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“Математическое моделирование в биомедицине”

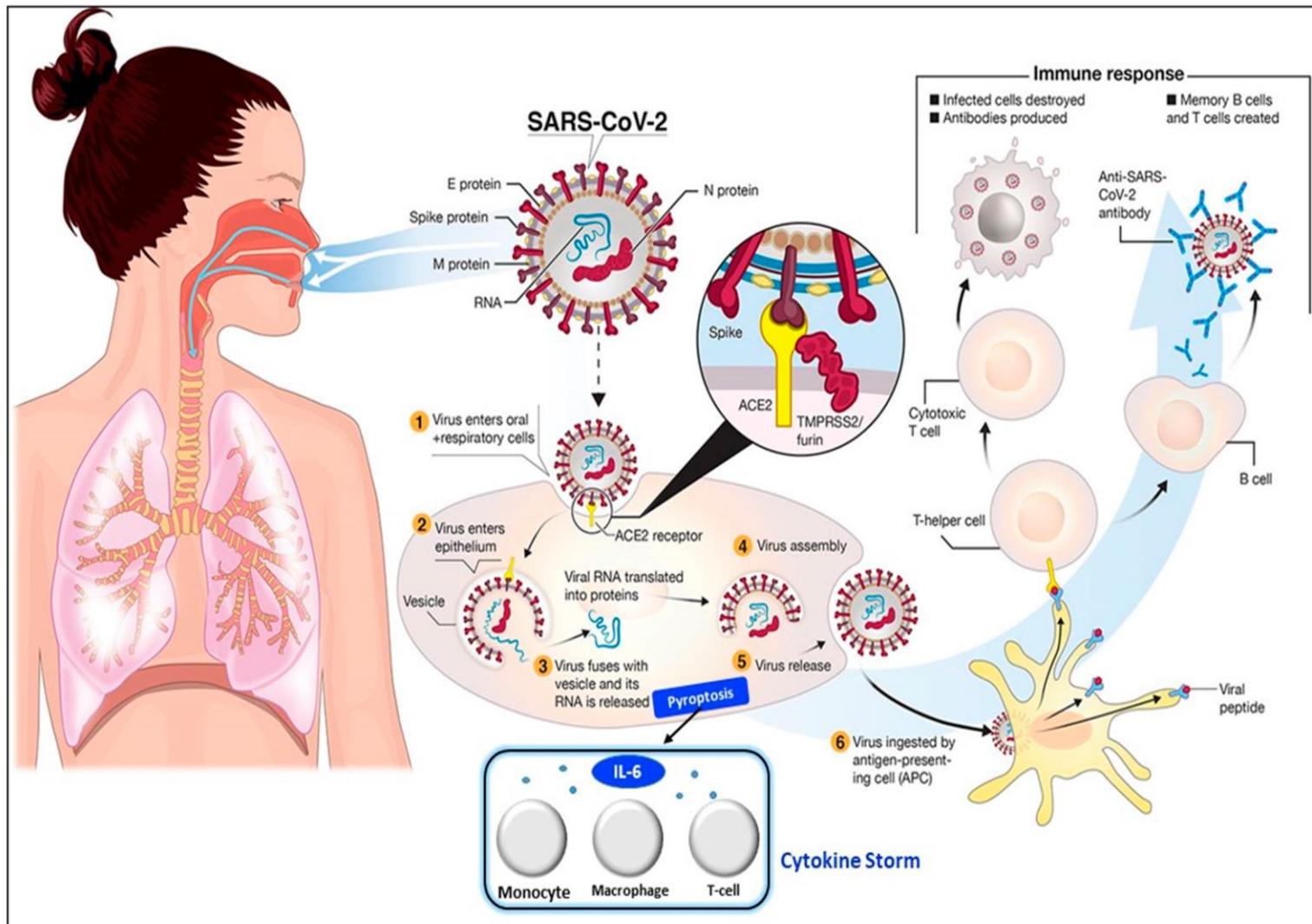
Математическое моделирование распространения вирусной инфекции с учётом воспаления

А. С. Мозохина, L. Ait Mahiout, B. A. Вольперт

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биологии и медицине»,

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Pathophysiology of the SARS-CoV-2 virus

