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# MODELLING THE INTERACTIONS BETWEEN THE PORCINE RESPIRATORY AND REPRODUCTIVE SYNDROME VIRUS AND ITS TARGET CELLS: CONDITIONS FOR VIRUS CLEARANCE

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## INTRODUCTION

### Background

- PRRSV = a major concern for swine industry
- Lacking of efficient vaccines for infection control

### Immune response

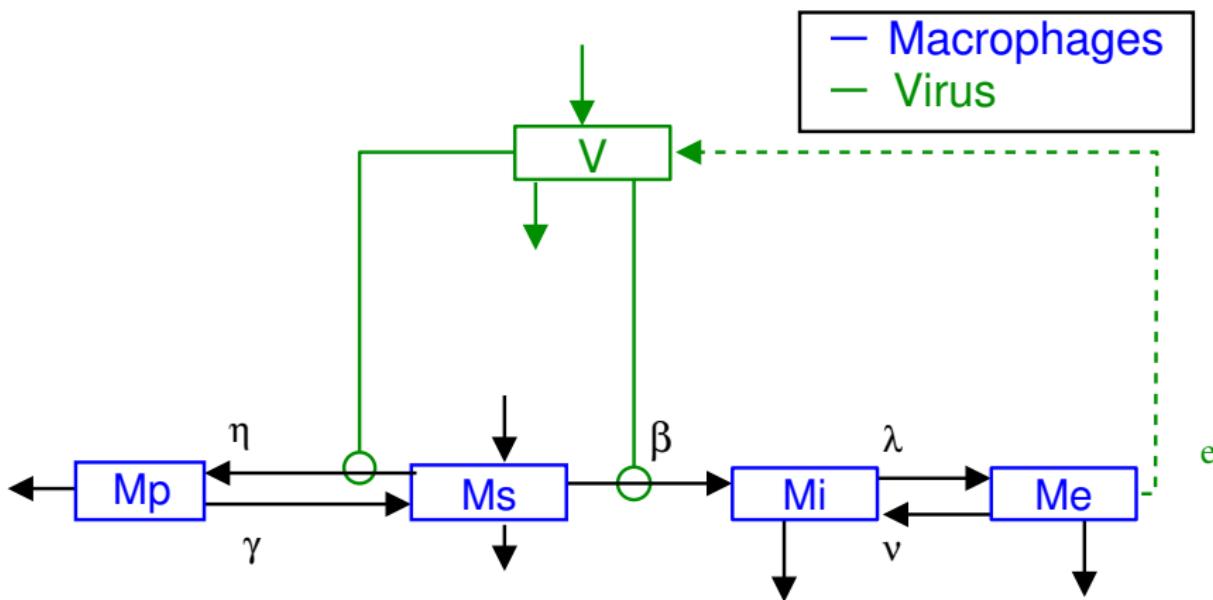
- High variability (host susceptibility & viral virulence)
- Mechanisms responsible for the resolution?  
⇒ MODELLING APPROACH

