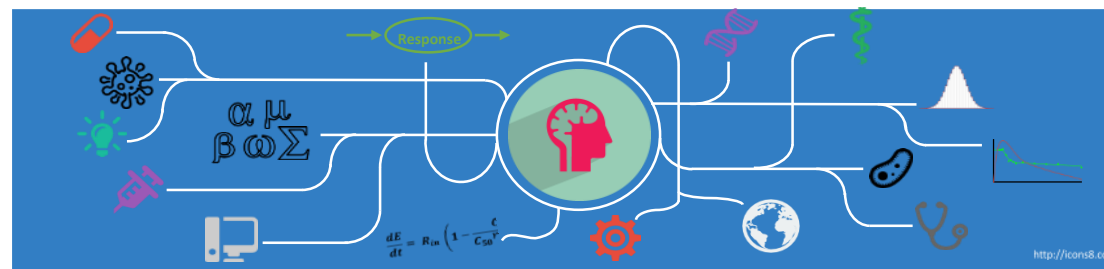


Ebola viral dynamics in nonhuman primates: insights into virus immuno-pathogenesis and antiviral strategies

V. Madelain¹, S. Baize², F. Jacquot³, C. Carbonnelle³, H. Raoul³,
X. de Lamballerie⁴, F. Mentré¹, J. Guedj¹

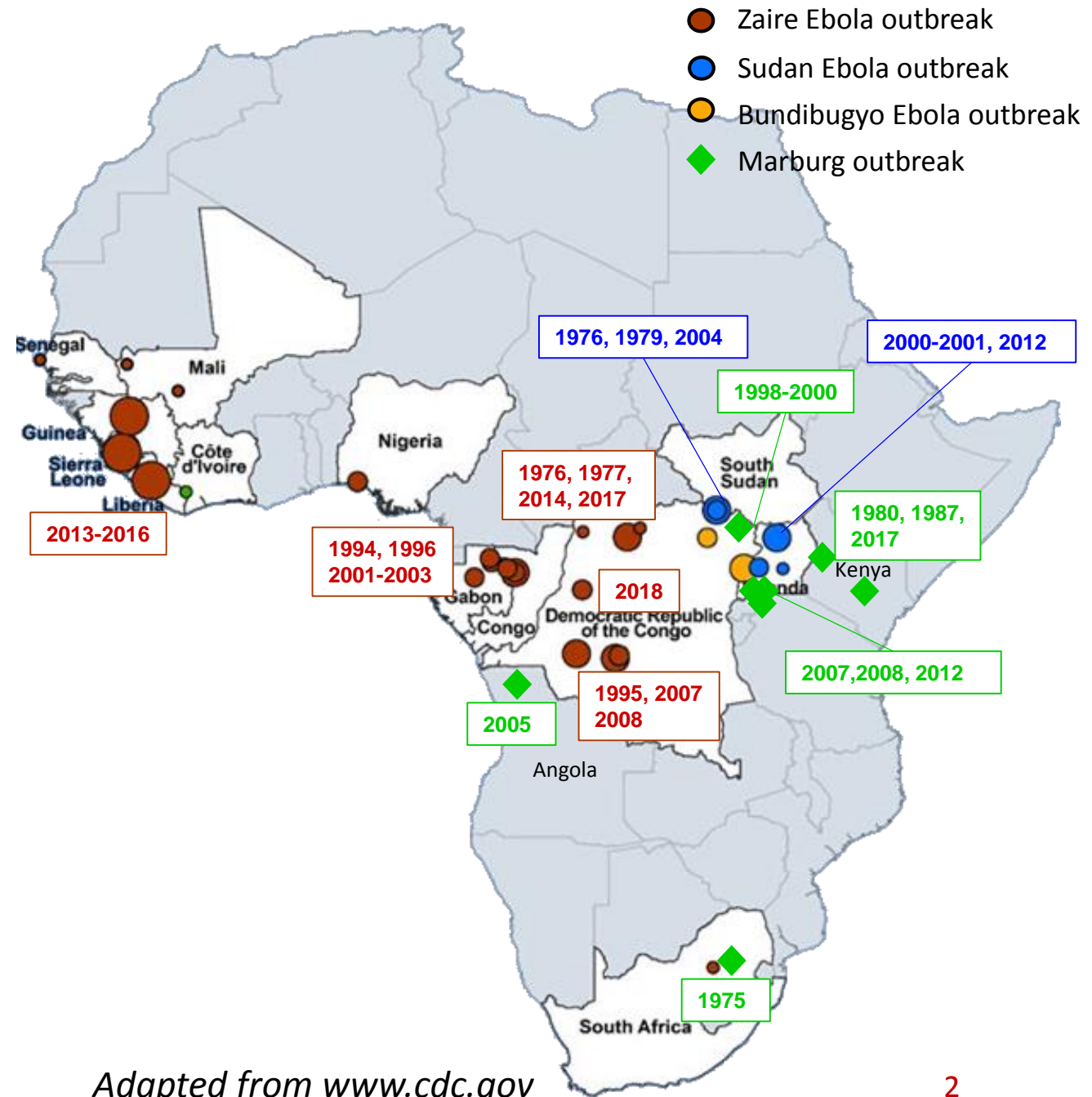
¹IAME, UMR 1137, INSERM – Paris Diderot University ²UBIVE, Institut Pasteur ³Laboratoire P4
Inserm, France ⁴UMR EPV Aix-Marseille University - IRD 190 - Inserm 1207 - EHESP

Page Meeting, June 1st 2018



Background

- Hemorrhagic fevers represent a constant threat to public health in Africa and beyond
- The 2013-2016 Ebola outbreak in West Africa confirmed its potential to develop into regional epidemics
- None of the antiviral drug tested could demonstrate a significant effect on survival rate



Adapted from www.cdc.gov

Background

- Hemorrhagic fevers represent a constant threat to public health in Africa and beyond
- The 2013-2016 Ebola outbreak in West Africa confirmed its potential to develop into regional epidemics
- None of the antiviral drug tested could demonstrate a significant effect on survival rate

Sissoko et al, Plos Med, 2016

Prevail study group, NEJM, 2016

Dunning et al, Plos Med, 2016

Dunning et al, Plos One, 2016

OPEN ACCESS

Citation: Sissoko D, Laouenan C, Folkesson E, M'Lebing A-B, Beavogui A-H, Balze S, et al. (2016) Experimental Treatment with Favipiravir for Ebola Virus Disease (the JIKI Trial): A Historically Controlled, Single-Arm Proof-of-Concept Trial in Guinea. *PLoS Med* 13(3): e1001967. doi:10.1371/journal.pmed.1001967

Academic Editor: Marc Lipsitch, Harvard School of Public Health, UNITED STATES

Received: September 18, 2015

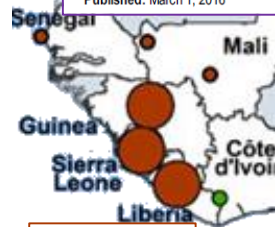
Accepted: January 21, 2016

Published: March 1, 2016

RESEARCH ARTICLE

Experimental Treatment with Favipiravir for Ebola Virus Disease (the JIKI Trial): A Historically Controlled, Single-Arm Proof-of-Concept Trial in Guinea

The NEW ENGLAND JOURNAL of MEDICINE



2013-2016

ORIGINAL ARTICLE

A Randomized, Controlled Trial of ZMapp for Ebola Virus Infection

The PREVAIL II Writing Group, for the Multi-National PREVAIL II Study Team*

OPEN ACCESS

Citation: Dunning J, Sahr F, Rajak A, Gannon F, Carson G, Idriss B, et al. (2016) Experimental Treatment of Ebola Virus Disease with TKM-130803: A Single-Arm Phase 2 Clinical Trial. *PLoS Med* 13(4): e1001997. doi:10.1371/journal.pmed.1001997

Academic Editor: Lorenz von Seiden, Mahidol-Oxford Tropical Medicine Research Unit, THAILAND

Received: October 9, 2015

Accepted: March 8, 2016

Published: April 19, 2016

RESEARCH ARTICLE

Experimental Treatment of Ebola Virus Disease with TKM-130803: A Single-Arm Phase 2 Clinical Trial

2007, 2008, 2012

OPEN ACCESS

Citation: Dunning J, Kennedy SB, Antierens A, Whitehead J, Ciglenecki I, Carson G, et al. (2016) Experimental Treatment of Ebola Virus Disease with Brincidofovir. *PLoS ONE* 11(9): e0162199. doi:10.1371/journal.pone.0162199

Editor: John W Glod, National Cancer Institute, UNITED STATES

Received: April 25, 2016

Accepted: August 15, 2016

Published: September 9, 2016

RESEARCH ARTICLE

Experimental Treatment of Ebola Virus Disease with Brincidofovir

Jake Dunning^{1‡}, Stephen B. Kennedy^{2‡}, Annick Antierens³, John Whitehead⁴, Iza Ciglenecki⁵, Gail Carson¹, Rupa Kanapathipilla², Lyndsey Castle¹, Rebecca Howell-Jones¹, Raul Pardinaz-Solis¹, Jennifer Grove¹, Janet Scott⁶, Trudie Lang¹, Piero Olliaro^{1,7}, Peter W. Horby^{1*}, for the RAPIDE-BCV trial team[†]

Adapted

ic.gov

ik
outbreak

1, 2012



Background

- Most of these candidate treatments, and others, demonstrated *in vivo* antiviral effect, and were found to significantly improve survival in NHP experiments