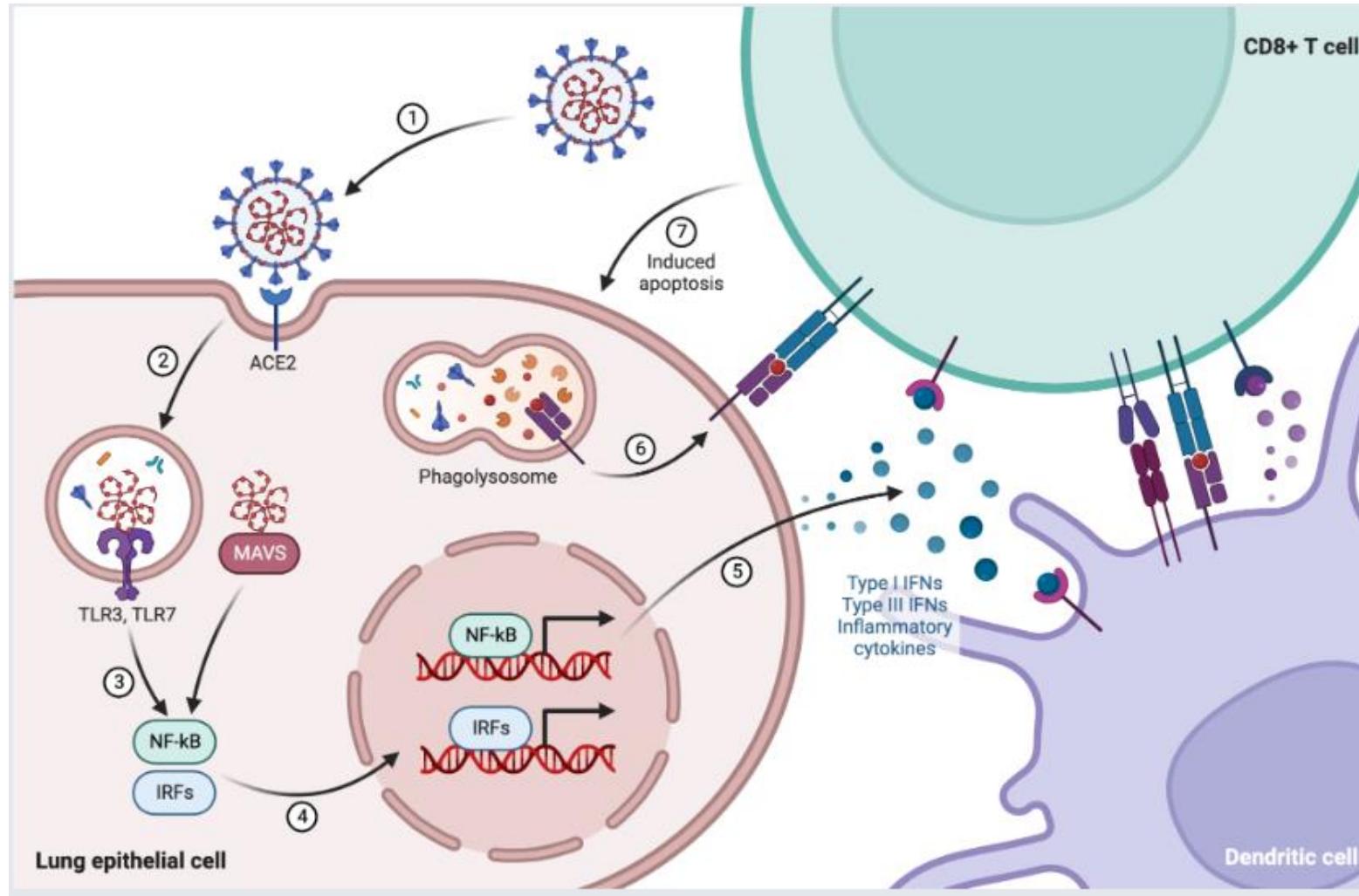


Infection-Mediated Clinical Biomarkers for a COVID-19 Electrical Biosensing Platform

by Reena Sri Selvarajan 1 ID, Subash C. B. Gopinath 2 ID,
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Virus infection - immune response - inflammation



Acute Immune Responses to
Coronaviruses

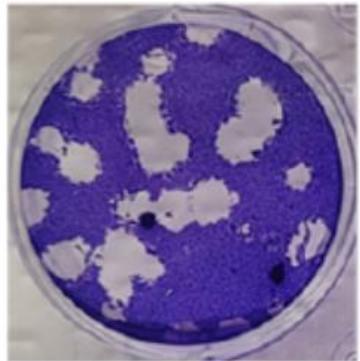


Gillian Dunphy

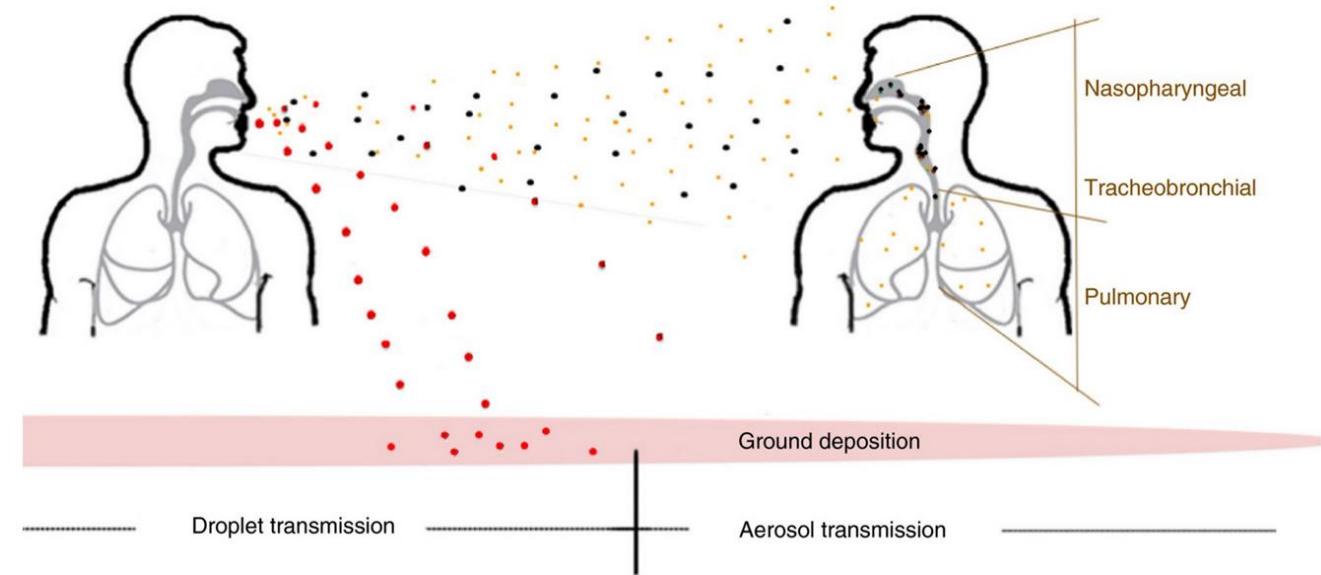
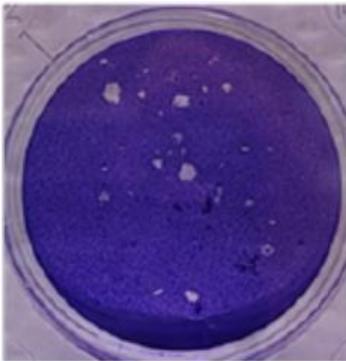
<https://app.biorender.com/b>

