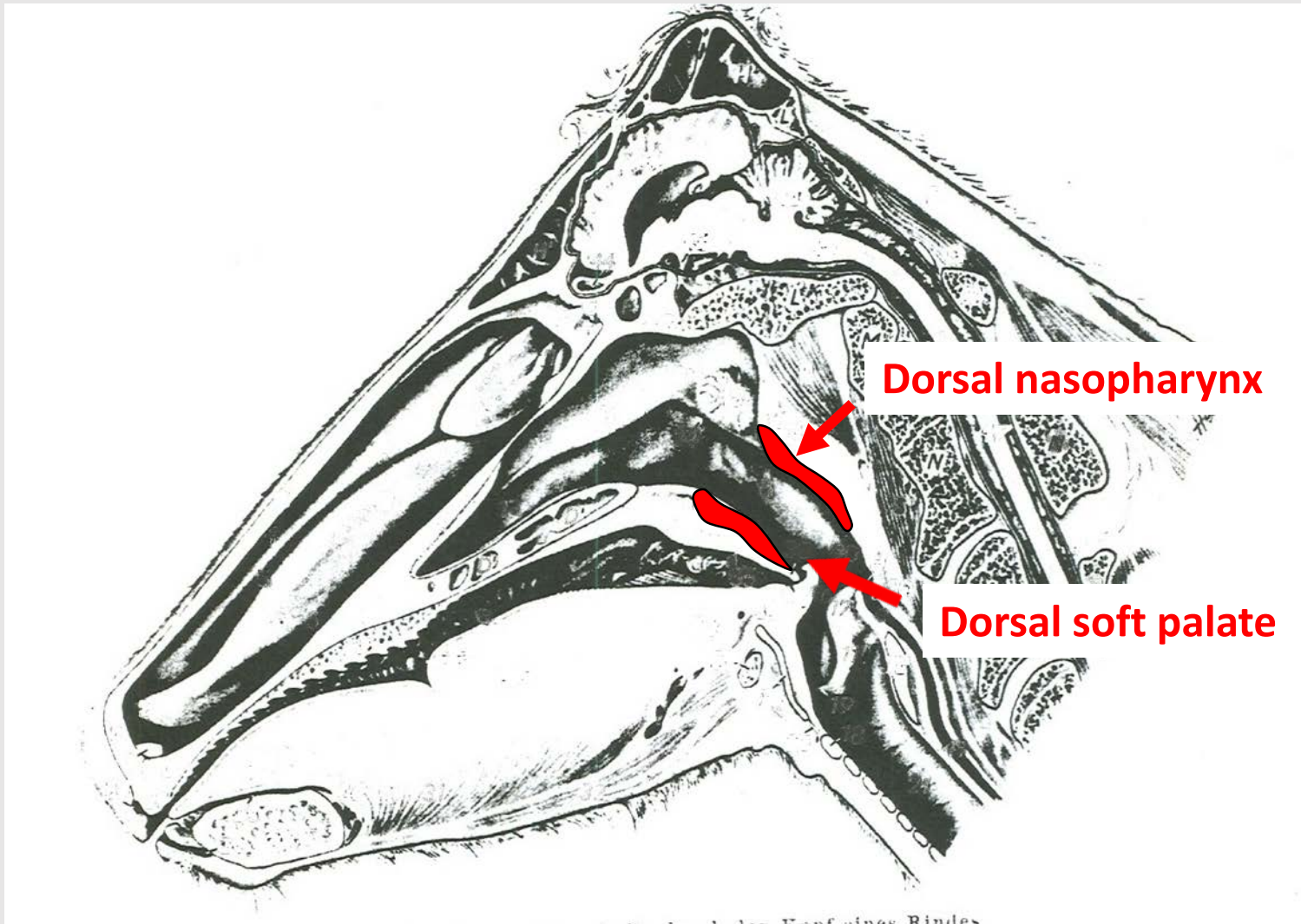


Bronsvort et al

Comprehensive

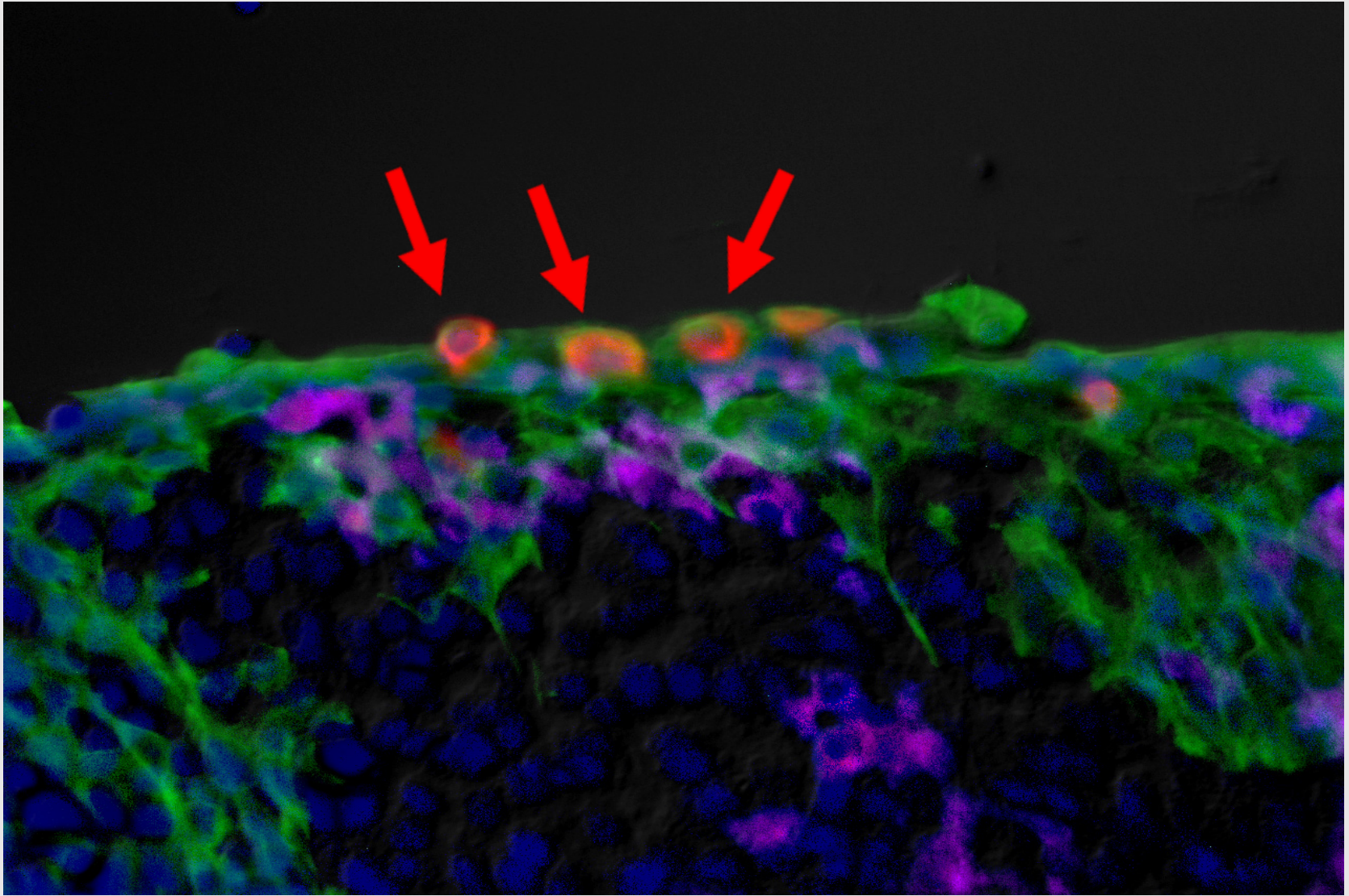
Sites of infection in carriers