Warren et al, Nature, 2014

Qiu et al, Nature, 2014

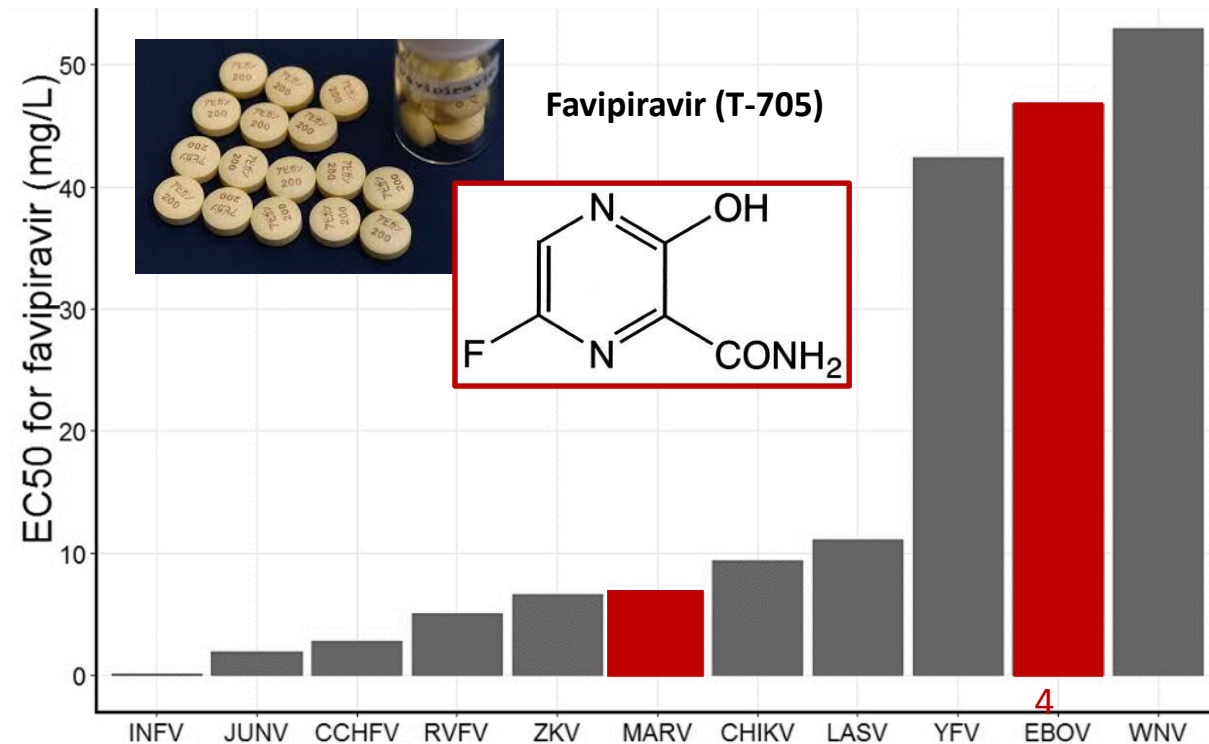
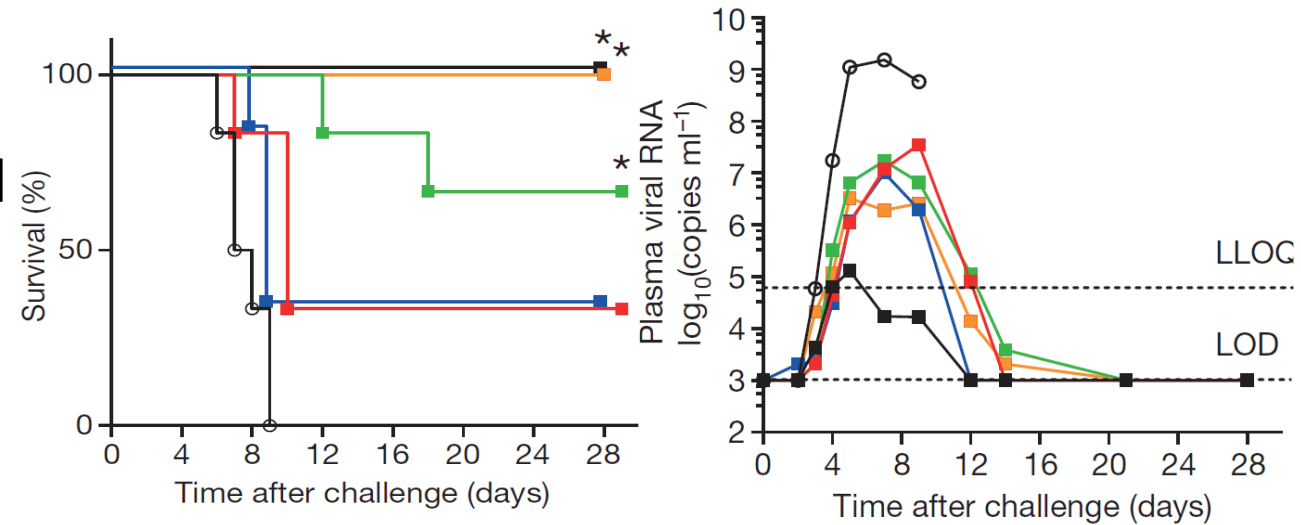
Thi et al, Nature, 2015

Warren et al, Nature, 2016

- Our group focused on favipiravir, a RNA polymerase inhibitor approved for influenza, with activity against various etiological agents of hemorrhagic fevers

Oestereich et al, Antiviral res, 2014

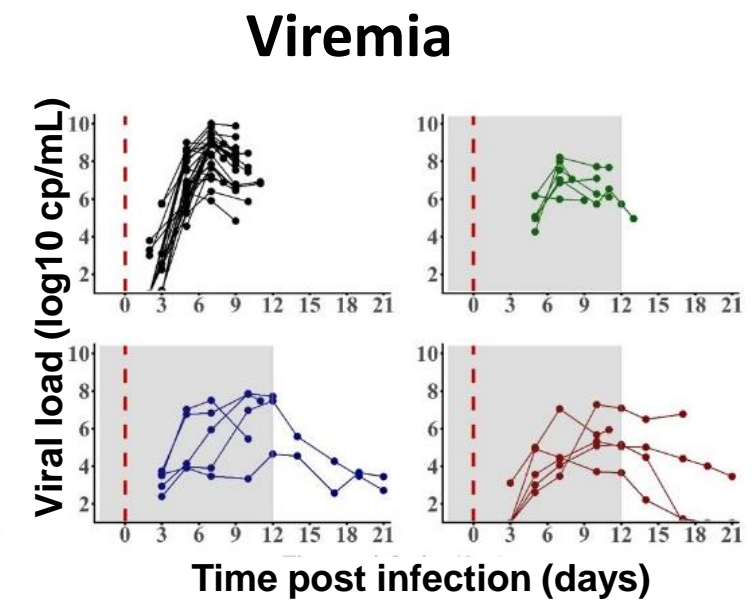
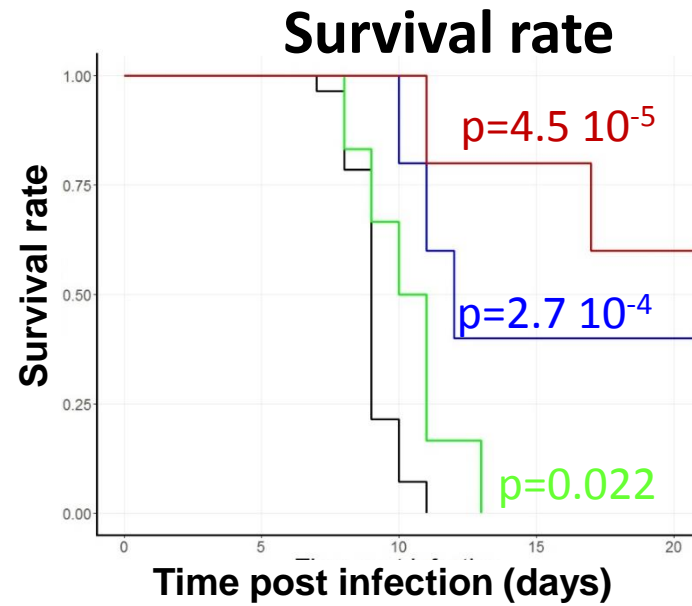
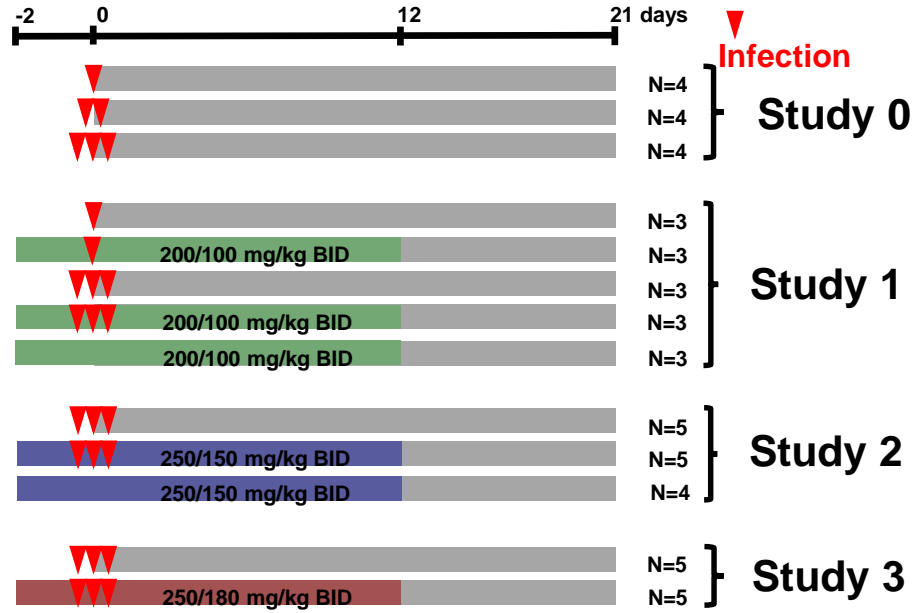
Guedj et al, Plos med, 2018



Objectives

- To develop a mathematical mechanistic model to characterize the pathogenesis and the determinants of the death of Ebola virus disease (EVD) in NHPs
- To assess the activity of favipiravir in EVD
- To predict the impact of treatment potency and timing of initiation on survival rate