Infection spreading virulence and infectivity

Delta



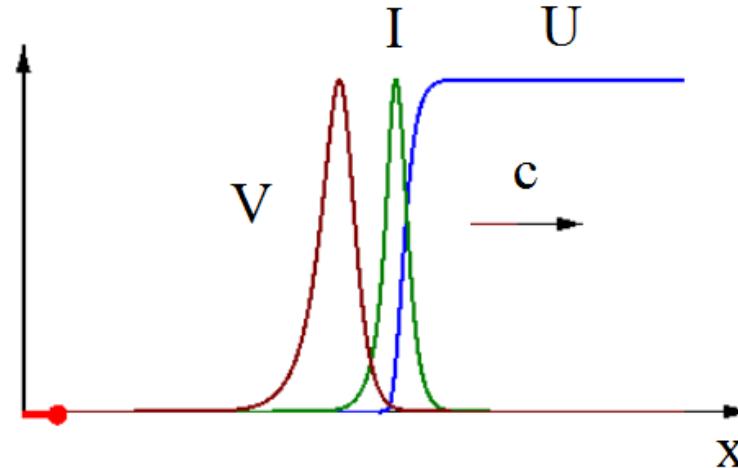
Omicron



$$\frac{\partial U}{\partial t} = -aUV,$$

$$\frac{\partial I}{\partial t} = aUV - \beta I,$$

$$\frac{\partial V}{\partial t} = D \frac{\partial^2 V}{\partial x^2} + bI_\tau - \sigma V$$



System characteristic	Biological interpretation
wave speed c	virulence (disease severity)
total viral load $\int V dx$	infectivity

Model from 3 equations

[Ait+22] L Ait Mahiout et al. “Virus replication and competition in a cell culture: Application to the SARS-CoV-2 variants”. en. In: *Appl. Math. Lett.* 133.108217 (Nov. 2022), p. 108217.

$$\begin{aligned}\frac{\partial U}{\partial t} &= -aUV, \\ \frac{\partial I}{\partial t} &= aUV - \beta I, \\ \frac{\partial V}{\partial t} &= D\frac{\partial^2 V}{\partial x^2} + bI_\tau - \sigma V\end{aligned}$$

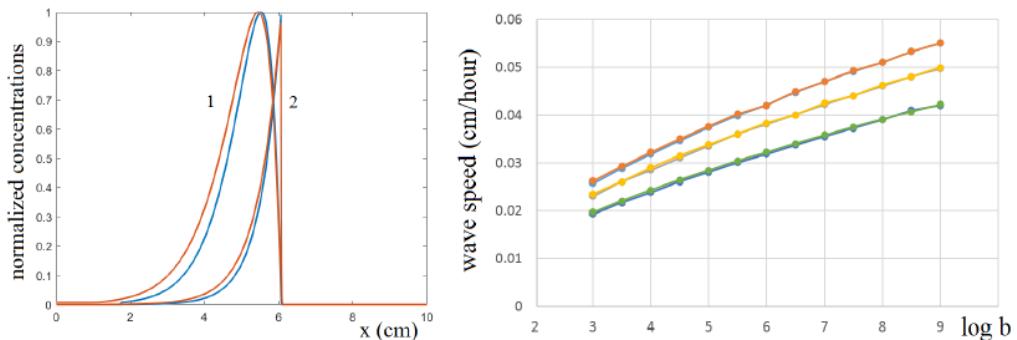


Figure 1: Left: virus concentration (curve 1) and concentration of infected cells (curve 2) in numerical simulations (red) and analytical approximation (blue) for the values of parameters: $a = 0.1$ (1/(hour · virus)), $b = 1000$ (copies/(hour · cell)), $\beta = 0.1$ (1/hour), $\sigma = 0.1$ (1/hour), $D = 10^{-4}$ (cm^2/hour). Analytical and numerical solutions for $v(\xi)$ are normalized by its maximum, $v_{\max} = 3672$ (copies/ml). Right: wave speed in numerical simulations and analytical formula (curves coincide) for the values of parameters: $a = 0.1$, $D = 0.001$, $\beta = 0$, $\tau = 5$ (hour) (upper curve), $\tau = 5$ (middle curve) and $\tau = 8$ (lower curve).

$$\begin{aligned}R_v &= abu_0/(\beta\sigma) \\ R_v(\omega - 1) &= \ln \omega \quad \omega = u_f/u_0. \\ V_x &\approx bcu_0/(\beta\sigma). \quad V_X \equiv \int_{-\infty}^{\infty} v(x)dx = -\frac{c}{a} \ln \omega. \\ c &= \min_{\mu > \mu_0} F(\mu) \equiv \frac{\sqrt{D}\mu}{\sqrt{\mu + \sigma - abu_0 e^{-\mu\tau}/(\mu + \beta)}} ,\end{aligned}$$