Anatomic localization of persistent FMDV: Nasopharynx



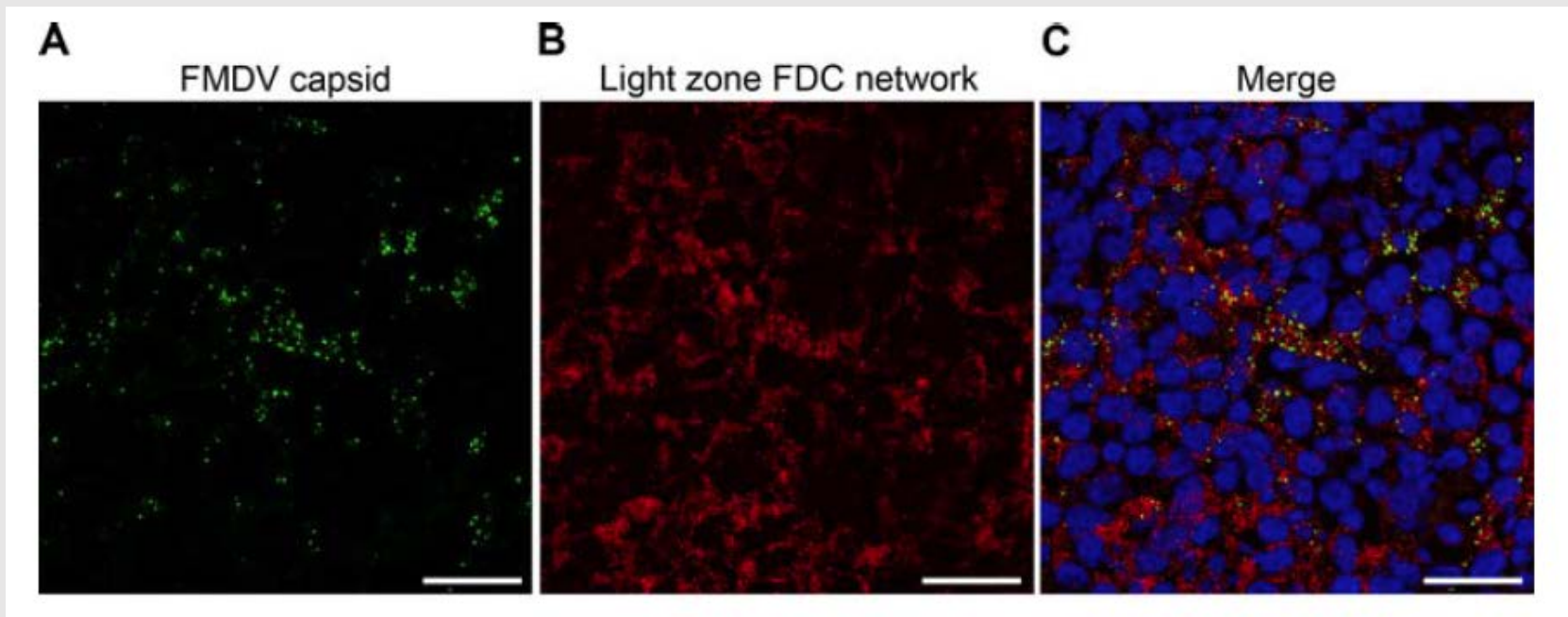
Not the oropharynx.