Study design and data

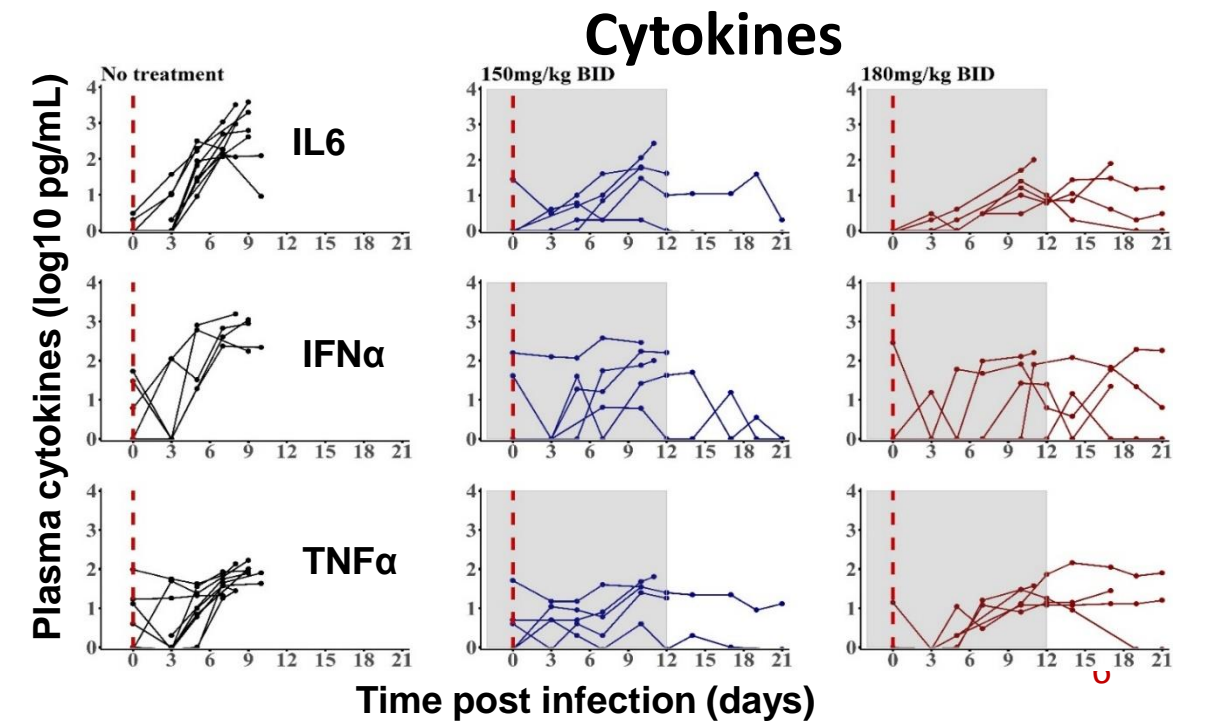
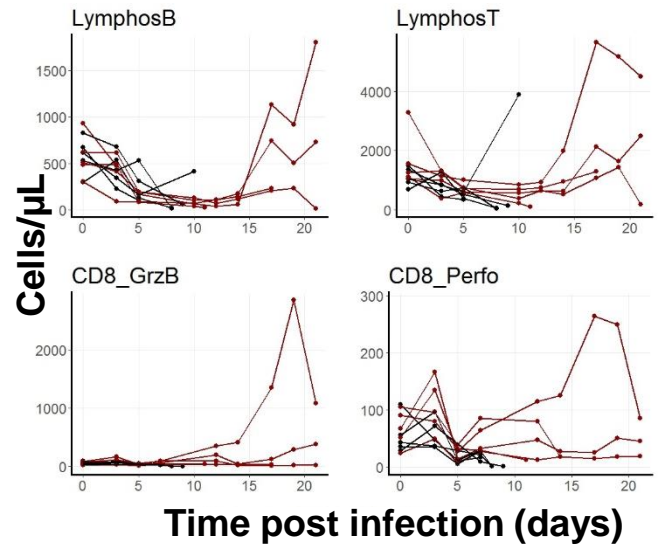


Guedj et al, Plos med, 2018

Dosing

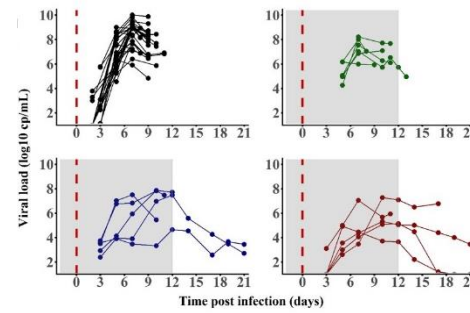
- No treatment
- 100 mg/kg BID
- 150 mg/kg BID
- 180 mg/kg BID

Cytometry



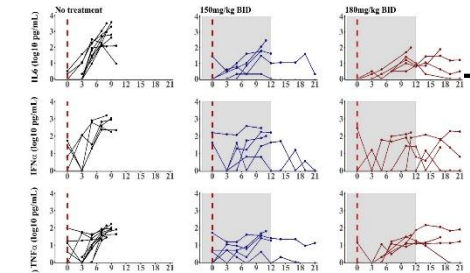
Model building strategy

Parameter estimation was performed using the SAEM algorithm implemented in Monolix software



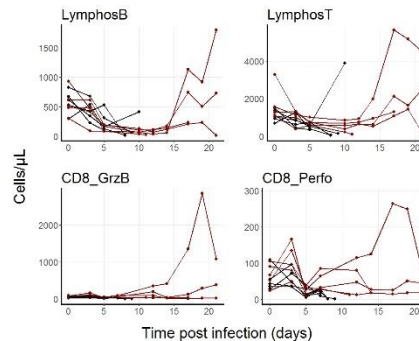
Viremia data

Target cell limited model



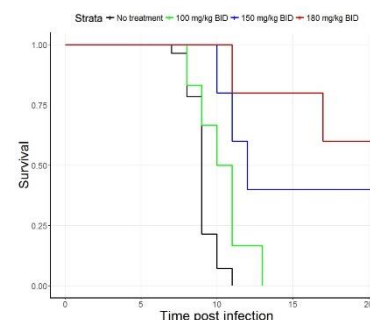
Cytokine data

Innate response model to describe the early stage of the infection



Cytometry data

Adaptive response model to describe the late stage of the infection



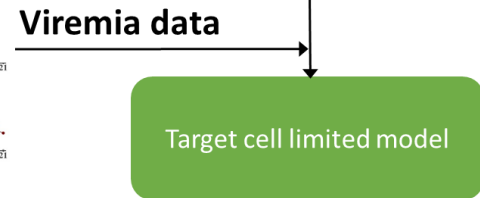
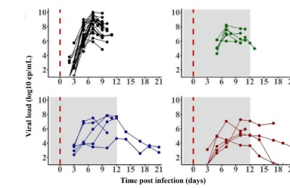
Time to death

Joint model to assess the impact of viral load and cytokine dynamics on survival

Improvement of viremia OFV

Target cell limited model

- Assumes that viral control results from target cell depletion
- Favipiravir concentrations are described by a previously developed PK model and inhibits viral production



$$\frac{dT}{dt} = -\beta TV$$

$$\frac{dI_1}{dt} = \beta TV - kI_1$$

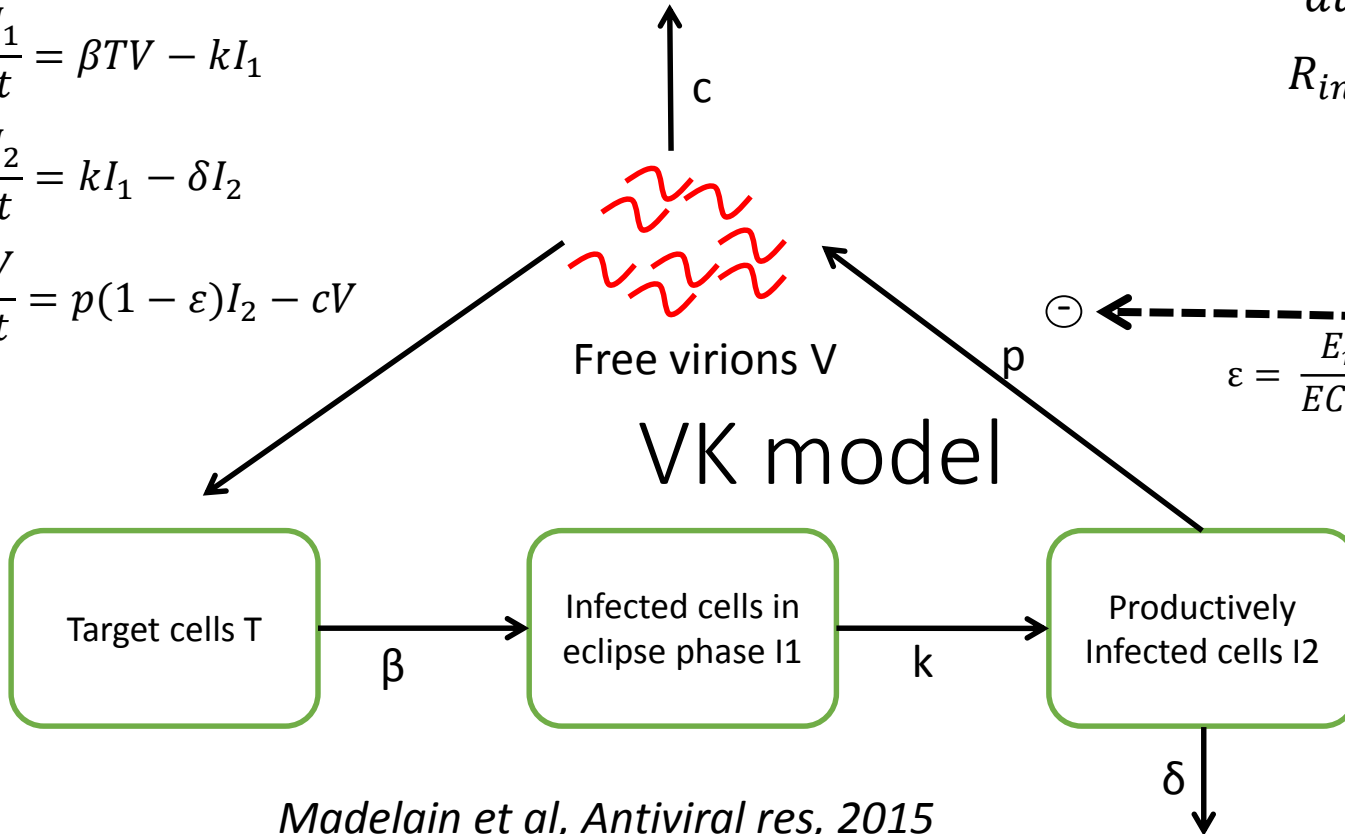
$$\frac{dI_2}{dt} = kI_1 - \delta I_2$$

$$\frac{dV}{dt} = p(1 - \varepsilon)I_2 - cV$$

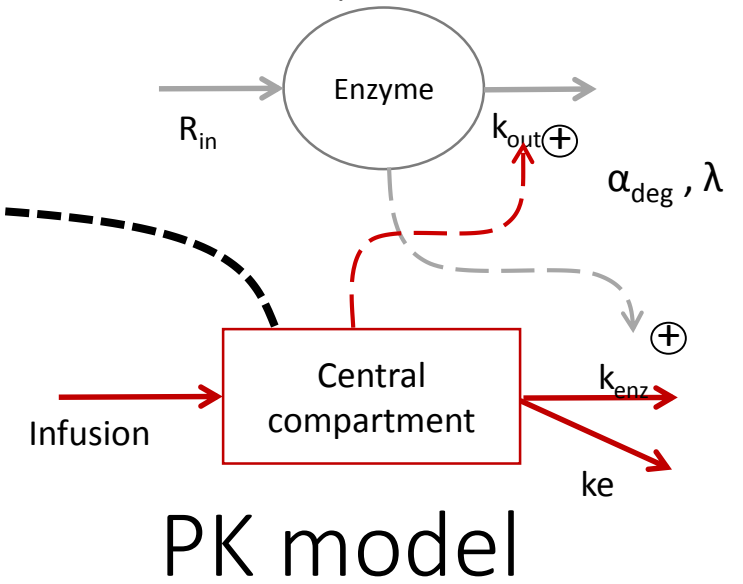
$$\frac{dA_c}{dt} = -k \times A_c - V_{max} \times A_e \times A_c$$