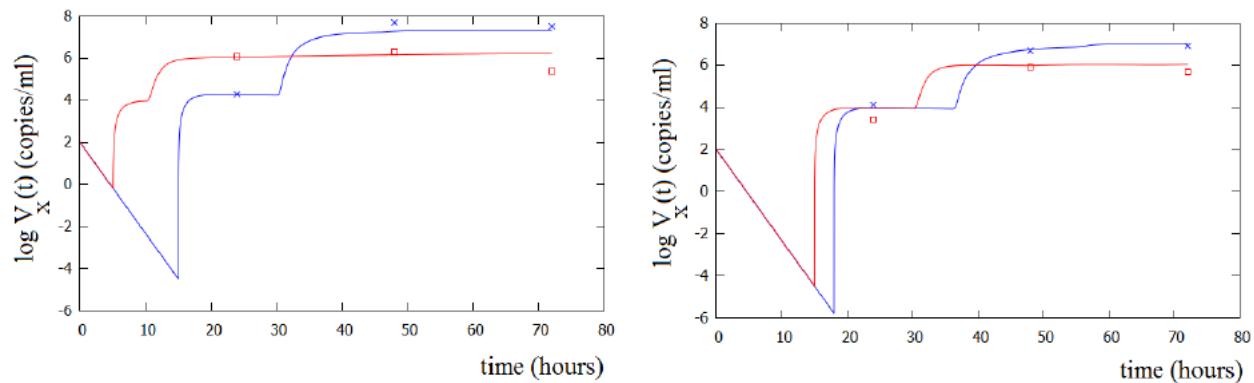


Figure 3: Left: human nasal cells, experimental results from [5] (dots) and numerical simulations with the values of parameters for Delta (blue): $a = 10^{-5}$, $b = 2 \cdot 10^7$, $\beta = 0.01$, $\sigma = 1$, $\tau = 15$; for Omicron (red): $a = 10^{-4}$, $b = 10^6$, $\beta = 0.01$, $\sigma = 1$, $\tau = 8$. Right: human lung cells, experimental results from [5] (dots) and numerical simulations with the values of parameters for Delta (blue): $a = 10^{-5}$, $b = 10^7$, $\beta = 0.01$, $\sigma = 1$, $\tau = 17$; for Omicron (red): $a = 10^{-4}$, $b = 10^6$, $\beta = 0.01$, $\sigma = 2$, $\tau = 15$. Common parameters: $D = 0.001$, $L = 10$, $x_0 = 1$, $v_0 = 1$, $U_0 = 1$. Units of parameters are given in Figure 1.

Model of the viral infection with respect to inflammation

$$\frac{\partial U}{\partial t} = k_1(U_0 - U) - k_2UV,$$

$$\frac{\partial W}{\partial t} = k_2UV - k_3SW - \sigma_1W,$$

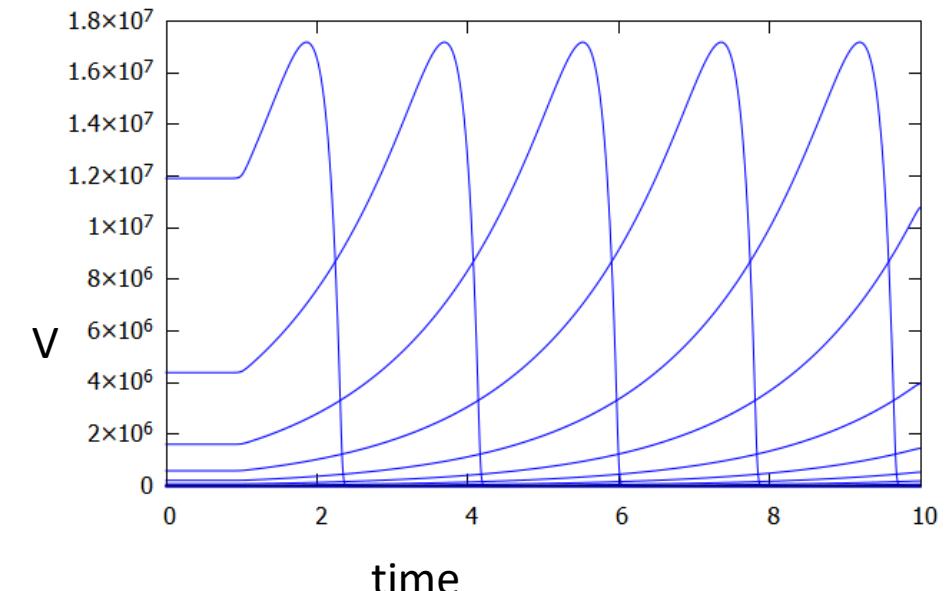
$$\frac{\partial V}{\partial t} = D_1 \frac{\partial^2 V}{\partial x^2} + bW - \sigma_3V,$$

$$\frac{dS(t)}{dt} = k_6J(W) - k_7J(W)S - \sigma_4S,$$

$$J(W) = \int_{-\infty}^{+\infty} W(x, t) dx.$$

- U - uninfected cells
- W - infected cells
- V - virus
- S - inflammatory cytokines

Citation: Mozokhina, A.;
Ait Mahiout, L.; Volpert, V. Modeling
of Viral Infection with Inflammation.
Mathematics 2023, 11, 4095.
[https://doi.org/10.3390/
math11194095](https://doi.org/10.3390/math11194095)



