



# Alternatives for in vivo Foot-and-mouth disease vaccines potency tests: Serotype dependency of the in vitro vaccine potency models



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## Introduction and objectives

Foot-and-mouth disease (FMD) vaccine potency testing is performed by experimentally infecting vaccinated cattle. The *in vivo* 50% Protective Dose (PD<sub>50</sub>) test is the standard European procedure for the quality control of FMD vaccines. Due to ethical reasons, current research alliances focus on the replacement of the *in vivo* viral challenge by *in vitro* alternatives. Previously, Goris *et al.* (Vaccine, 2008) have validated an *in vitro* model for the FMDV reference strain O<sub>1</sub> Manisa. The present study aims to develop comparable models for serotype A and to test both models for serotype-dependency.

## First step: replicates of the in vivo PD<sub>50</sub> test

Five replicate *in vivo* PD<sub>50</sub> tests are performed with a FMDV A Iran 96 vaccine. Twenty-one days post vaccinal (21DPV) serum samples are collected to perform the serological analyses on which the alternative *in vitro* vaccine potency models will be based. Each PD<sub>50</sub> tests is performed with 17 naïve cattle that are randomly divided into 2 control animals and 3 groups of 5 animals, each group receiving a different vaccine dose. At 21 DPV all animals are challenged with homologous FMD A Iran 96 virus (A96). Eight days post challenge (8DPC) they are clinically inspected for generalisation of FMD and animals are classified as protected or unprotected against challenge. Then the Kärber (1931) formula is used to calculate the PD<sub>50</sub> for the vaccine batch.

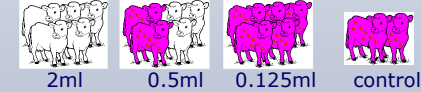
### Day 0: Vaccination

Vaccine dose groups



### 21 DPV: Virus challenge (10<sup>4</sup> BID<sub>50</sub>)

Vaccine dose groups



### 8 DPC: Clinical inspection of cattle

PD<sub>50</sub> result

## Second step : in vitro vaccine potency model

All 21 DPV sera are analysed (serum neutralisation (SNT A96) & liquid phase blocking ELISA (LPBE A96)) and with the resulting serum titres (t<sub>i</sub>) a logistic correlation model is built. This model correlates the serum titres with a probability of being protected against virus challenge. This results in a cut off titre (t<sub>co</sub>) which will be used to classify the animal sera. Then the Kärber formula is used to calculate the EPD<sub>50</sub> for the vaccine batch.

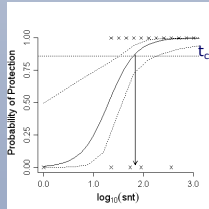
### Day 0: Vaccination

Vaccine dose groups



### 21 DPV:

Serum sampling



Serum titre Classification with Logistic regression model

t<sub>i</sub> ≥ t<sub>co</sub> p : protected  
t<sub>i</sub> < t<sub>co</sub> p : unprotected

Estimated EPD<sub>50</sub> result

## Third step : comparison PD<sub>50</sub> vs. EPD<sub>50</sub>

*In vitro* vaccine potency models are built for 2 different serological assays and 2 different laboratories. The EPD<sub>50</sub> results are estimated using the cut off from the A96 model. Alternatively the EPD<sub>50</sub> is also estimated against the cut off from an earlier developed O<sub>1</sub> Manisa model (Goris *et al.*, 2008). The table below shows the mean EPD<sub>50</sub> estimates for the 5 trials for each serological test (Test) and model cut off (Model). There is a difference between the mean *in vivo* PD<sub>50</sub> (20.5 with 95%CI between 14.9 and 27.4) and the estimated values, except for the LPBE A96 test from VAR which confidence intervals have a good overlap of the *in vivo* results.

Laboratory	Test	Model (t <sub>co</sub> )	Mean EPD <sub>50</sub>	95% CI
ARRIAH	SNT A96	SNT A96 (1.34)	11.1	9.2 - 19.1
VAR	SNT A96	SNT A96 (1.53)	10.2	8.4 - 19.1
ARRIAH	SNT A96	SNT O <sub>1</sub> (1.51)	5.8	5.0 - 14.0
VAR	SNT A96	SNT O <sub>1</sub> (1.68)	8.7	6.9 - 14.6
ARRIAH	LPBE A96	LPBE A96 (1.34)	6.0	5.0 - 11.2
VAR	LPBE A96	LPBE A96 (1.73)	14.7	12.5 - 21.1
ARRIAH	LPBE A96	LPBE O <sub>1</sub> (1.65)	2.1	1.7 - 3.1
VAR	LPBE A96	LPBE O <sub>1</sub> (1.95)	13.1	11.2 - 20.5

## Conclusion

For each serological assay the corresponding alternative *in vitro* vaccine potency model has to be validated for every individual laboratory and its serotype dependency has to be investigated.

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