$$\frac{dA_e}{dt} = R_{in} - k_{out} \times (1 + C_c \times k_{el} \times e^{-\lambda \times t}) \times A_e$$

$$R_{in} = k_{out} \times A_{e0} \quad ; \quad C_c = \frac{A_c}{V}$$



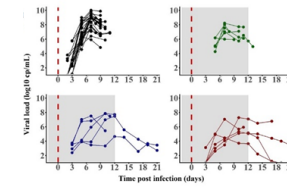
Madelain et al, Antiviral res, 2015



$$\varepsilon = \frac{E_{max} C_c}{EC_{50} + C_c}$$

Madelain et al, AAC, 2017

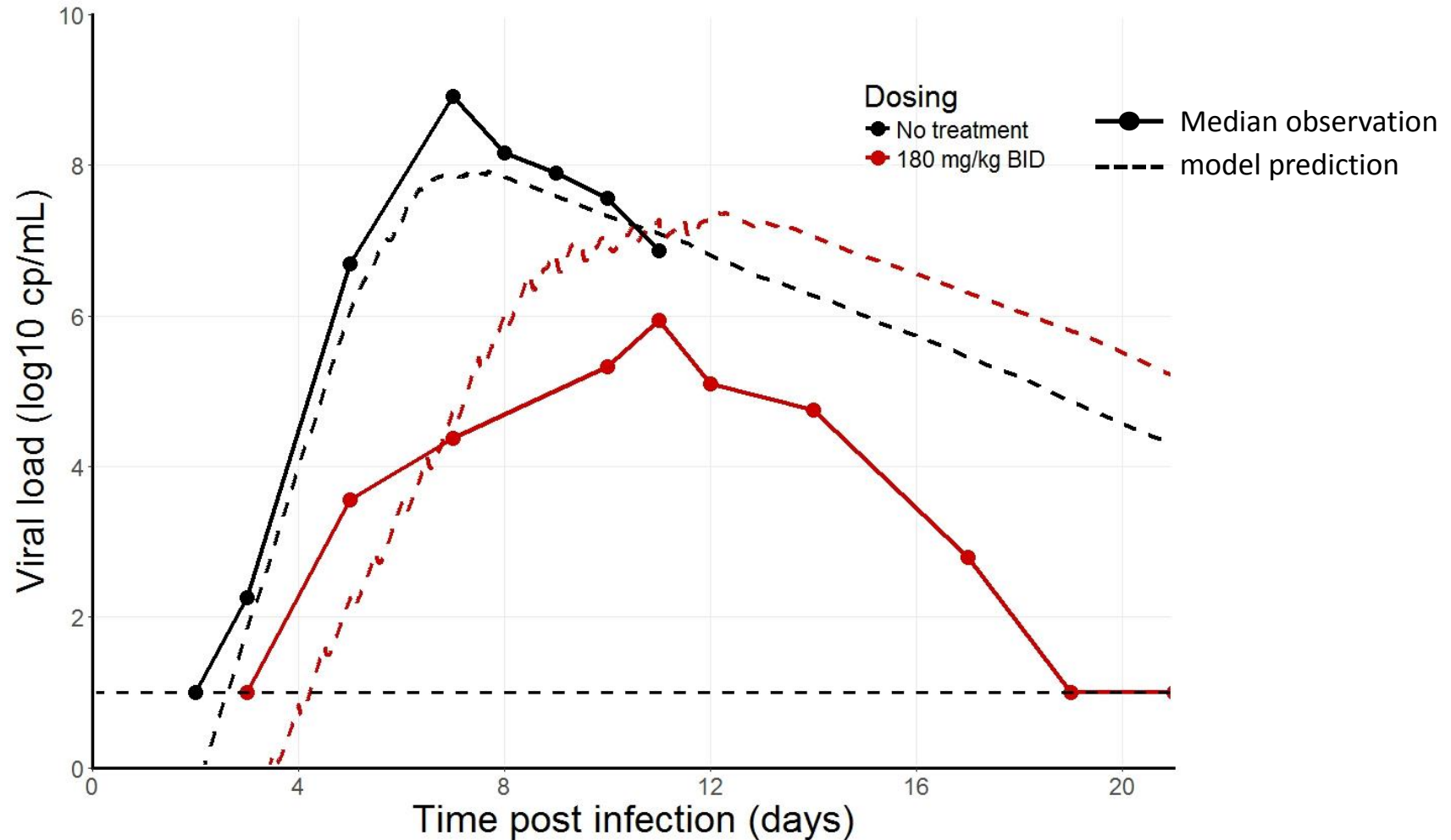
Target cell limited model



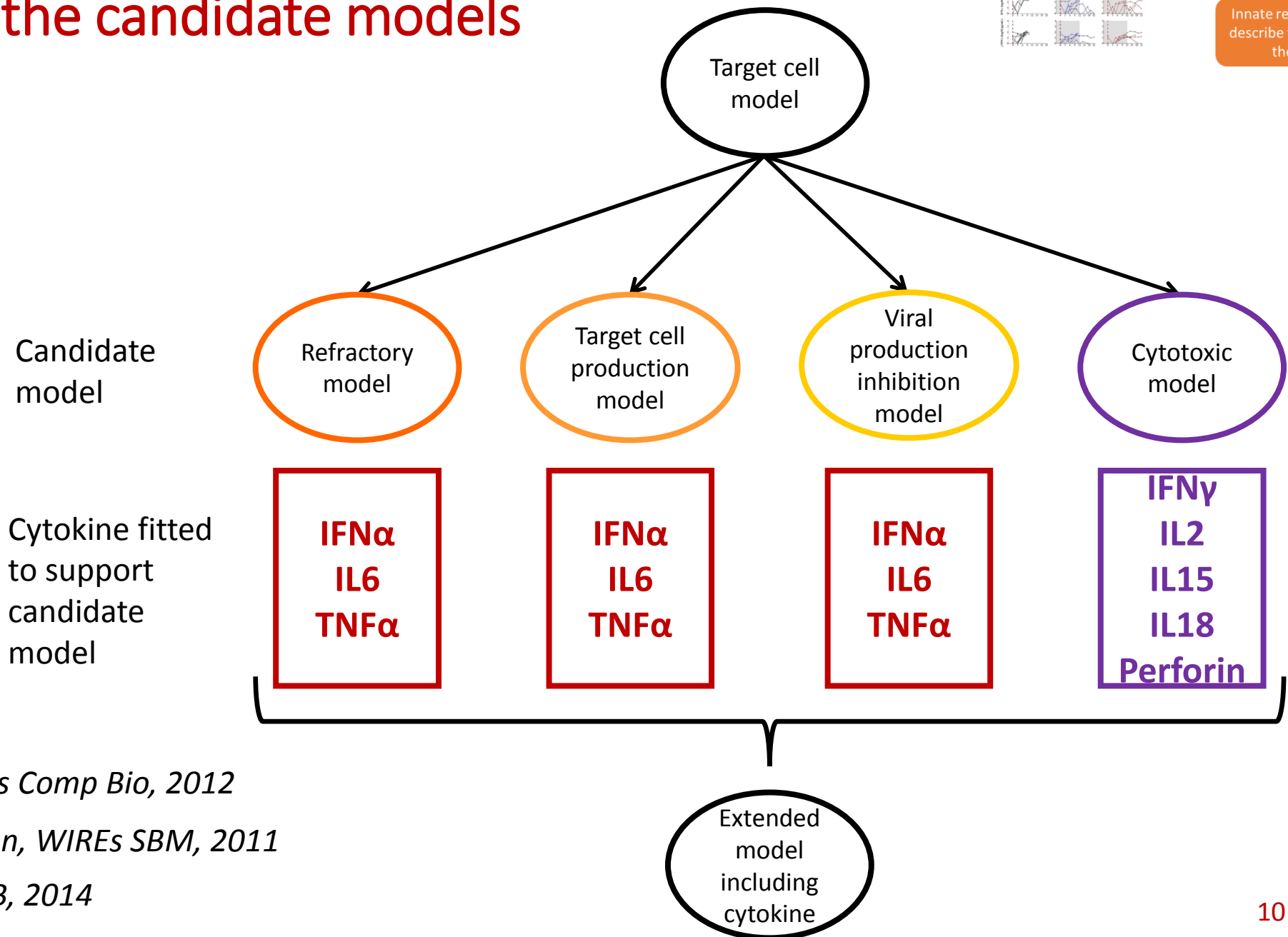
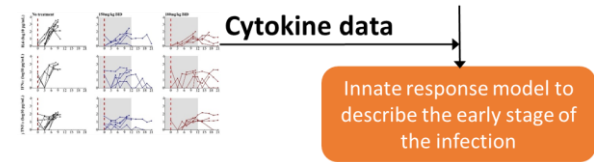
Viremia data

Target cell limited model

- Simulations showed that the model underestimates the effect of treatment on peak viremia