Case 1: clearance of cytokines = 0

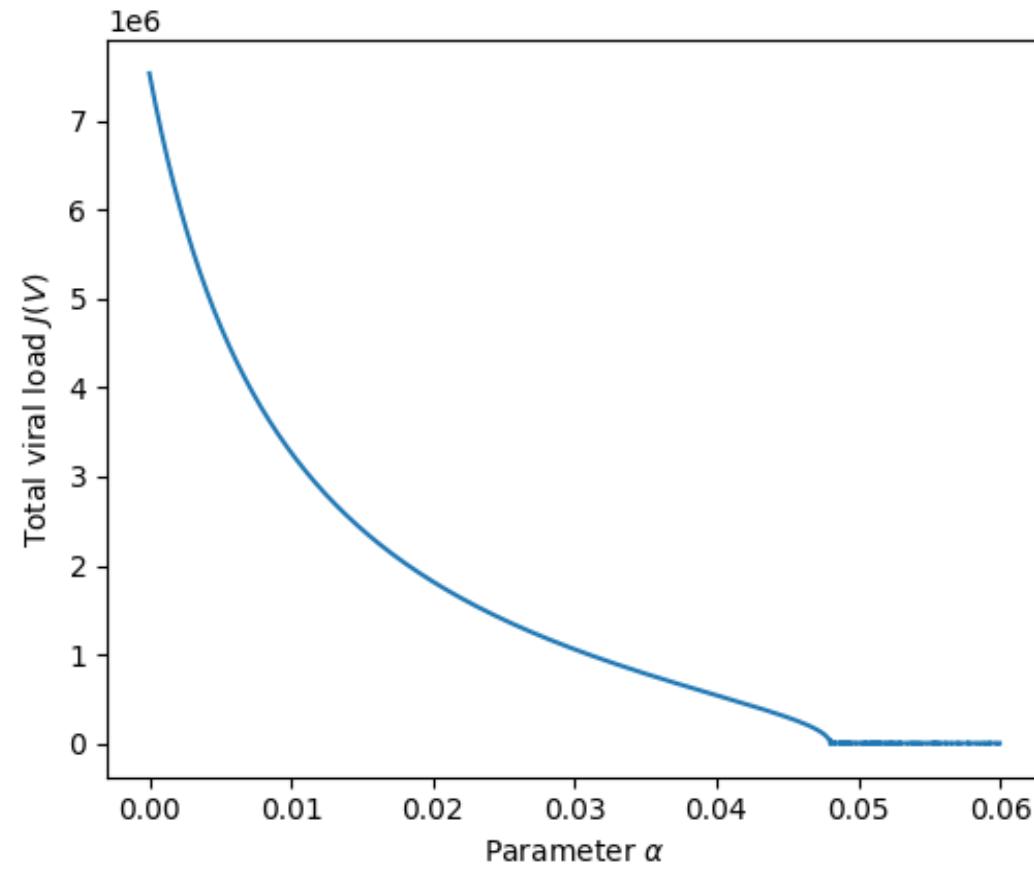
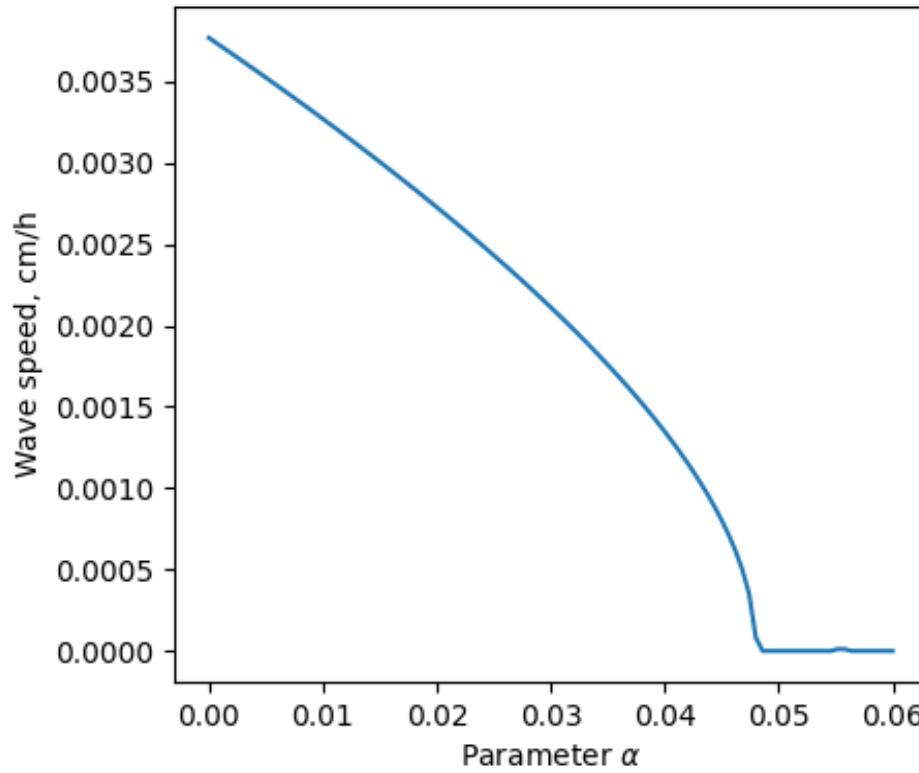
- Swell / edema
- Stationary points: $W = 0, V = 0, S$ and U are arbitrary.
- Virus replication number: $R_v^{(2)} = \frac{k_2 b U_0}{\sigma_3 (k_3 S_0 + \sigma_1)} < 1.$
- Wave
 - total viral load: $\ln X = R_v(X - 1), \quad R_v = \frac{k_2 b u_0}{\sigma_3 (\alpha + \sigma_1)}, \quad \alpha = k_3 k_6 / k_7. \quad X = u_f / u_0$
 $J(v) = -\frac{c \ln X}{k_2} \approx \frac{c R_v}{k_2}.$
 - for $\alpha \rightarrow 0$ all formulas are the same as for UIV system
 - for $\alpha > \frac{k_2 b u_0}{\sigma_3} - \sigma_1$ there is no solution of the equation for X
- wave speed: $c^2 = \min_{\mu > \mu_0} \frac{D_1(\mu + \sigma_1 + \alpha)\mu^2}{(\mu + \sigma_3)(\mu + \sigma_1 + \alpha) - b k_2 u_0}.$