Anatomic localization of persistent FMDV in the bovine nasopharyngeal mucosa



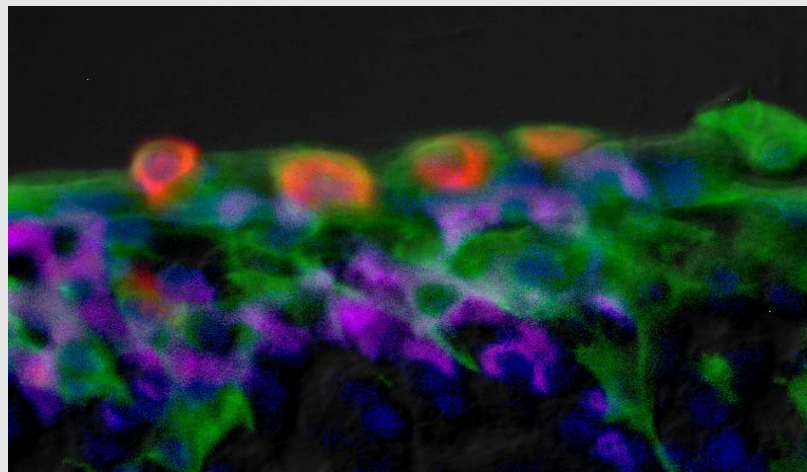
Cytokeratin, FMDV VP1, FMDV 3D, MHC II

Anatomic localization of persistent FMDV



Submandibular LN, 38 dpc

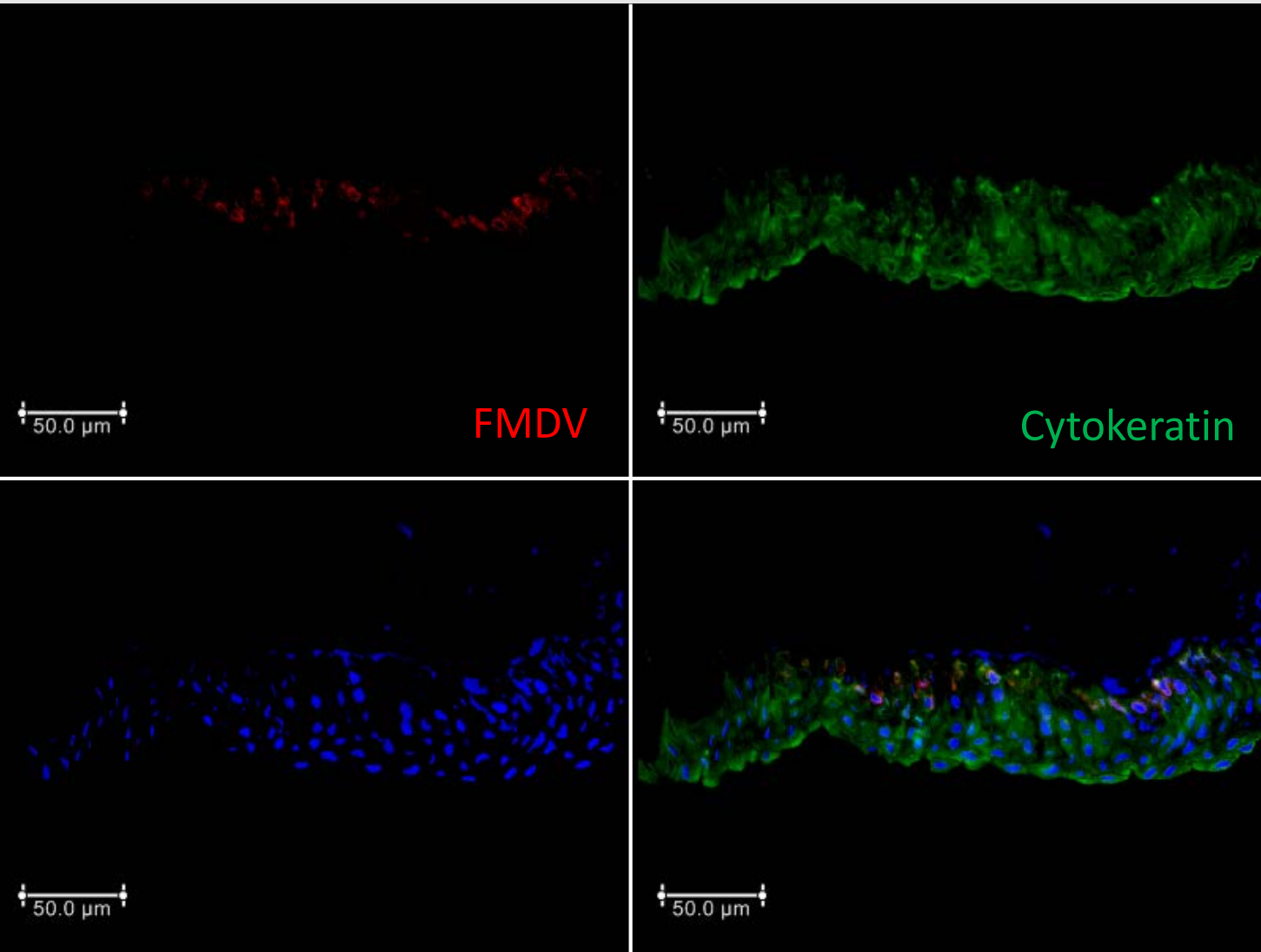
Not mutually exclusive findings.



Juleff et al., Plos ONE (2008)