Evaluation of the candidate models



Pawelek et al, Plos Comp Bio, 2012

Smith and Perelson, WIREs SBM, 2011

Li and Handel, JTB, 2014

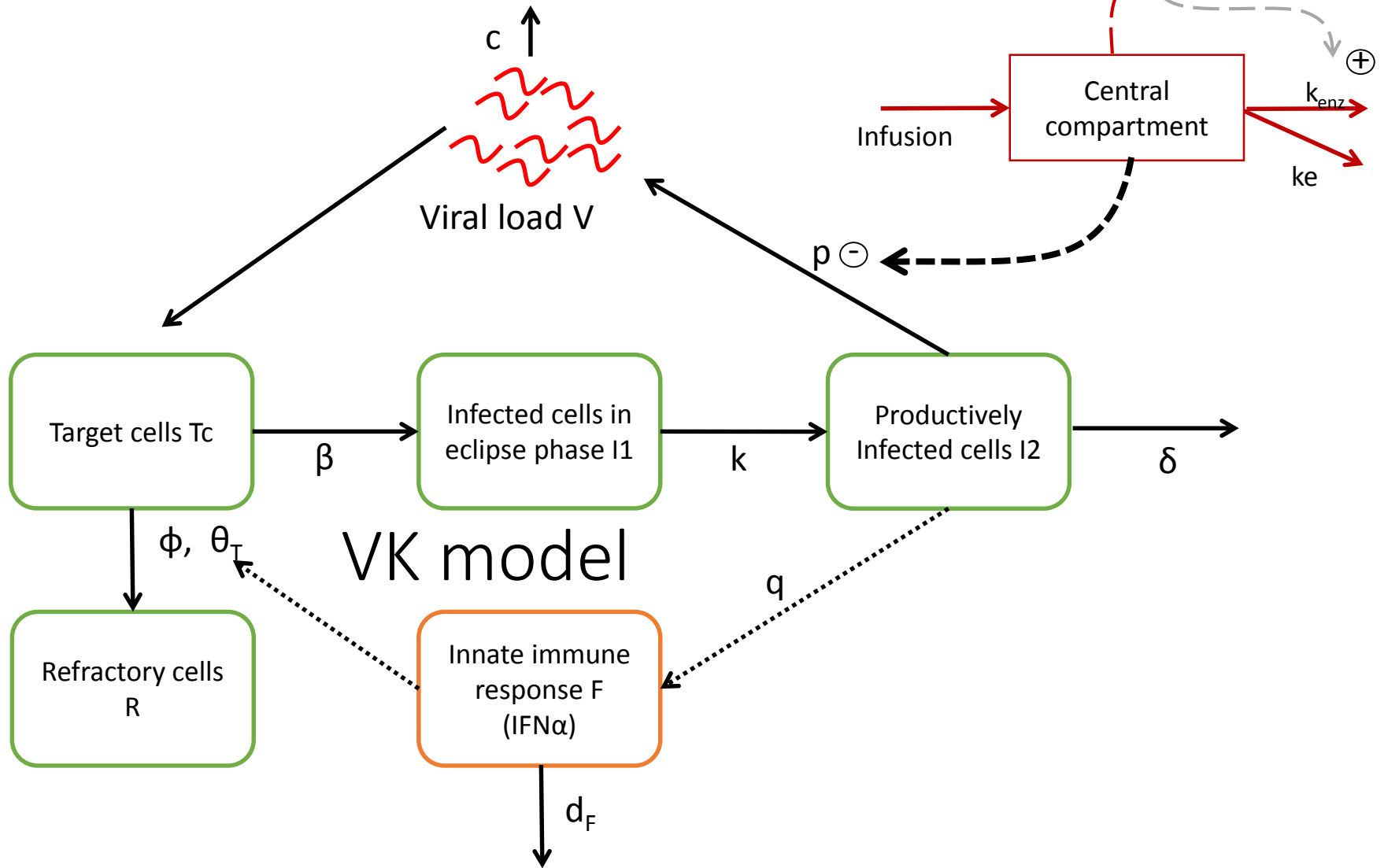
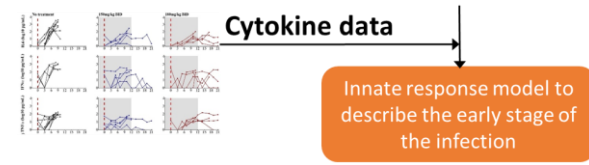
Selection of the candidate models



| cytokine | model | effect of cytokine | -2logL viremia | Additional parameters |
|-------------------------------|----------------------|--|----------------|-----------------------|
| None (step 1) | target cell limited | none | 645.7 | – |
| IL6 | refractory | increase refractory cell production | 618.0 | 4 |
| IL6 | target cell increase | increase of target cells | 644.8 | 4 |
| IL6 | prod inhibition | non linear inhibition of viral production | 645.7 | 4 |
| IFNα | refractory | increase refractory cell production | 622.4 | 4 |
| IFN α | target cell increase | increase of target cells | 641.8 | 4 |
| IFN α | prod inhibition | non linear inhibition of viral production | 692.8 | 4 |
| TNFα | refractory | increase refractory cell production | 621.8 | 4 |
| TNF α | target cell increase | increase of target cells | 635.1 | 4 |
| TNF α | prod inhibition | non linear inhibition of viral production | 633.3 | 4 |
| IFN γ | cytotox | increase infected cell elimination | 634.5 | 4 |
| IL2 | cytotox | increase infected cell elimination | 636.3 | 5 |
| perforin | cytotox | increase infected cell elimination | 634.6 | 4 |
| IL15 | cytotox | increase infected cell elimination | 634.6 | 5 |
| IL18 | cytotox | increase infected cell elimination | 634.8 | 4 |

→ Refractory models relying on IFN α , IL6 and TNF α and assuming a conversion of target cells into refractory cells provided the best description of viremia

Model including IFN α dynamics



$$\frac{dT}{dt} = -\beta TV - \frac{\phi TF}{F + \theta_T}$$

$$\frac{dI_1}{dt} = \beta TV - kI_1$$

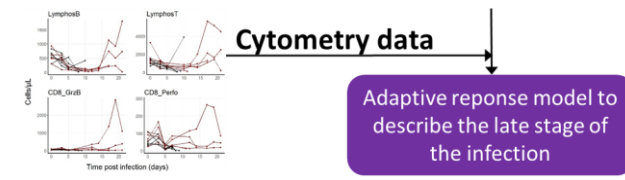
$$\frac{dI_2}{dt} = kI_1 - \delta I_2$$

$$\frac{dR}{dt} = \frac{\phi TF}{F + \theta_T}$$

$$\frac{dV}{dt} = p(1 - \varepsilon)I_2 - cV$$

$$\frac{dF}{dt} = qI_2 - d_F F$$

Selection of the candidate models

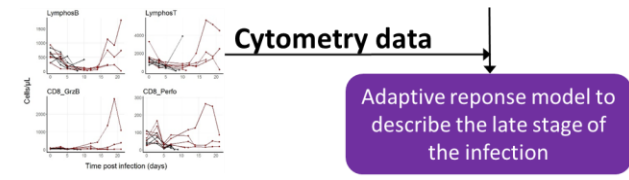
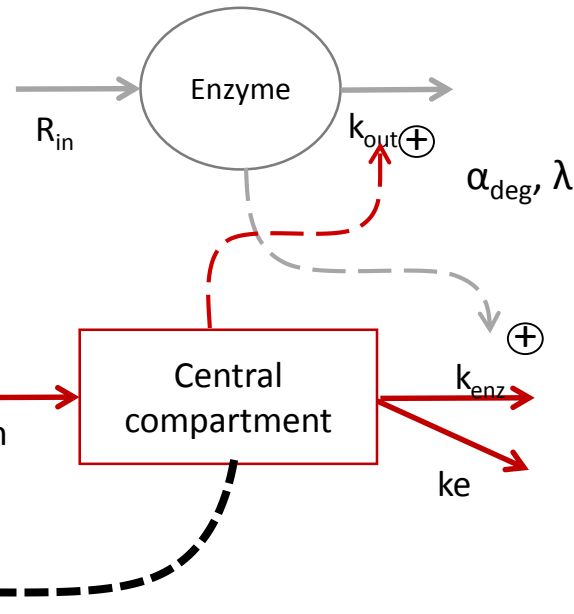
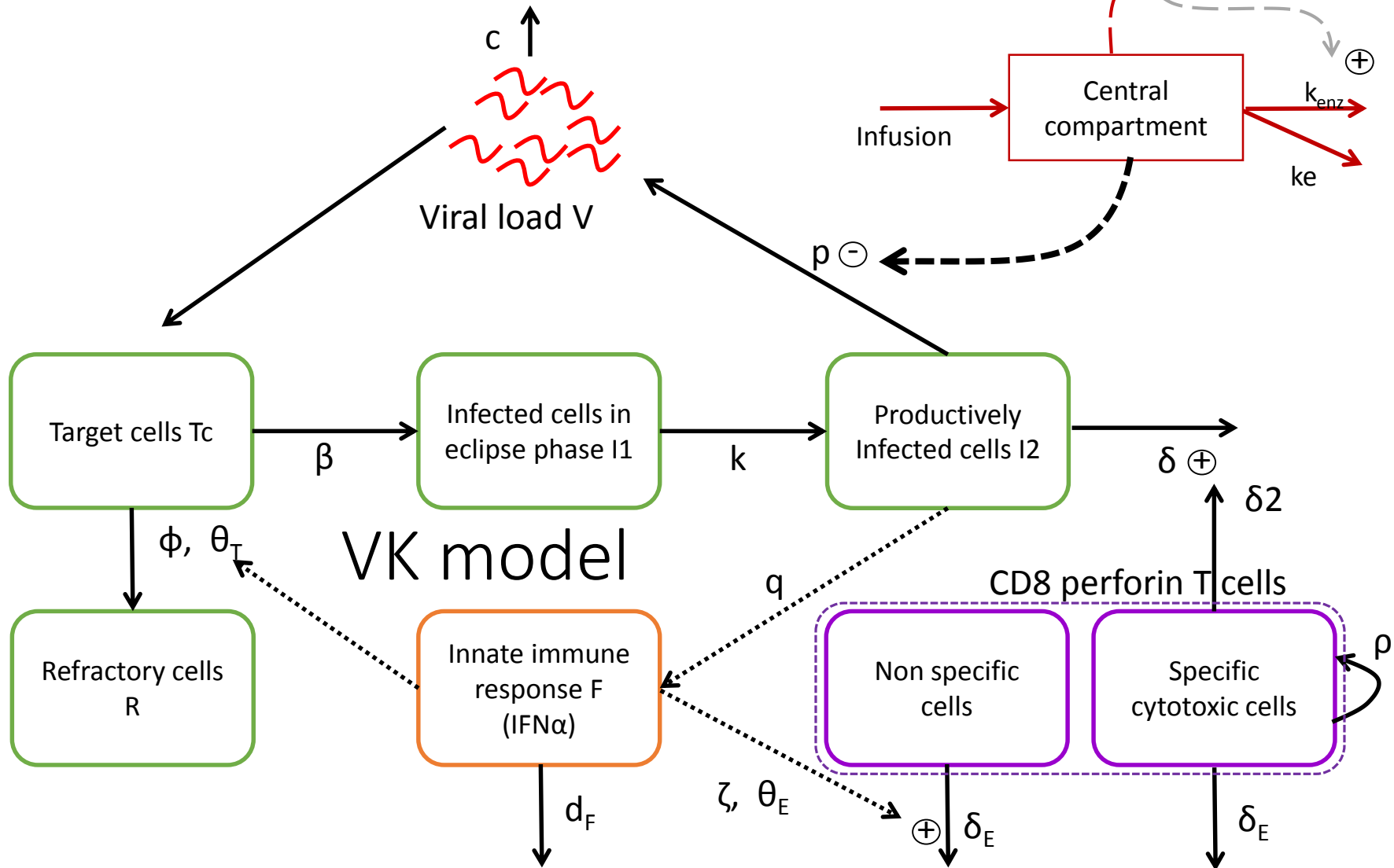