Dependence of the total viral load and the wave speed on the effective cell death due to inflammation

for $\alpha = 0$ – the same as in UIV without inflammation

$vel=0.0037683636560039402$

$jv=7513497.885188183$



inflammation reduces both infectivity and severity of the disease

The values of the wave speed and the total viral load are bounded from above by the values obtained in UIV system

Case 2: clearance of cytokines $\neq 0$

- Stationary points: $W = 0, V = 0, S = 0$ and any U .
- Virus replication number: $R_v^{(1)} = \frac{k_2 b U_0}{\sigma_1 \sigma_3} < 1$. - the same as for UIV system
- Wave
 - Total viral load: $\ln X(\ln X - q) = R_v(X - 1)(\ln X - r)$, $R_v = \frac{k_2 b u_0}{\sigma_3 (\alpha + \sigma_1)}$,
 $q = \frac{bk_2\sigma_1\beta}{\sigma_3 c(\alpha + \sigma_1)}$, $r = \frac{bk_2\beta}{\sigma_3 c}$, $\beta = \sigma_4/k_7$.

Proposition 1. Equation (2.17) has a solution $X \in (0, 1)$ if and only if $R_v^{(1)} > 1$.

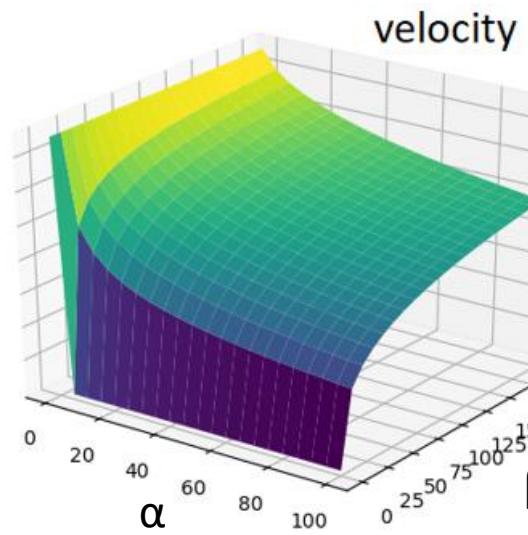
- wave speed: $c^2 = \min_{\mu > \mu_0} \frac{D_1[(\mu + \sigma_1)(J(w) + \beta) + \alpha J(w)]\mu^2}{(\mu + \sigma_3)[(\mu + \sigma_1)(J(w) + \beta) + \alpha J(w)] - bk_2 u_0 (J(w) + \beta)}$,

For $\alpha \rightarrow 0$ ($\beta \rightarrow 0$ or β is fixed): the UIV case

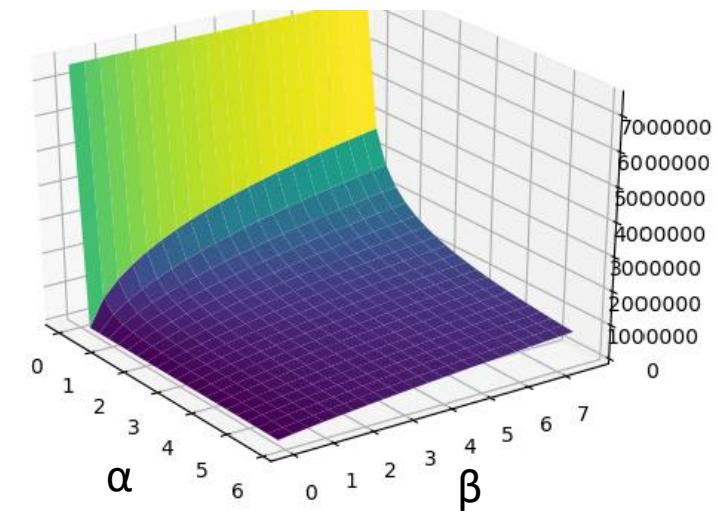
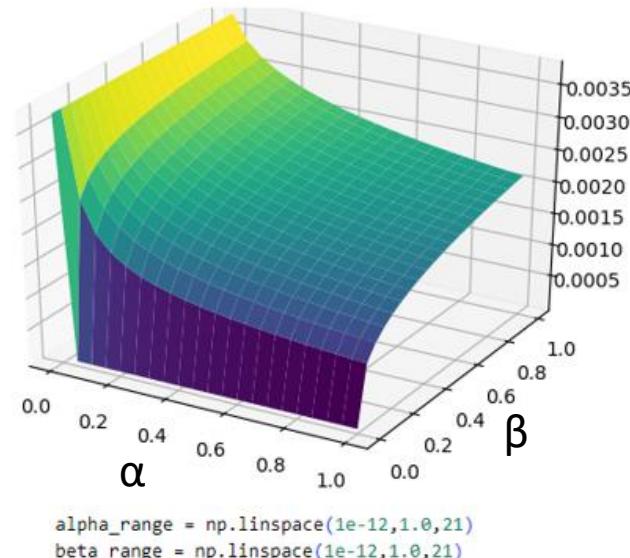
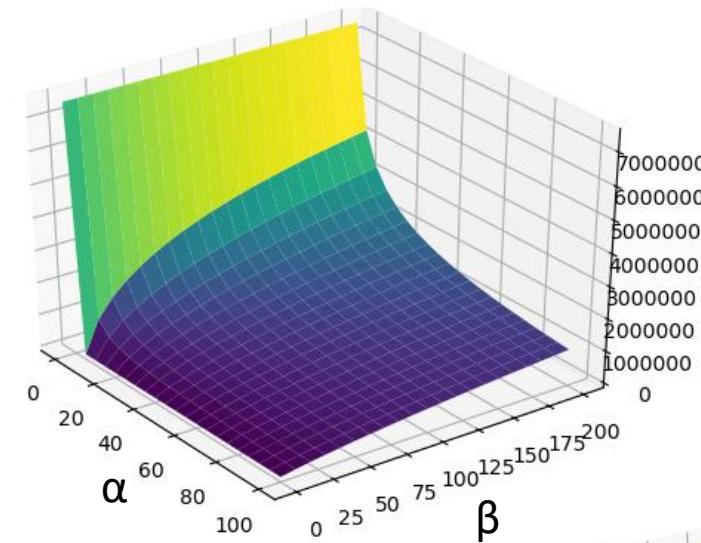
For $\beta \rightarrow 0$ (α is fixed, $\neq 0$): the previous ($\sigma_4=0$) case - all the dynamics depends only on α