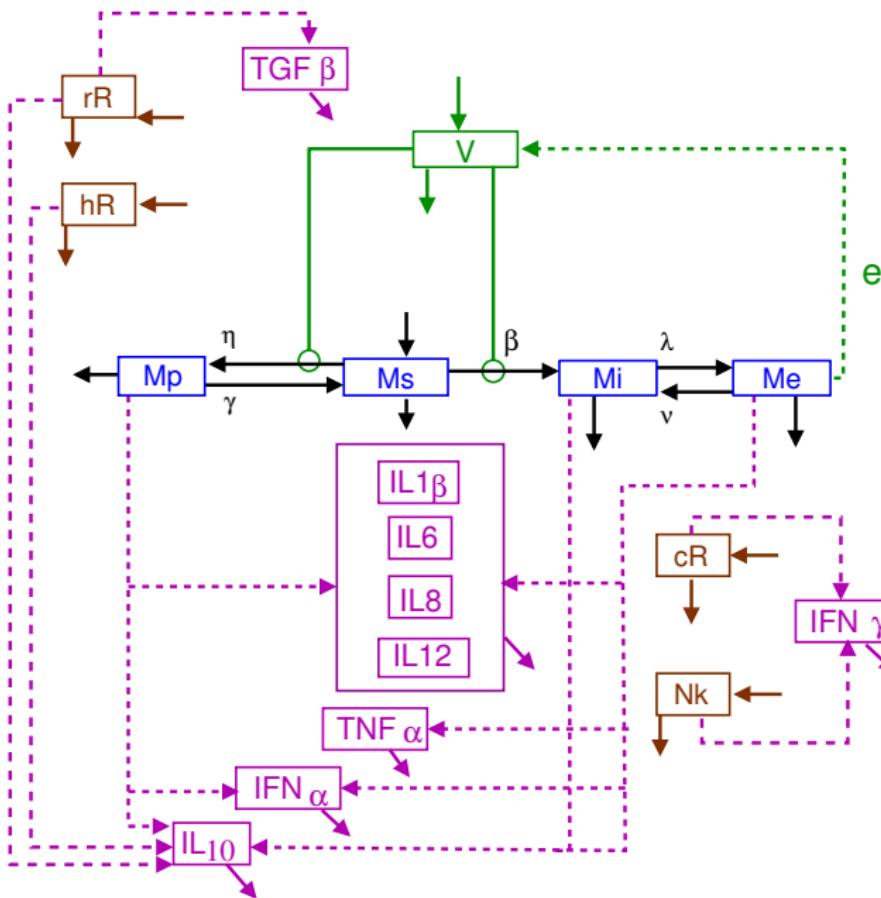
### Macrophage – virus interactions

- Target cells : pulmonary macrophages
- Macrophages = key immune cells (phagocytosis, inflammation, adaptive response activation)

MODEL OF THE IMMUNE RESPONSE centered on the macrophage-virus interactions in the lung ⇒ Exploration of control measures.

## CONCEPTUAL MODEL







## Model

- Formalised by ODE in a deterministic mathematical model

$$\dot{I}_\gamma = \rho_{I_\gamma} [ cR + N_k ] \frac{[1 + \kappa_1(L_{12}, T_\alpha)] [1 + \kappa_1(L_{12}, I_\alpha)] [1 + \kappa_1(I_\gamma)]}{[1 + \kappa_1(L_{10})] [1 + \kappa_1(T_\beta)]} - \mu_C I_\gamma$$

- 18 state variables & 30 parameters

## Calibration

- compiled data from litterature (experimental and modelling studies)
- lack of data to inform parameter values and high response variability

## Sensitivity analysis

- identify the parameter values resulting in realistic outputs
- identify the key parameters which have a strong influence on model outputs

## METHOD

### Factorial design

- Production of a fractional factorial design using the R library PLANOR  
(Monod *et al.*) ⇒ 2187 combinations

### Output selection

Viral titer, % infected macrophages, % phagocytizing macrophages,  
NK & cellular response cells

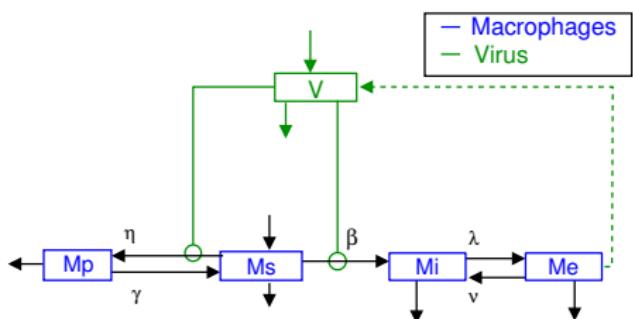
### Sensitivity analysis

Mutivariate method: R library MULTISENSI, Generalized Sensitivity  
Method (*Lamboni and Monod*)

## RESULTS : SUMMARY

Table of the parameter influences on model outputs: positive effects (++) , negative effects (--) , no effect ( )