Adaptive response models including CD8 T cells expressing cytotoxic activity NKp80, perforin and granzymeB were compared

| model | CD8 T cell data | -2logL viremia | Additional parameters |
|------------------------------|----------------------|----------------|-----------------------|
| target cell limited (step 1) | – | 645.7 | – |
| Refractory (step 2) | – | 618 | 4 |
| refractory | CD8 perforin+ | 605.5 | 12 |
| refractory | CD8 granzymeB+ | 611.6 | 12 |
| refractory | CD8 NKp80+ | 608.4 | 12 |

→ CD8 T cell population expressing perforin was selected and included

Model including CD8 dynamics



$$\frac{dT}{dt} = -\beta TV - \frac{\phi TF}{F + \theta_T}$$

$$\frac{dI_1}{dt} = \beta TV - kI_1$$

$$\frac{dI_2}{dt} = kI_1 - \delta I_2 - \kappa I_2 E_2$$

$$\frac{dR}{dt} = \frac{\phi TF}{F + \theta_T}$$

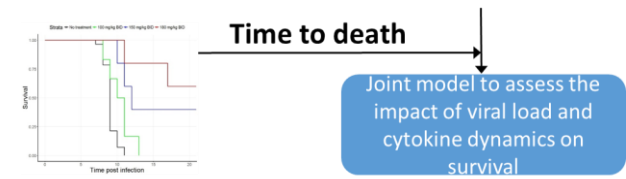
$$\frac{dV}{dt} = p(1 - \epsilon)I_2 - cV$$

$$\frac{dF}{dt} = qI_2 - d_F F$$

$$\frac{dE_1}{dt} = \sigma - \frac{\zeta FE_1}{F + \theta_E} - \delta_E E_1$$

$$\frac{dE_2}{dt} = \rho E_2 \left(1 - \frac{E_2}{E_0}\right) - \delta_E E_2$$

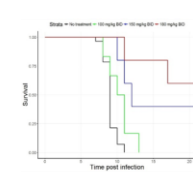
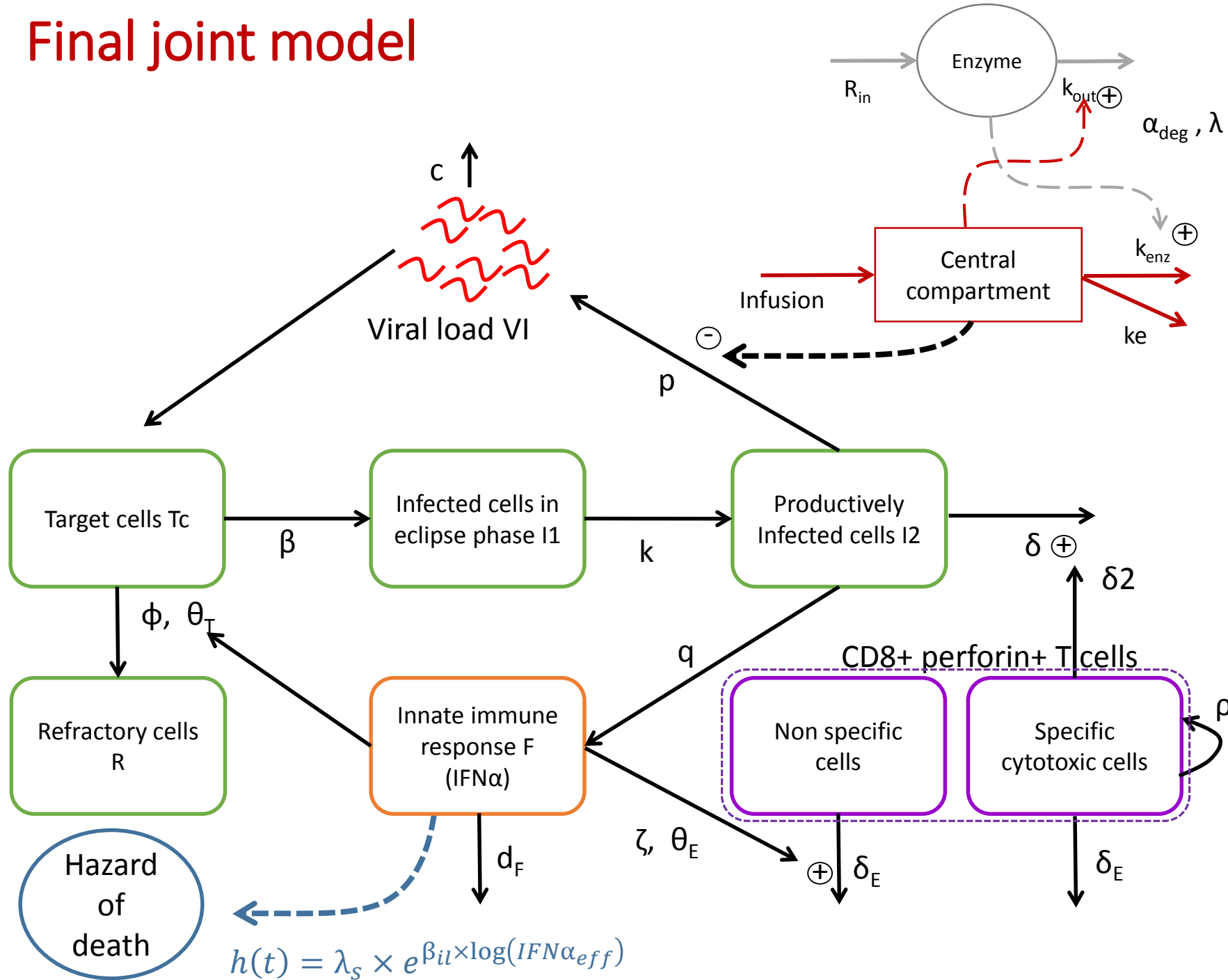
Impact of viral and cytokine dynamics on survival



Extension to a joint model to assess the impact of viremia and cytokines on the hazard rate, where:

- $h(t)$ is the hazard rate : $h(t) = \lambda_0 \times e^{\sum_k \beta_k \times \log(X_k(t))}$
where $X_k(t)$ is current or lag-value of viral load, IL6, IFN α or TNF α
- $S(t)$ is the probability to be alive up to time t : $S(t) = e^{-\int_0^t h(u) du}$

Final joint model



Time to death

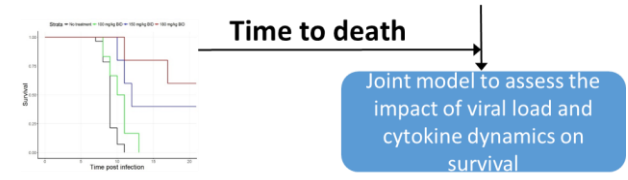
Joint model to assess the impact of viral load and cytokine dynamics on survival

| Model compartment driving the hazard of death | BIC |
|---|---------------|
| viremia | 1545.0 |
| IL6 | 1506.5 |
| IFNα | 1506.5 |
| TNF α | 1526.8 |

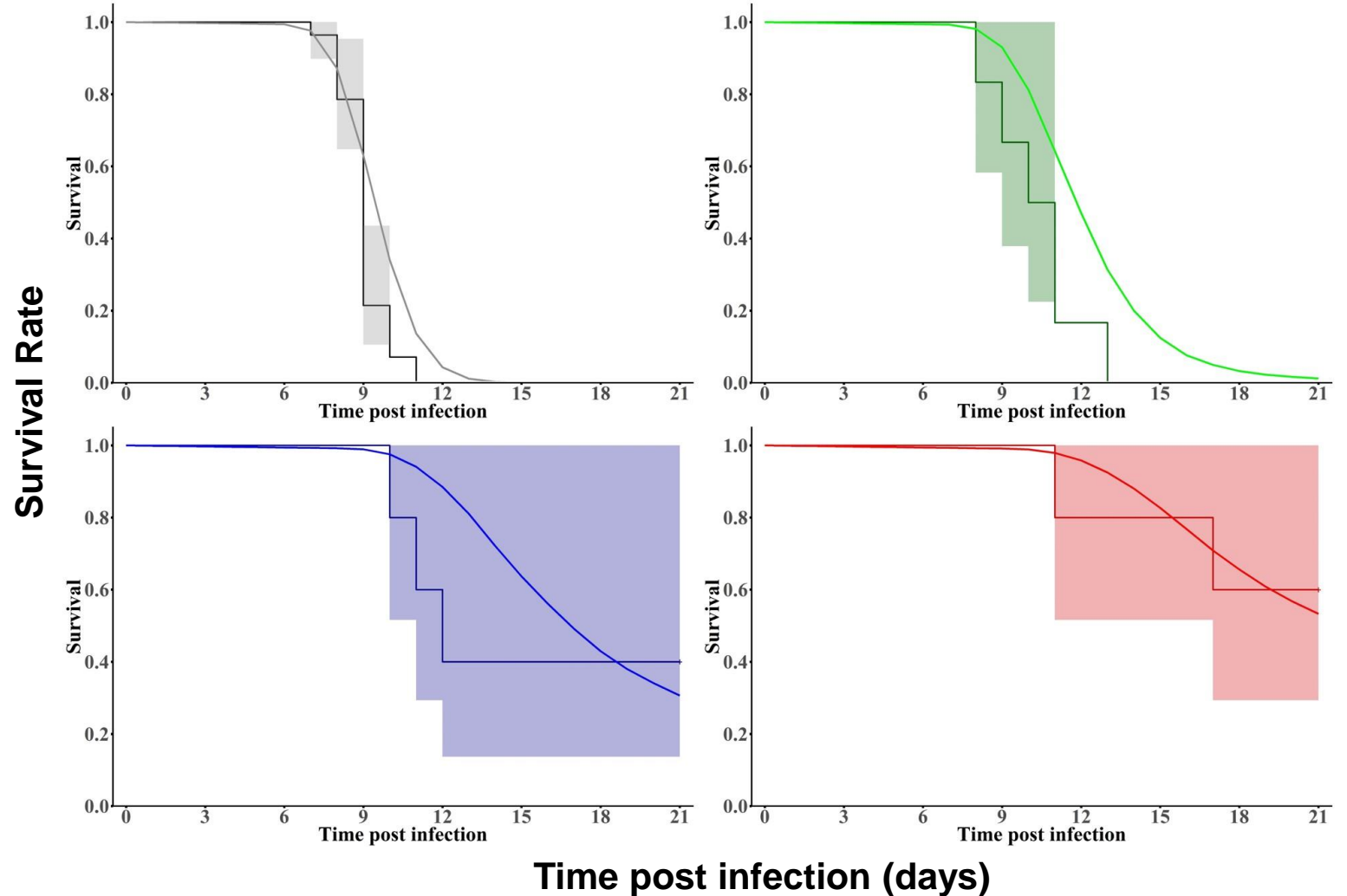
Best fit to the survival data was obtained with a model including IFN α or IL6

$$h(t) = \lambda_s \times e^{\beta_{il} \times \log(IFN\alpha_{eff})}$$

Model predictions of survival rate



- The joint model recapitulates the survival rate at day 21 in each treatment group