For $\beta \rightarrow \infty$ (α is fixed): the UIV case where the R_v inversely depends on alpha

Dependence of the total viral load and the wave speed on the inflammation parameters

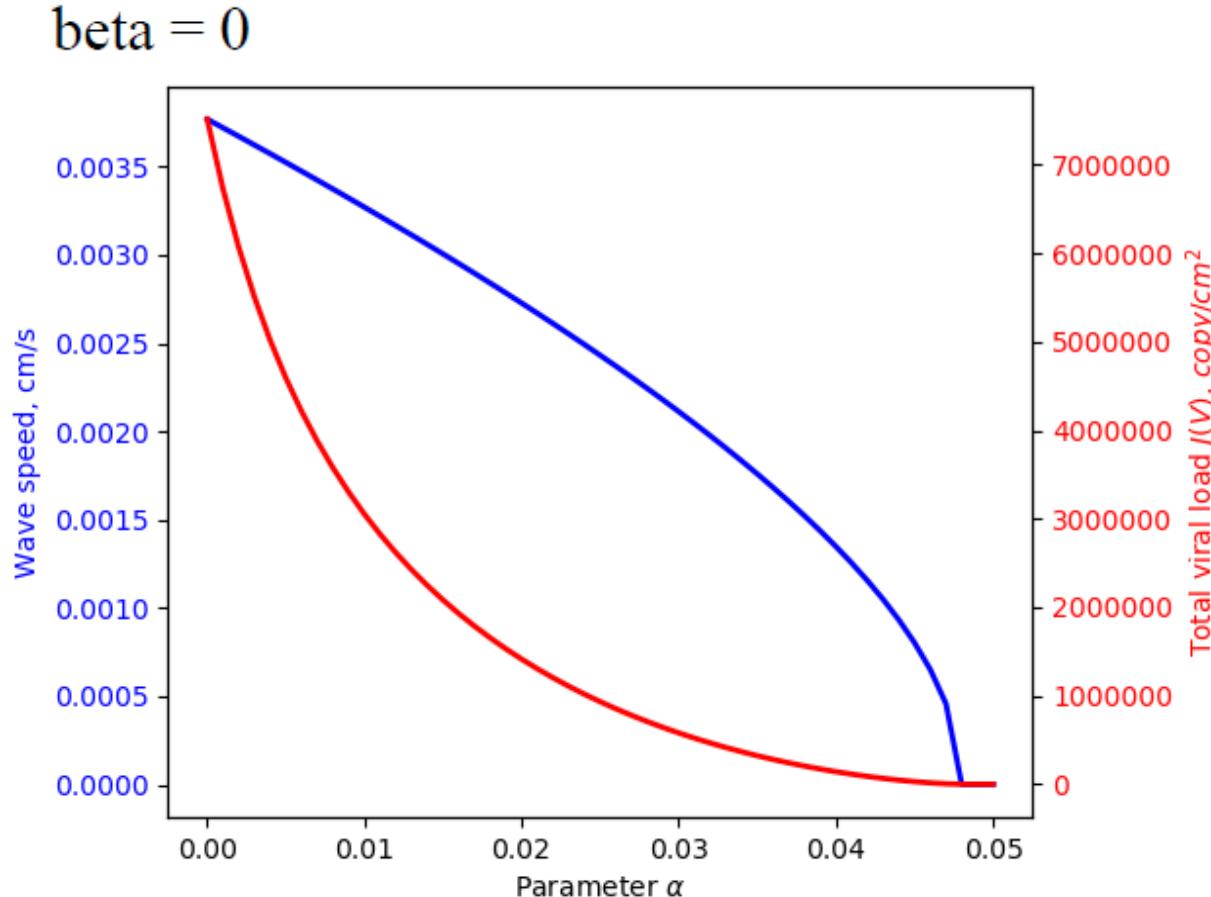


Total viral load



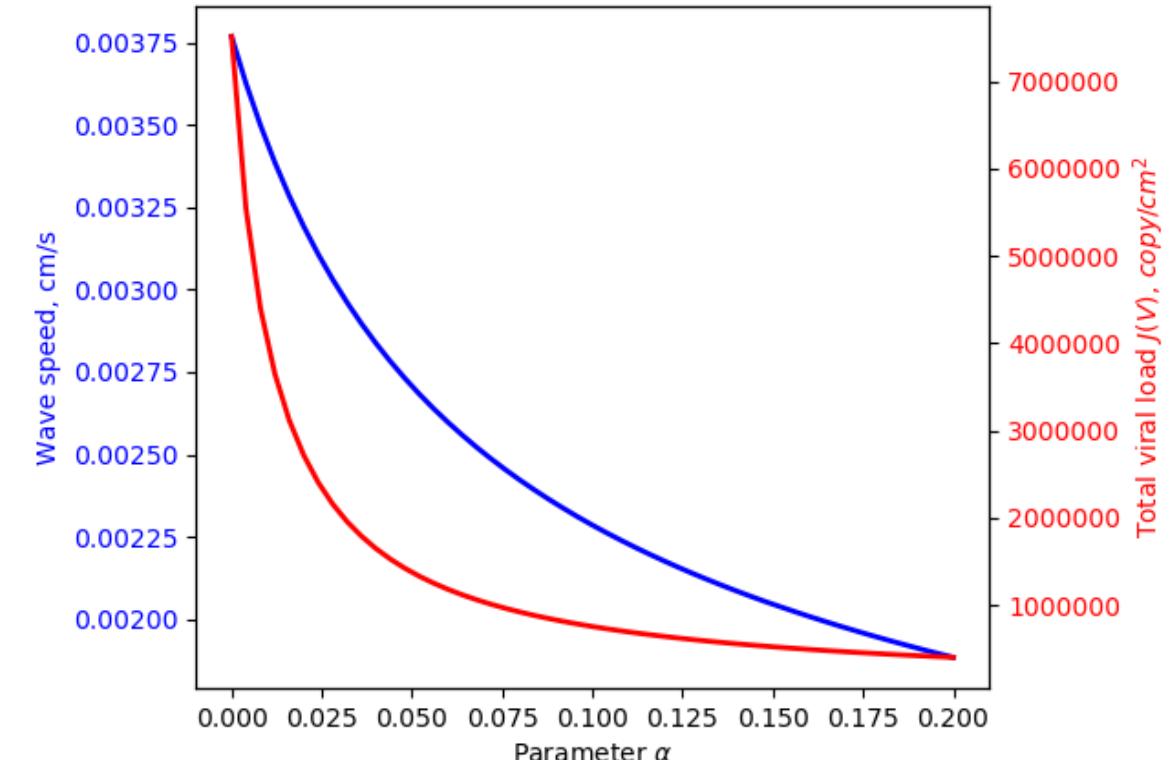
Dependence of the wave speed and total viral load on the effective cell death induced by inflammation (α)

For $\alpha \rightarrow 0$ ($\beta \rightarrow 0$ or β is fixed): the UIV case:



vel=0.0037683636560039402
jv=7513497.885188183

beta = 0.1