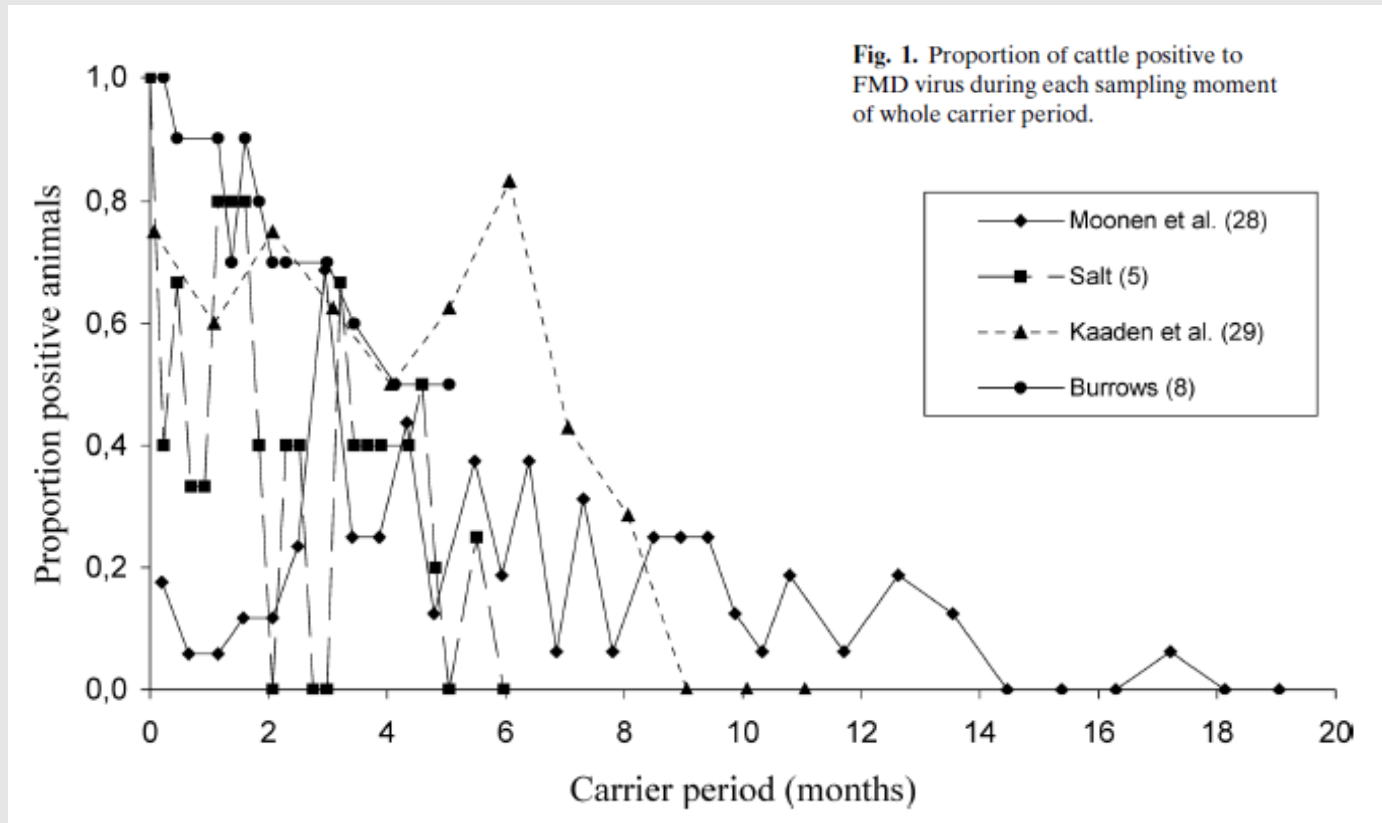
Anatomic confirmation from the field;

Nasopharynx of Asian buffalo



Duration of carrier state

Duration of FMDV persistence (extinction)

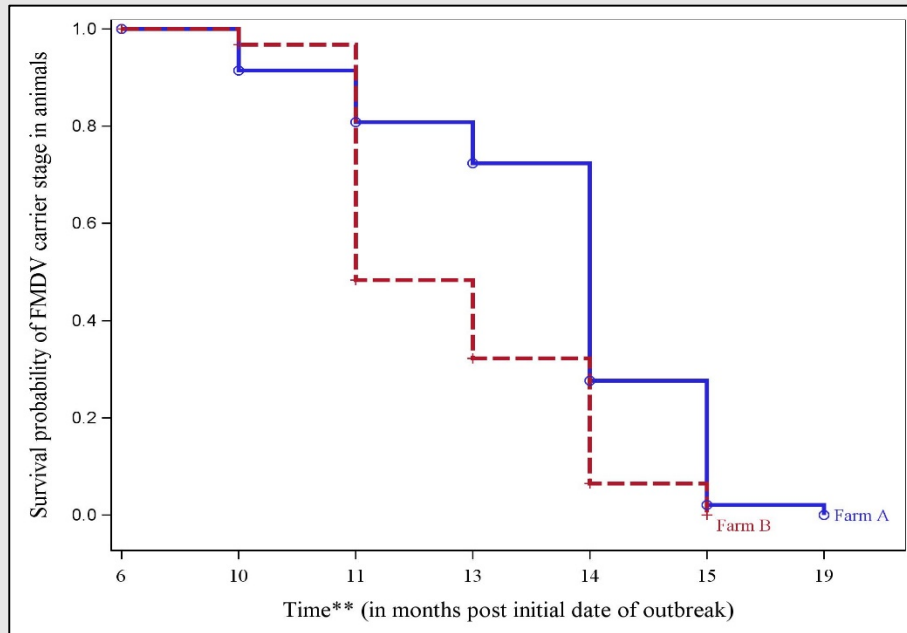


Meta-analysis of published experimental studies

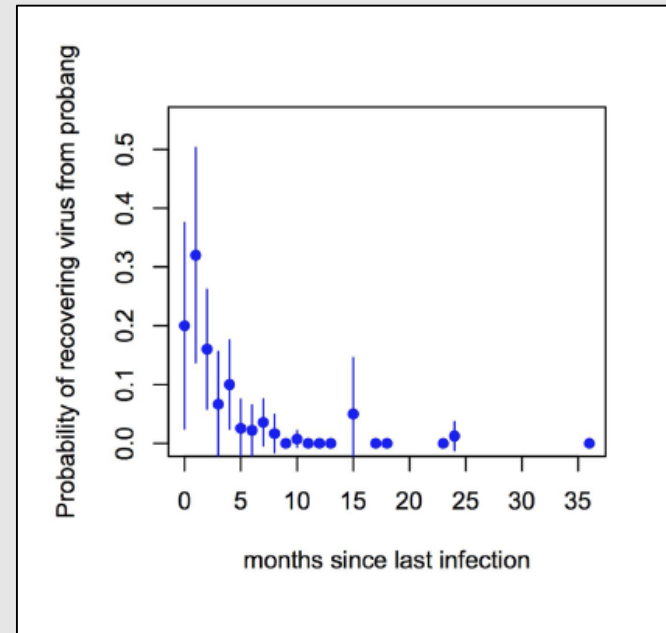
Decrease in proportion positive animals: 0.11 per carrier month

Duration of FMDV persistence: field confirmation

India (field-based)



Cameroon (field-based)



Hayer et al., Duration of FMDV carrier state based on detection of FMDV RNA in probang samples from two farms in Uttarakhand, India

Bronswort et al., Unlikely to detect probang-positive animals (by VI) >12 months post-outbreak

Viral genomic changes during the carrier state

Experimental studies

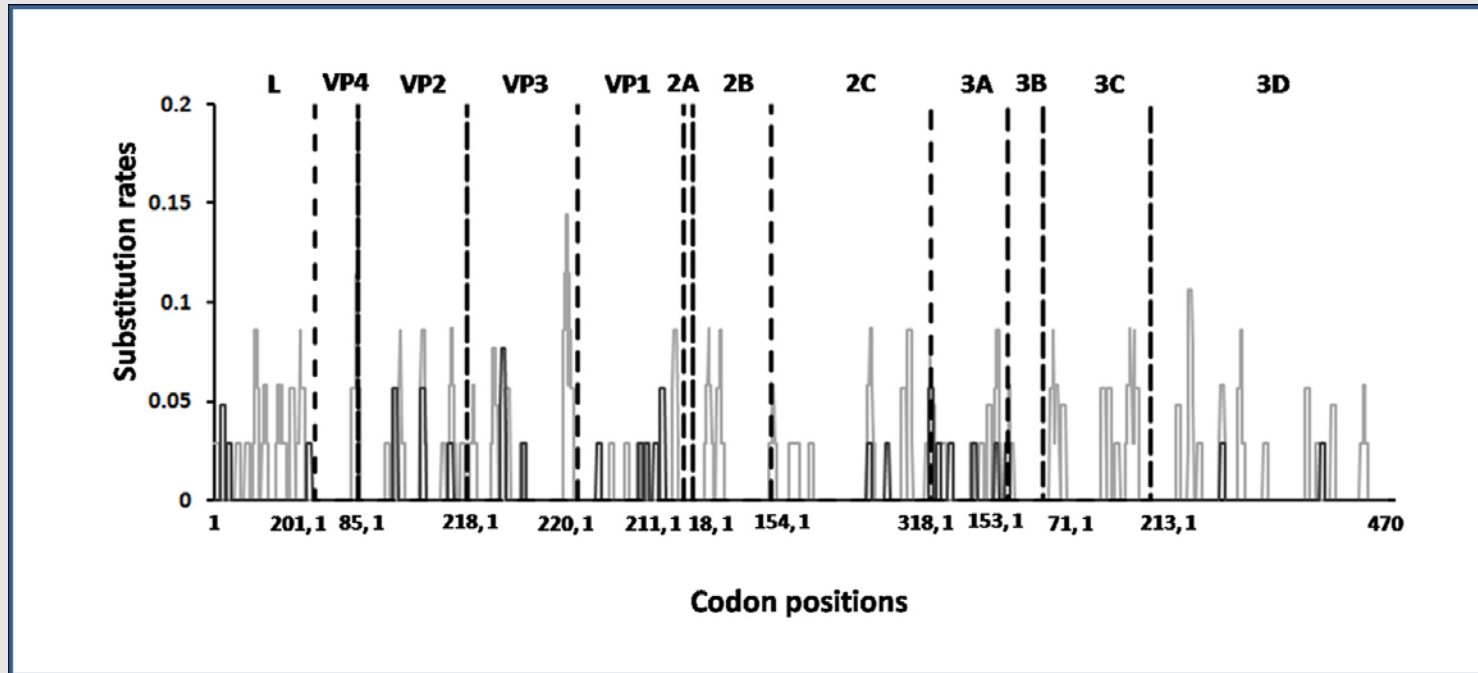
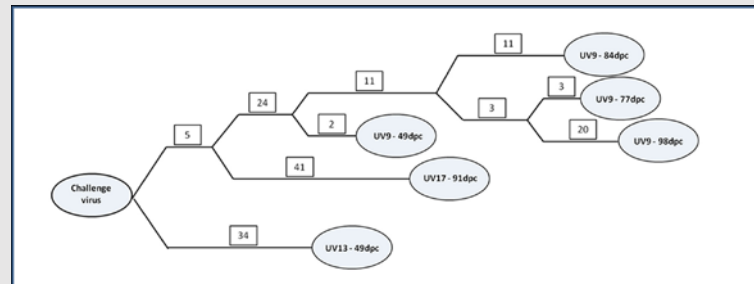
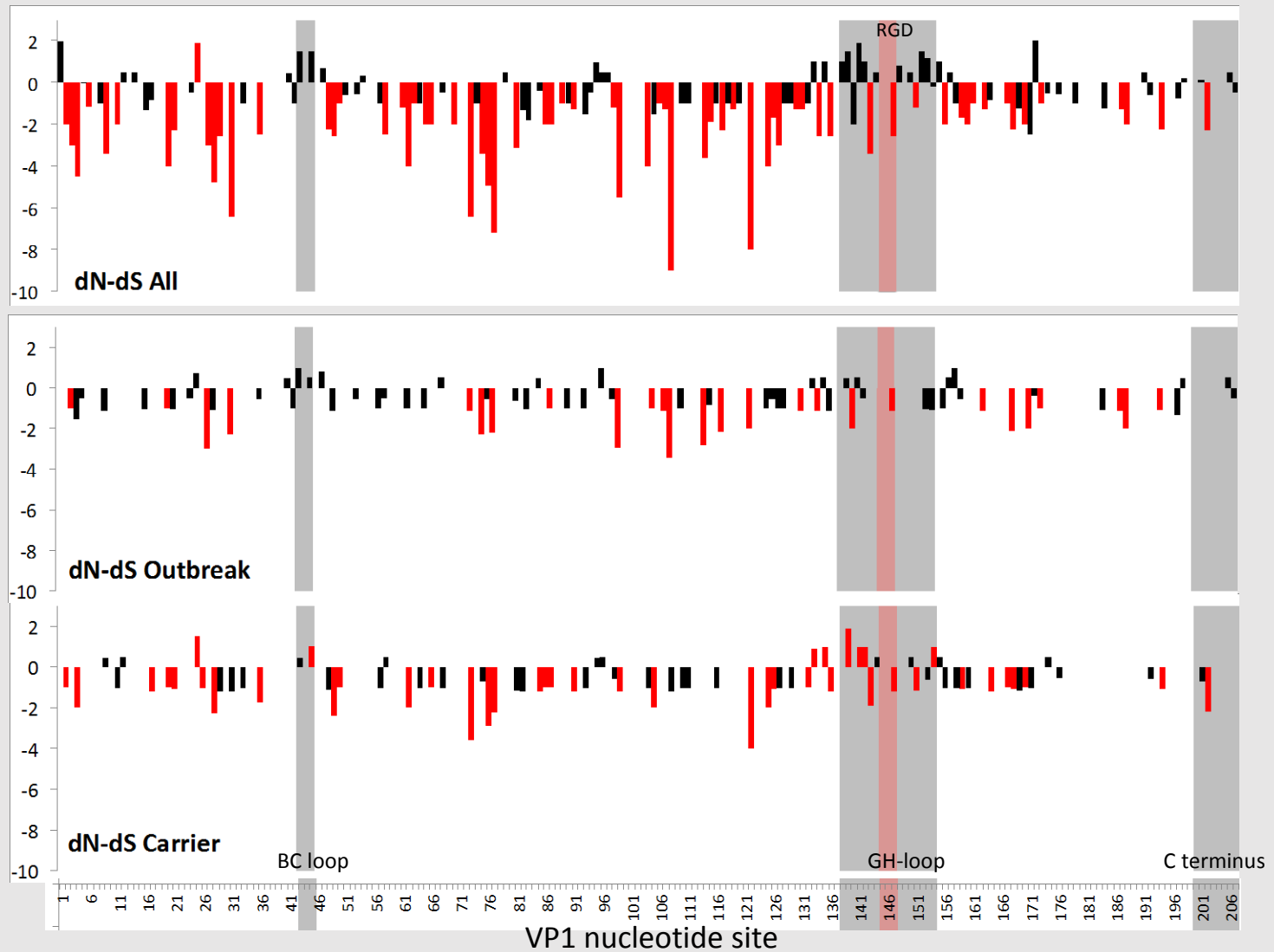