Validation of model predictions using remdesivir data

Warren et al, Nature, 2016

LETTER

doi:10.1038/nature17180

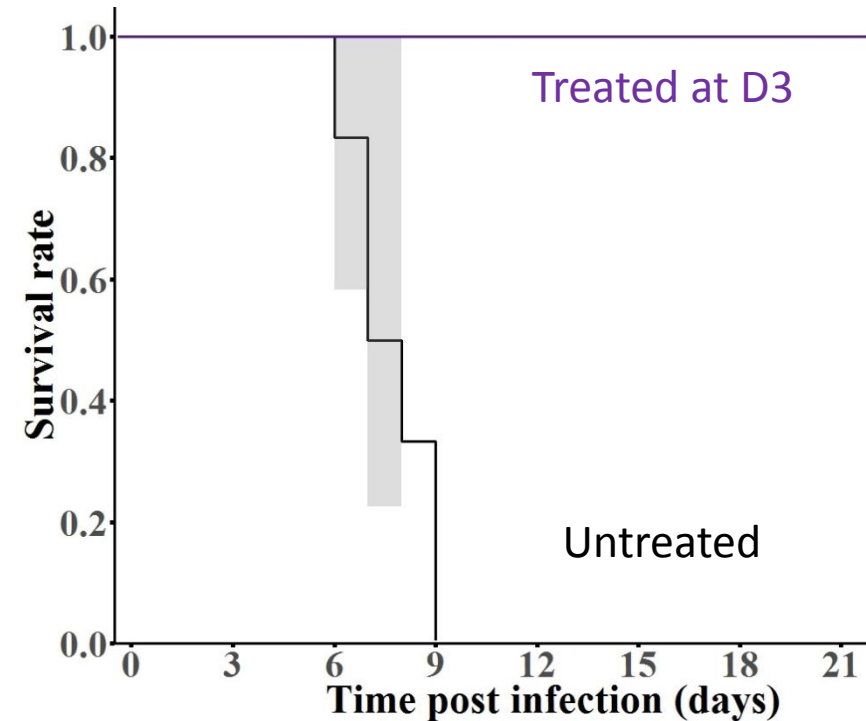
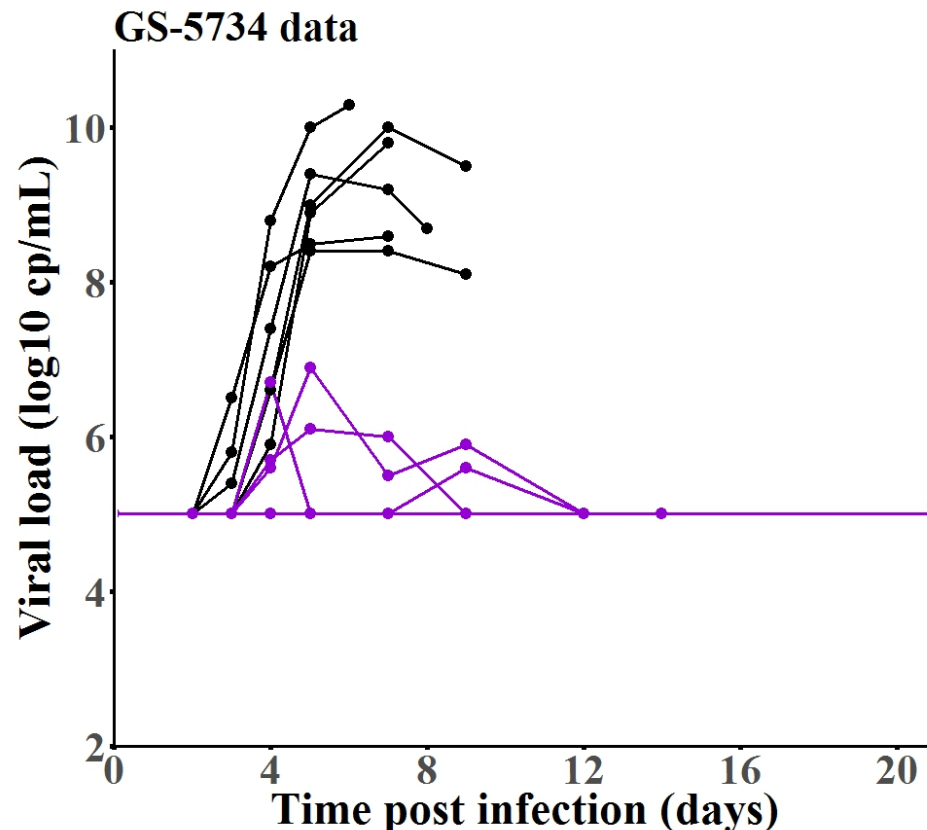
Therapeutic efficacy of the small molecule GS-5734 against Ebola virus in rhesus monkeys

including testes, eyes, and brain. In a rhesus monkey model of EVD, once-daily intravenous administration of 10 mg kg^{-1} GS-5734 for 12 days resulted in profound suppression of EBOV replication and protected 100% of EBOV-infected animals against lethal disease, ameliorating clinical disease signs and pathophysiological markers, even when treatments were initiated three days after virus exposure

We used the model (with variation of R_0 parameter) to fit viral load data only:

→ Found $\varepsilon = 0.9$

→ With this level of effectiveness 100% survival predicted by the model



Validation of model predictions using remdesivir data

LETTER

Warren et al, Nature, 2016

doi:10.1038/nature17180

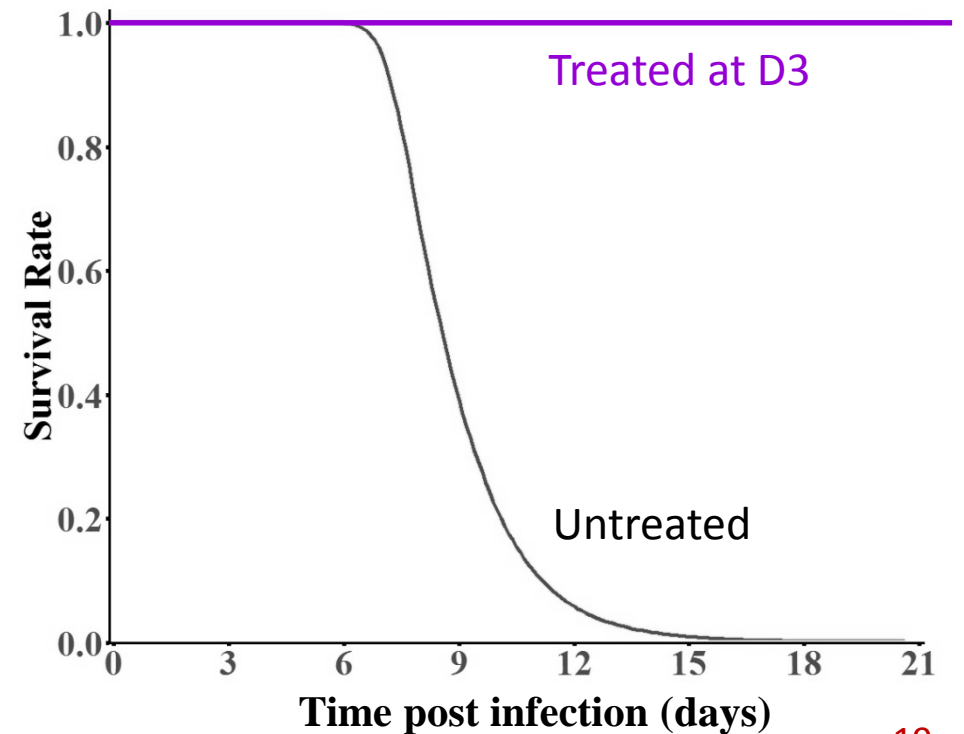
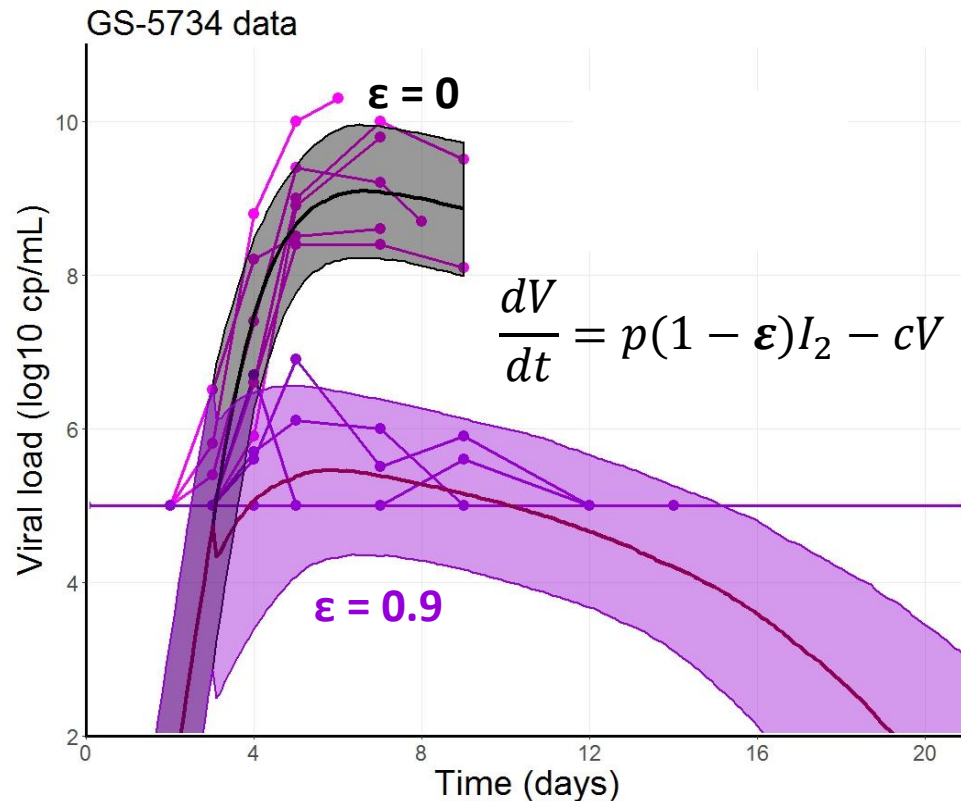
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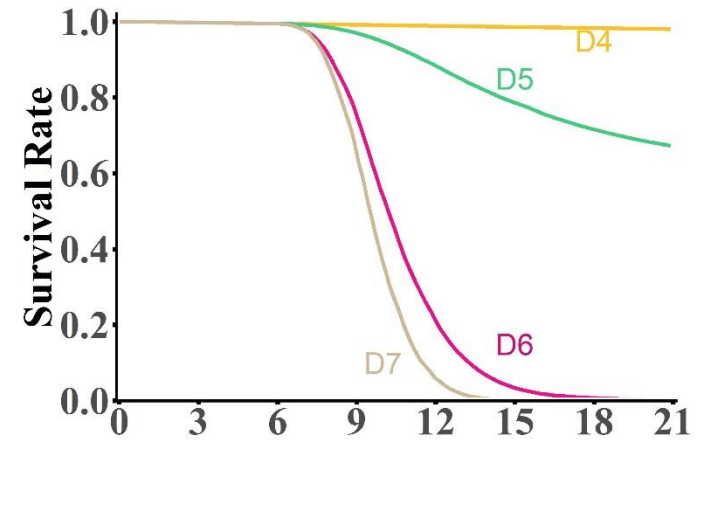
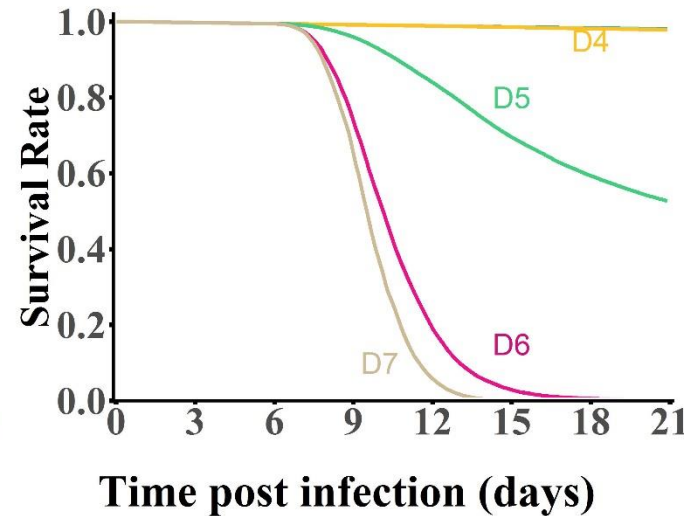
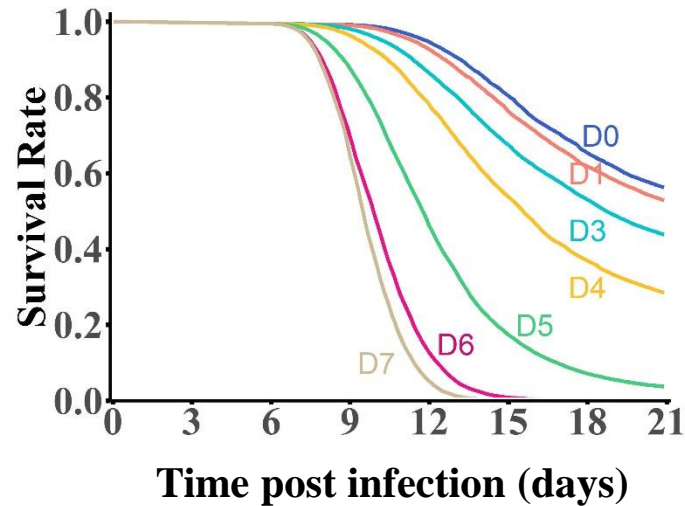
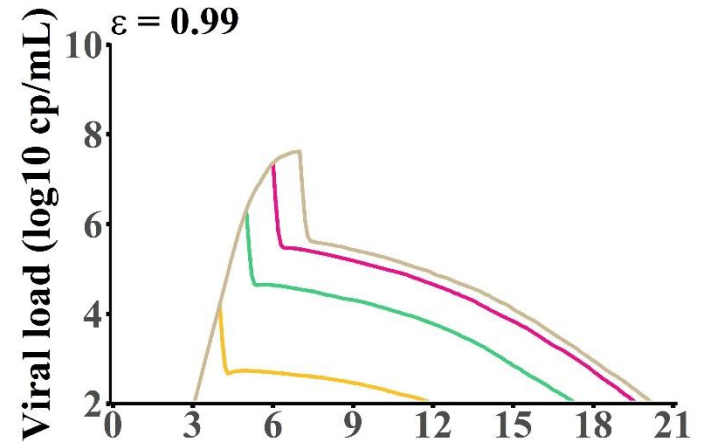
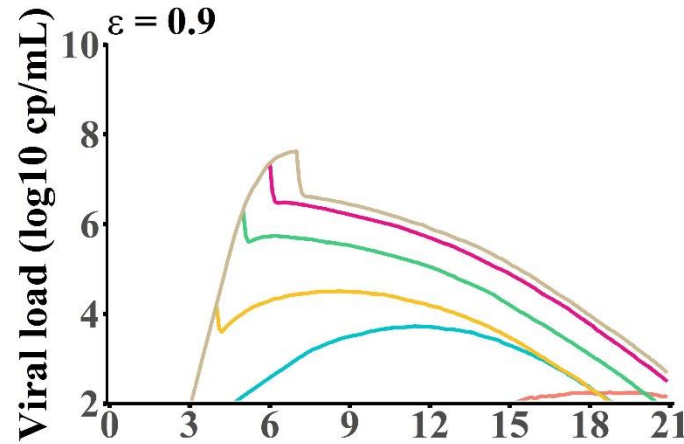
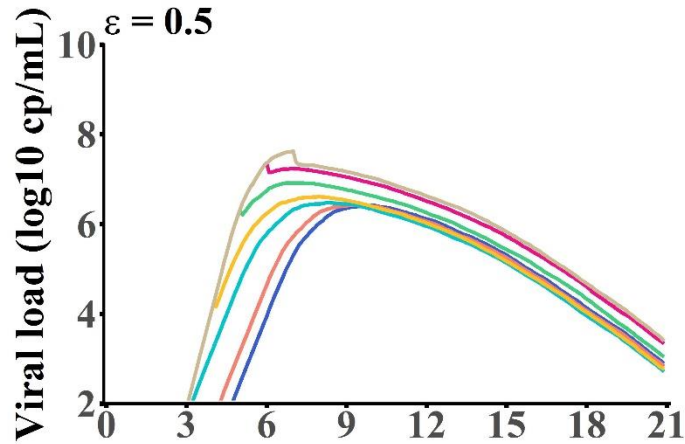
We used the model (with variation of R_0 parameter) to fit viral load data only:

→ Found $\epsilon = 0.9$

→ With this level of effectiveness 100% survival predicted by the model



Impact of the timing of treatment initiation on survival



Treatment initiation: Day 0 Day 1 Day 3 Day 4 Day 5 Day 6 Day 7

Summary on EVD modeling

- Best fit to the data was obtained with models assuming that pro-inflammatory cytokines (IFN α , IL6) were associated with
 - control of viremia via the reduction of target cell population during acute infection
 - time-to-death with a stronger impact than viral load
- Favipiravir initiated two days prior the infection had a moderate impact on viral replication with average $\epsilon = 50\%$ at 180 mg/kg BID
- Model predicts that antiviral drugs may improve survival rate in NHPs only if initiated before the cytokine storm

Patient admission occurred most often close to the viremia peak

RESEARCH ARTICLE

Experimental Treatment with Favipiravir for Ebola Virus Disease (the JIKI Trial): A Historically Controlled, Single-Arm Proof-of-Concept Trial in Guinea

ORIGINAL ARTICLE

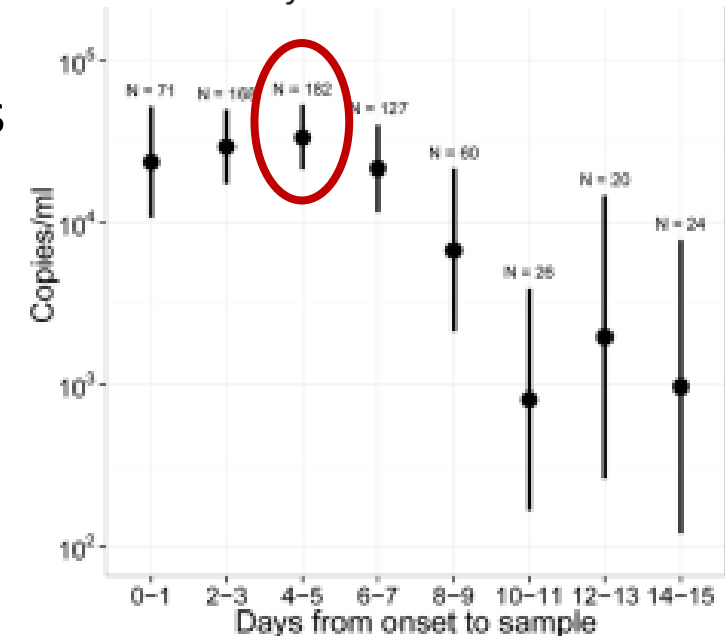
A Randomized, Controlled Trial of ZMapp for Ebola Virus Infection

The PREVAIL II Writing Group, for the Multi-National PREVAIL II Study Team*

RESEARCH ARTICLE

Use of Viremia to Evaluate the Baseline Case Fatality Ratio of Ebola Virus Disease and Inform Treatment Studies: A Retrospective Cohort Study

- Time from symptom onset to inclusion were 4.2 and 3.5 days in the two main clinical trials during the last EVD outbreak
- Retrospective analysis indicated that maximal level of viremia was observed 4 to 5 days after symptoms onset
- These results suggest that majority of patients initiated treatment close to their time to viremia peak
- This supports the design of prophylaxis or post exposure trials for the evaluation of direct antiviral in future outbreaks



Sissoko et al, Plos med, 2016

Prevail study group, NEJM, 2016

Faye et al, Plos med, 2015

Acknowledgements

My PhD supervisor:
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Our collaborators:

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Thi Huyen Tram Nguyen

Hervé Raoul
Frédéric Jacquot
Caroline Carbonelle

Xavier de Lamballerie
Géraldine Piorkowski

Sylvain Baize
Stéphanie Reynard
Alexandra Fizet



And all the Reaction consortium



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Our Sponsors:

