

# FAO World Reference Laboratory for Foot-and-Mouth Disease (WRLFMD)

## Genotyping Report

Report Date for this Batch: 25 April 2013

FMDV type SAT 2

Country: Cameroon

Period: 2012

No. of samples: 9

BATCH: WRLFMD/2012/00038



The contents of this report are copyright and should not be reproduced without permission

© The Pirbright Institute

















# FAO World Reference Laboratory for FMD Genotyping Report

IAH-P-EP-MEG-FOR-005-3

Page 1 of 1

Serotype: SAT2 WRLFMD Ref No: CAR/24/2012 Batch No: WRLFMD/2012/00038 Sender Ref: DNL20603 Location: Adamawa, Cameroon Date collected: 23/06/2012 Date received by WRLFMD: 17/08/2012 Date received for sequencing: 16/04/2013 Species: Bovine Material used: BTy1 16/01/2013; BTy2 04/02/2013 Region sequenced: VP1 RT-PCR primers: SAT2-1C445F/SAT-2B208R SAT2-P1-1223F/SAT-2B208R	Report date: 24/04/2013 Reported by: N.J. Knowles Checked by: D.P. King  Topotype: VII Genotype/strain: Lib-12 Sequence filename: CAR12-24.SEQ Date sequence last updated: 24/04/2013 No. of Nt determined: 648 No. of ambiguities: 0 Gene length: 648 Total no. of comparisons: 632 Min. no. of nt for comparison: 600 Total turn-around time: 250 days Sequencing time: 8 days
Comments:	

<b>Most Closely Related Viruses</b>									
Pos.	Virus name	Filename	No. nt comp.	No. nt match.	No. of ambig.	% Id.	% Diff.	Topotype	Strain
1	SAT2/CAR/10/2012	CAR12-10	648	647	0	99.85	0.15	VII	Lib-12
2	SAT2/CAR/2/2012	CAR12-02	648	647	0	99.85	0.15	VII	Lib-12
3	SAT2/CAR/3/2012	CAR12-03	648	647	0	99.85	0.15	VII	Lib-12
4	SAT2/CAR/4/2012	CAR12-04	648	647	0	99.85	0.15	VII	Lib-12
5	SAT2/CAR/9/2012	CAR12-09	648	647	0	99.85	0.15	VII	Lib-12
6	SAT2/CAR/1/2012	CAR12-01	648	646	0	99.69	0.31	VII	Lib-12
7	SAT2/CAR/8/2012	CAR12-08	648	643	0	99.23	0.77	VII	Lib-12
8	SAT2/LIB/39/2012 (JX570633)	LIB12-39	648	642	0	99.07	0.93	VII	Lib-12
9	SAT2/LIB/40/2012 (JX570634)	LIB12-40	648	642	0	99.07	0.93	VII	Lib-12
10	SAT2/LIB/41/2012 (JX570635)	LIB12-41	648	642	0	99.07	0.93	VII	Lib-12
<b>Most Closely Related Reference Viruses</b>									
(see <a href="http://www.wrlfmd.org/fmd_genotyping/prototypes.htm">http://www.wrlfmd.org/fmd_genotyping/prototypes.htm</a> )									
Pos.	Virus name	Filename	No. nt comp.	No. nt match.	No. of ambig.	% Id.	% Diff.	Topotype	Strain
1	SAT2/CAR/8/2005 (JX570616)	CAR05-08	648	585	0	90.28	9.72	VII	unnamed
2	SAT2/SAU/6/2000 (AF367135)	SAU00A06	647	580	1	89.64	10.36	VII	unnamed
3	SAT2/ZAI/1/74 (DQ009737)	ZAI74-AA	642	482	0	75.08	24.92	VIII	unnamed
4	SAT2/KEN/3/57 (AJ251473)	KEN57G03	648	481	0	74.23	25.77	IX	unnamed
5	SAT2/UGA/51/75 (AY343963)	UGA75-AA	648	481	0	74.23	25.77	XII	unnamed
6	SAT2/ETH/2/91 (AY343938)	ETH91-AB	648	479	0	73.92	26.08	XIV	unnamed
7	SAT2/RWA/1/2000* (AF367134)	RWA00-01	648	479	0	73.92	26.08	VIII	unnamed
8	SAT2/UGA/19/98 (AY343969)	UGA98-AA	648	478	0	73.77	26.23	X	unnamed
9	SAT2/KEN/2/84 (AY343941)	KEN84-AC	648	475	0	73.30	26.70	IX	unnamed
10	SAT2/ZAI/1/82 (AF367100)	ZAI82-01	647	474	1	73.26	26.74	X	unnamed

nt, nucleotides

\*, not a WRLFMD reference number

Assembled with Report Generator v4.1

© The Pirbright Institute (not to be reproduced without permission)



# Report on FMDV SAT 2 in Cameroon in 2012

(cattle probang samples)

Batch: WRLFMD/2012/00038

◆ indicates viruses in this batch

Software: MEGA 5.1

Analysis

Analysis ----- Phylogeny Reconstruction

Scope ----- All Selected Taxa

Statistical Method ----- Neighbor-joining

Phylogeny Test

Test of Phylogeny ----- Bootstrap method

No. of Bootstrap Replications ----- 1000

Substitution Model

Substitutions Type ----- Nucleotide

Model/Method ----- Kimura 2-parameter model

Substitutions to Include ----- d: Transitions + Transversions

Rates and Patterns

Rates among Sites ----- Uniform rates

Pattern among Lineages ----- Same (Homogeneous)

Data Subset to Use

Gaps/Missing Data Treatment ----- Pairwise deletion

Codons Included ----- 1st+2nd+3rd+Non-Coding

No. of Sites : 651

No Of Bootstrap Reps = 1000

Only bootstrap values of 70% and above are shown

\*, not a WRLFMD Ref. No.

N.J. Knowles, K. Bachanek-Bankowska & J. Wadsworth,  
25 April 2013

© The Pirbright Institute

(not to be reproduced without permission)