Fig 2. Synonymous (grey) and non-synonymous (black) changes in the ORF of the genomes of the FMD viruses obtained from carrier animals.

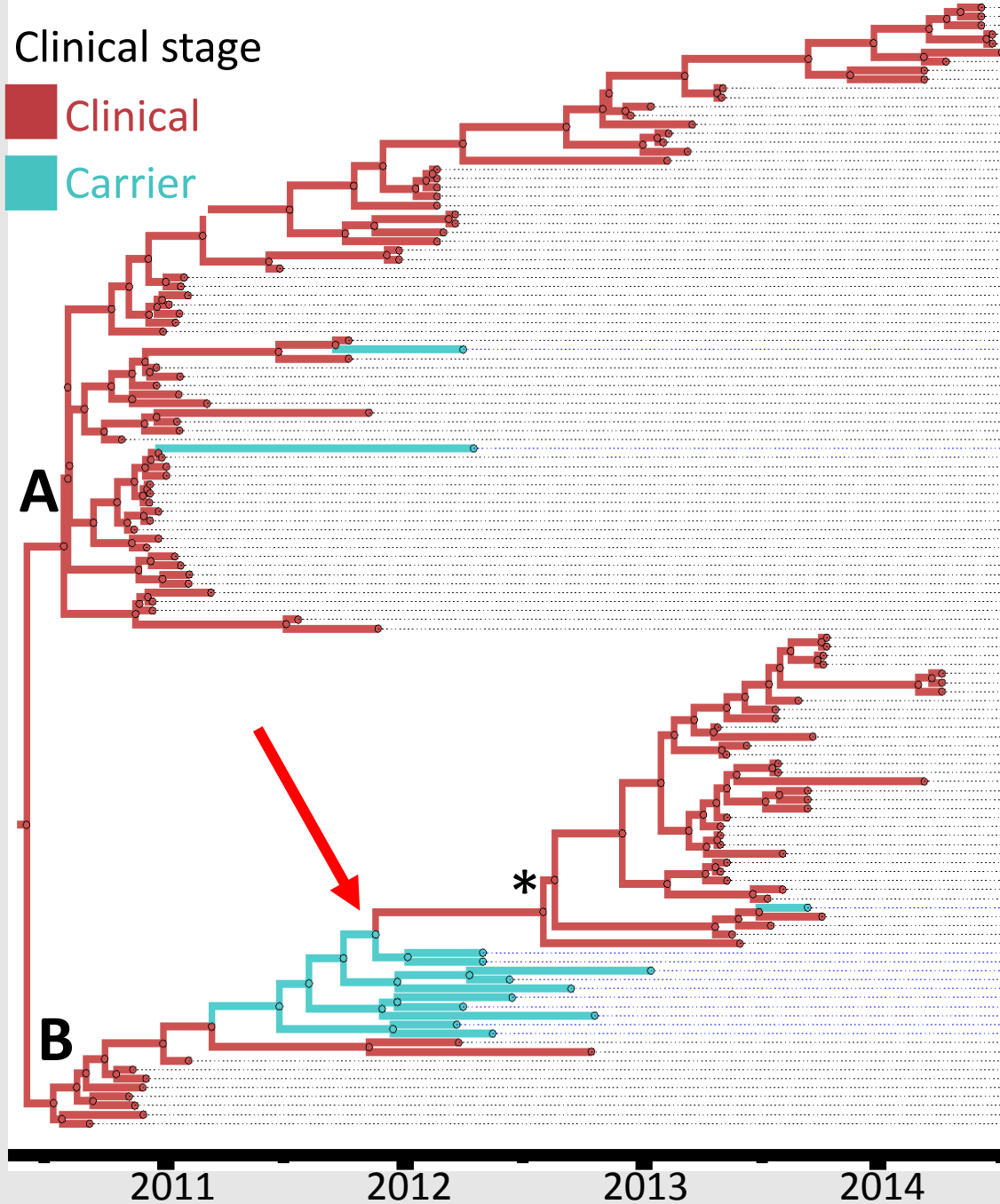


Genomic variation; carriers from the field Vietnam



Clinical stage

- Clinical
- Carrier



- ### Location_area
- CenHigh
 - China
 - Kazakhstan
 - Mekong_River_Delta
 - NCCoast
 - Northeast
 - Northwest
 - Red_River_Delta
 - South_Central_Coast
 - Southeast