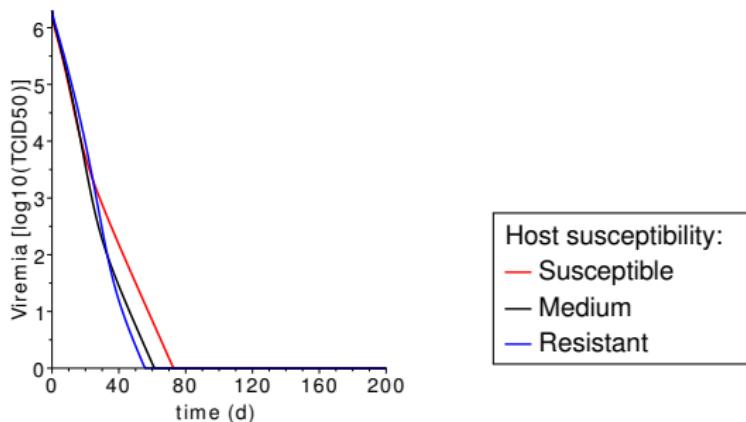
	Virus	% infection	% phagocytosis	NK	cellular R
Inoculation $V_0$	++				
Excretion $e$	++	++	--		++
Infectiosity $\beta$		++	--	++	++
Phagocytosis $\eta$	--		++		



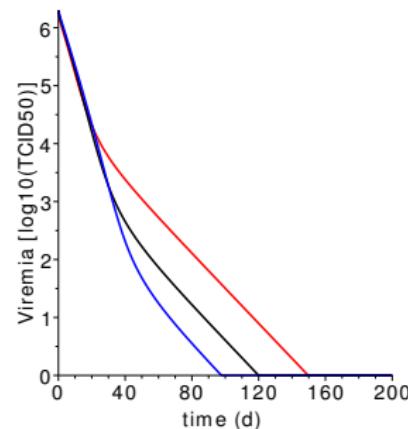
## HOST SUSCEPTIBILITY AND VIRAL VIRULENCE

**Host susceptibility**  $\leftrightarrow$  capacities of cytokine synthesis by immune cells

**Viral virulence**  $\leftrightarrow$  excretion and phagocytosis rate



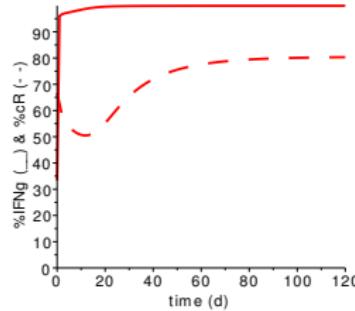
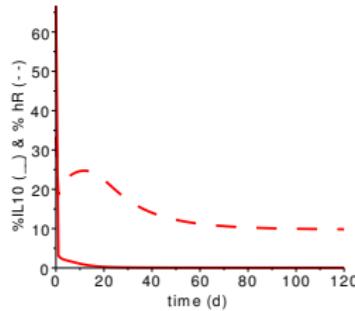
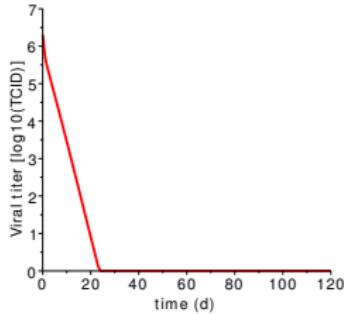
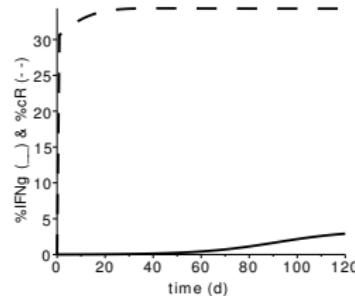
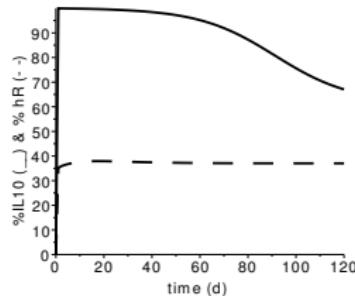
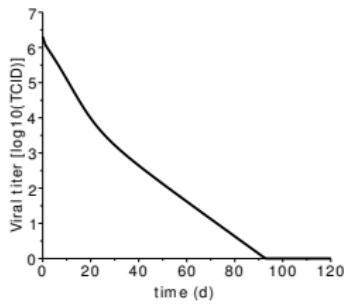
**Figure:** infection by viral strain with low virulence



**Figure:** infection by viral strain with high virulence

## VACCINATION EFFECT

Comparison of the infection dynamic without vaccination (—) and with vaccination (—)



## CONCLUSIONS & PERSPECTIVES

- **An original model** thanks to the detailed macrophage – virus interactions and the cytokine effects.
- **A model tool:** we can vary the parameter values to
  - a) explore the infection dynamic depending on viral strain and host susceptibility
  - b) test hypothesis (vaccination, ...) to improve infection resolution
  - c) explore the respective influence of immune mechanisms on viral clearance (not presented here)
- **Perspectives:** model extension to the individual scale

# THANK YOU FOR YOUR ATTENTION !

## *References*

### **(1)** Planor library :

MONOD H., KOBELINSKY A., AND BOUVIER A. : Automatic generation of regular factorial designs : the PLANOR R library. (2011)

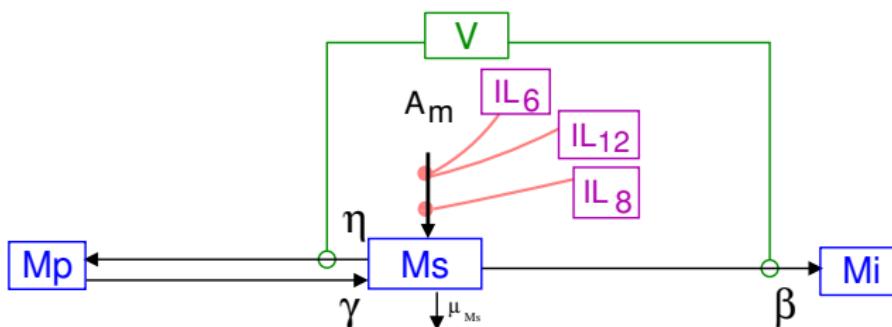
<http://hal.archives-ouvertes.fr/docs/00/71/75/56/ANNEX/planorVignette.pdf>

### **(2)** Multisensi Package :

LAMBONI M. AND MONOD H. : Package 'multisensi' (2012).

<http://cran.r-project.org/web/packages/multisensi/index.html>

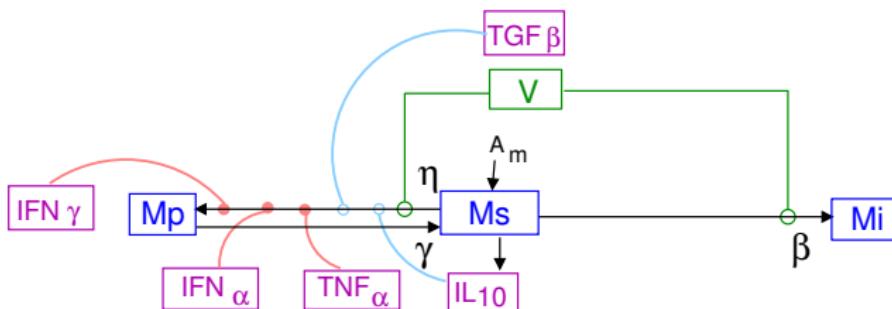
## EXAMPLE: MACROPHAGES RECRUITMENT



$$\dot{M}_s = A_m (1 + \kappa (IL_{12}, IL_6)) (1 + \kappa (IL_8)) \quad \text{macrophage recruitment}$$

- > *activatory* cytokines: additive effect on the basic rate  $A_m$
- >  $\kappa$  : Michaëlis-Menten function of the cytokine concentration  
(limited number of cytokine receptors)

## EXAMPLE: PHAGOCYTOSIS START

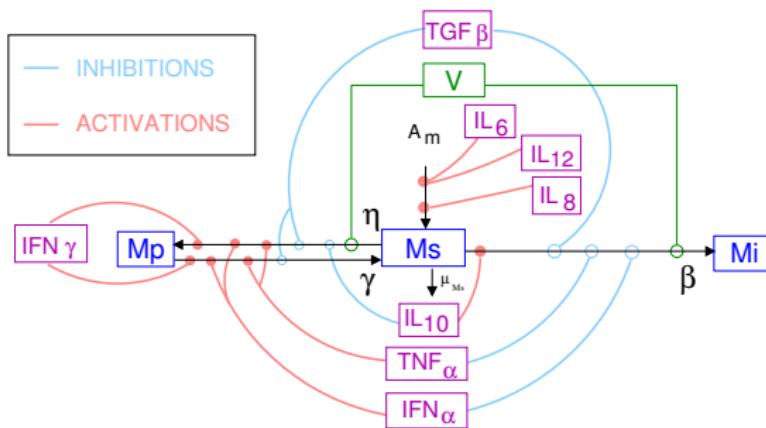


$\dot{M}_s$  = macrophage recruitment

$$-\eta M_s V \frac{(1 + \kappa(TNF_\alpha))(1 + \kappa(IFN_\gamma))(1 + \kappa(IFN_\alpha))}{(1 + \kappa(IL10))(1 + \kappa(TGF_\beta))} \quad \text{phagocytosis start}$$

- > *Inhibitory* cytokines: symmetric effect

## EXAMPLE: SUSCEPTIBLE MACROPHAGES DYNAMIC



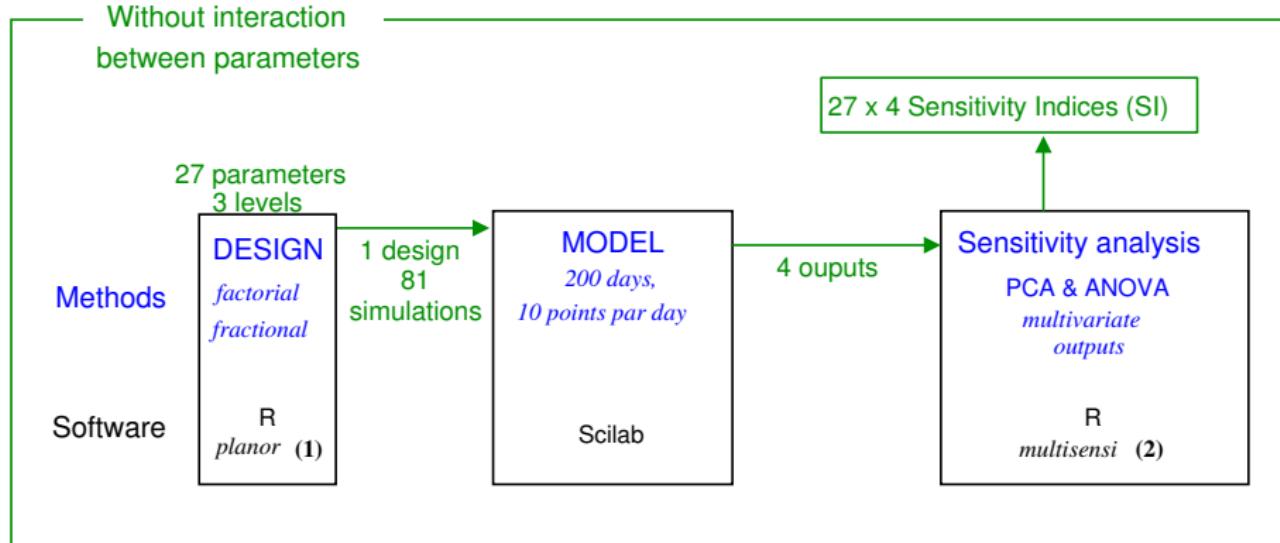
$\dot{M}_s$  = macrophage recruitment

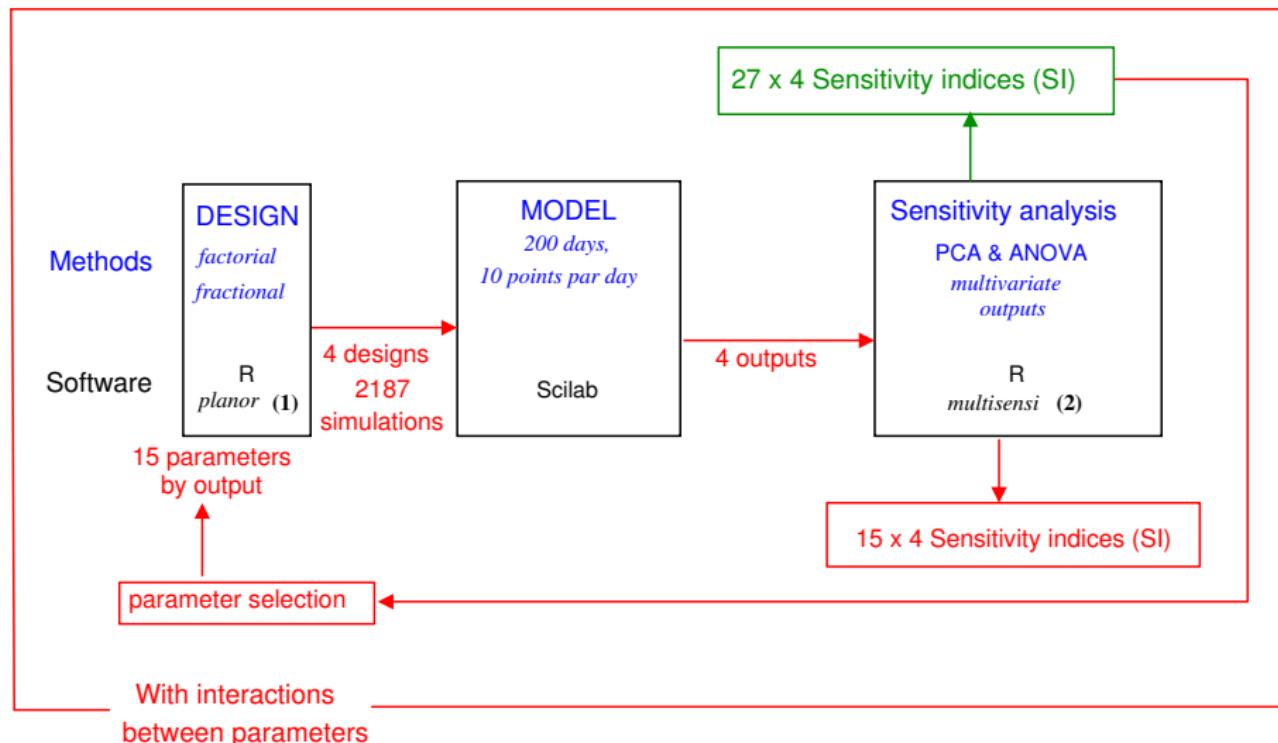
— phagocytosis start

+ phagocytosis end

— infection

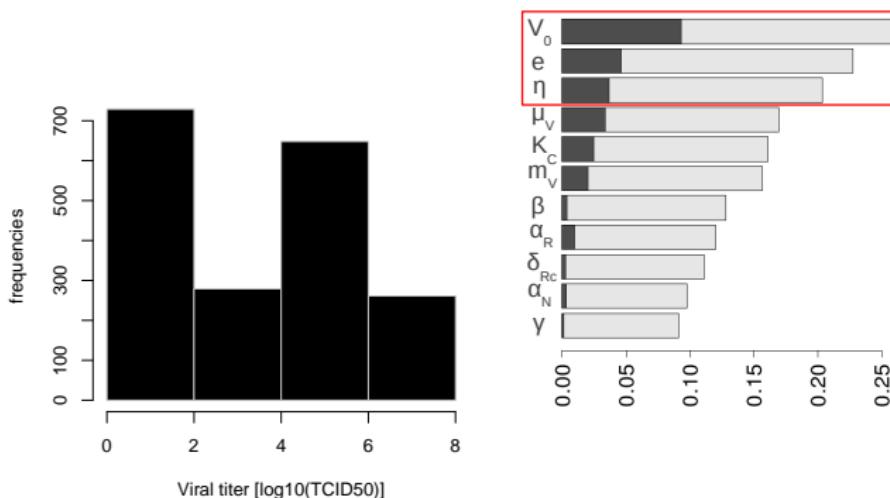
— macrophage death





## SA RESULTS : VIRUS

*A single virus inoculation at the beginning of the simulations (V0)*



## SA RESULTS : VIRUS