- ### Host species
- Buffalo
 - Cattle
 - Pig

- ### Clinical stage
- Outbreak
 - Carrier

2010/03/2011 Buffalo LangSon 2011/01/01
2010/25/2010 Cattle Hoabinh 2011/01/14
2010/27/2010 Cattle Hoabinh 2011/02/27
2010/28/2011 Buffalo Ha Giang 2011/02/17
2010/03/2011 Buffalo LangSon 2011/03/01
2010/10/2011 Cattle SonLa 2011/04/04
2010/10/2011 Cattle SonLa 2011/01/08
Kurchnum/08/2011 UK Kurchnum 2011/08/15
2013/10/30
2013/08/13
2013/05/17
2013/10/09
2013/08/17
2013/01/01
2013/05/31
2013/10/01
2013/05/21
3/08/24
3/01
8/06
3/3
3/5
3/7
2011/11/2012 Pig BaMie 2011/11/05
2011/12/2011 Cattle Ha Giang 2011/03/01
2011/12/2011 Cattle Quang Tri 2011/10/01
2011/11/2014 Cattle Quang Tri 2013/10/01
2011/15/2013 Cattle NinhThuan 2013/05/31
2011/11/2013 Cattle QuangNam 2013/05/21

- Location
- Host species
- Clinical stage

Transmission during the carrier state

Transmission from FMDV carriers

Table II. List of Experiments Used for the Estimation of Transmission Rate Parameter β

Ref. No. ^a	Number of			N ^b	Contact Days	Species
	Cases	Carriers	Susceptibles			
(7)	0	6	4	10	35	Cattle to cattle
(30)	0	14	8	22	43	Cattle to cattle
(28)	0	17	1	18	549	Cattle to cattle
(31)	0	1	1	2	61	Cattle to cattle
(31)	0	2	2	4	274	Cattle to cattle
(31)	0	2	2	4	274	Cattle to cattle
(31)	0	2	1	3	274	Cattle to cattle
(11)	0	1	2	3	42	Cattle to cattle
(11)	0	1	2	3	42	Cattle to cattle
(11)	0	1	2	3	42	Cattle to cattle
(11)	0	1	2	3	28	Cattle to cattle
(11)	0	1	2	3	28	Cattle to cattle
(11)	0	1	2	3	28	Cattle to cattle
(31)	0	2	1	3	84	Sheep to cattle
(9)	0	5	6	11	548	Buffalo ^c to cattle
(32)	0	1	2	3	175	Buffalo to cattle
(32)	0	1	2	3	152	Buffalo to cattle
(33)	0	6	6	12	456	Buffalo to cattle
(34)	1	3	4	7	168	Buffalo to cattle
(35)	0	6	3	9	731	Buffalo to cattle
(36)	2 ^d	3	2	5	312	Buffalo to cattle
(37)	0	1	7	8	122	Buffalo to buffalo
(36)	1	3	1	4	198	Buffalo to buffalo
(38)	0	4	6	10	30	Cattle to pigs
(38)	0	4	6	10	30	Cattle to pigs
(38)	0	4	6	10	30	Cattle to pigs
(38)	0	4	6	10	30	Cattle to pigs
(38)	0	4	6	10	30	Cattle to pigs
(38)	0	4	6	10	30	Cattle to pigs
(39)	2	3	6	9	75	Cattle to pigs
(39)	0	2	6	8	86	Cattle to pigs
(39)	0	3	4	7	44	Cattle to pigs
(39)	0	1	4	5	34	Cattle to pigs
(39)	0	1	4	5	35	Cattle to pigs
(39)	0	2	4	6	29	Cattle to pigs
(31)	0	2	4	6	91	Cattle to pigs

^aNumber of reference (in list). In one paper, more than one experiment or experimental unit could be included.

^bN = Total number of animals (carriers + susceptible).

^cAfrican buffalo (*Syncerus caffer*).

^dIt was assumed that only one susceptible became infected due to contact with the carrier, and the second was infected by the first contact-infected animal.

Tenzin et al., 2008: Meta-analysis of FMDV transmission from persistently infected cattle and buffalo
0.0148 infections per carrier per month

Transmission from FMDV carriers

RESEARCH ARTICLE


Virus Excretion from Foot-And-Mouth Disease Virus Carrier Cattle and Their Potential Role in Causing New Outbreaks

Aravindh Babu R. Parthiban, Mana Mahapatra, Simon Gubbins, Satya Parida*

The Pirbright Institute, Ash Road, Pirbright, Surrey GU24 0NF, United Kingdom

* satya.parida@pirbright.ac.uk

No transmission from persistently infected cattle to (4) contact exposed cattle

 OPEN ACCESS

Citation: Parthiban ABR, Mahapatra M, Gubbins S, Parida S (2015) Virus Excretion from Foot-And-Mouth Disease Virus Carrier Cattle and Their Potential Role in Causing New Outbreaks. PLoS ONE 10(6): e0128815. doi:10.1371/journal.pone.0128815

Editor: Jagadeesh Bayry, Institut National de la Santé et de la Recherche Médicale (INSERM), FRANCE

Received: March 31, 2015

Accepted: April 30, 2015

Published: June 25, 2015

Copyright: © 2015 Parthiban et al. This is an open access article distributed under the terms of the

ences in viral shedding between carrier and non-carrier animals, quantify the transmission rate of FMDV infection from carriers to susceptible animals and identify potential viral determinants of viral persistence. We collected nasal and saliva samples from 32 vaccinated and 7 unvaccinated FMDV carrier cattle and 48 vaccinated and 13 unvaccinated non-carrier cattle (total n=100) during the acute phase of infection (up to 28 days post-challenge) and then from limited number of animals up to a maximum 168 days post-challenge. We demonstrate that unvaccinated cattle excrete significantly higher levels of virus for longer periods compared with vaccinated cattle and this is independent of whether or not they subsequently become carriers. By introducing naïve cattle in to the FMDV carrier population we show the risk of new outbreaks is clearly very low in controlled conditions, although there could still be a potential threat of these carrier animals causing new outbreaks in the field situation. Finally, we compared the complete genome sequences of viruses from carrier cattle with the challenge virus and found no evidence for viral determinants of the carrier state.

breaks is
outbreak
the differ-

Transmission from FMDV carriers



AMERICAN
SOCIETY FOR
MICROBIOLOGY

Journal of
Virology



Differential Persistence of Foot-and-Mouth Disease Virus in African Buffalo Is Related to Virus Virulence

Francois Maree,^{a,b} Lin-Mari de Klerk-Lorist,^c Simon Gubbins,^d Fuquan Zhang,^d Julian Seago,^d Eva Pérez-Martín,^d Liz Reid,^d Katherine Scott,^a Louis van Schalkwyk,^c Roy Bengis,^c Bryan Charleston,^d Nicholas Juleff^d

Transboundary Animal Disease Programme, ARC-Onderstepoort Veterinary Institute, Onderstepoort, South Africa^a; South Africa Department of Microbiology and Plant Pathology, University of Pretoria, Pretoria, South Africa^b; Office of the State Veterinarian, Department of Agriculture, Forestry and Fisheries, Skukuza, South Africa^c; The Pirbright Institute, Woking, Surrey, United Kingdom^d

NO transmission from persistently infected African buffalo to contact exposed cattle

Transmission from FMDV carriers?



Field trial: Vietnam

- **NO transmission from persistently infected cattle to contact exposed cattle**

Transboundary and Emerging Diseases

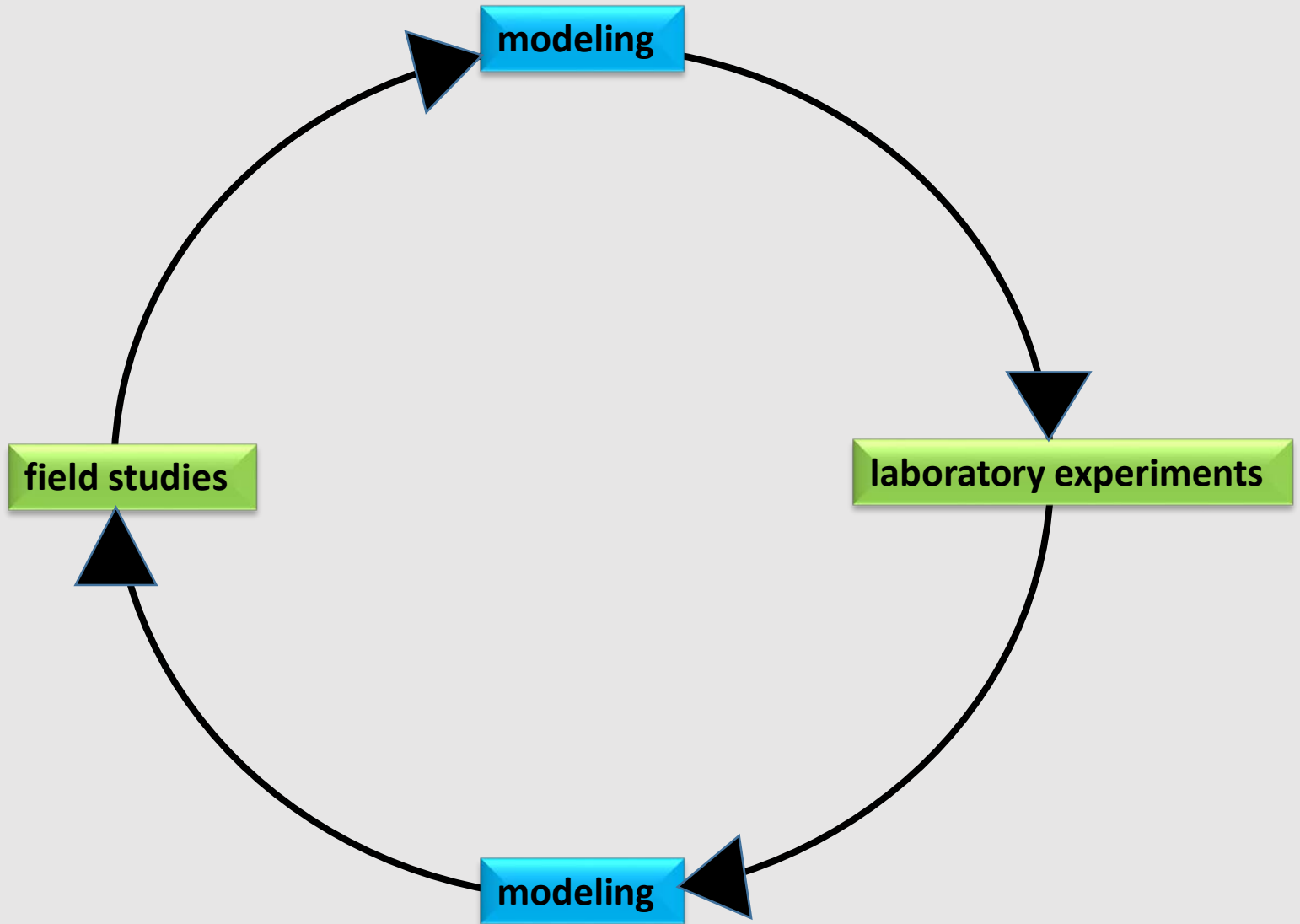
Transboundary and Emerging Diseases

ORIGINAL ARTICLE

An Integrative Analysis of Foot-and-Mouth Disease Virus Carriers in Vietnam Achieved Through Targeted Surveillance and Molecular Epidemiology

H. C. de Carvalho Ferreira^{1,2}, S. J. Pauszek¹, A. Ludi^{1,2,*}, C. L. Huston³, J. M. Pacheco¹, V. T. Le⁴, P. T. Nguyen⁴, H. H. Bui⁴, T. D. Nguyen⁵, T. Nguyen⁵, T. T. Nguyen⁶, L. T. Ngo⁴, D. H. Do⁶, L. Rodriguez¹ and J. Arzt¹

Keeping it real



Acknowledgements

Co-authors:

- Luis Rodriguez
- Carolina Stenfeldt
- Barbara Brito

Analytics:

- Shivdeep Hayer
- Kimberly VanderWaal
- Amy Delgado

International Collaborators:

- Do Huu Dung
- Ngo Thanh Long
- Rajeev Ranjan
- Jitendra K. Biswal
- Bramhadev Pattnaik

Funding:

- FADRU-ARS-USDA (base support)
- DTRA-DOD
- CEAH-APHIS-USDA (IAA)
- Science and Technology Directorate, DHS (IAA; primary animal work)



Keeping it real