effective cell death decreases both the severity and infectivity of the disease

The values of the wave speed and the total viral load are bounded from above by the values obtained in UIV system

Dependence of the wave speed and total viral load on the effective cytokine clearance (β)

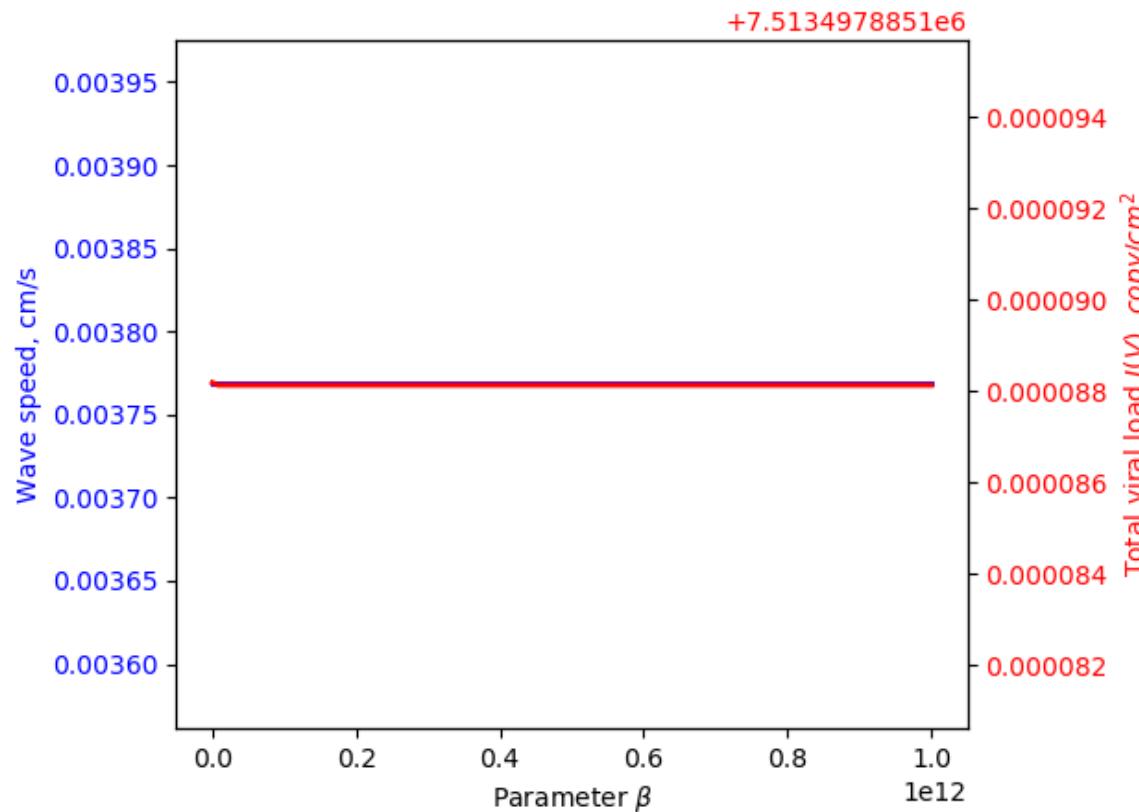
For $\beta \rightarrow 0$ (α is fixed): the previous ($\sigma_4=0$) case - all the dynamics depends only on α

For $\beta \rightarrow \infty$ (α is fixed): the UIV case where the R_v inversely depends on alpha

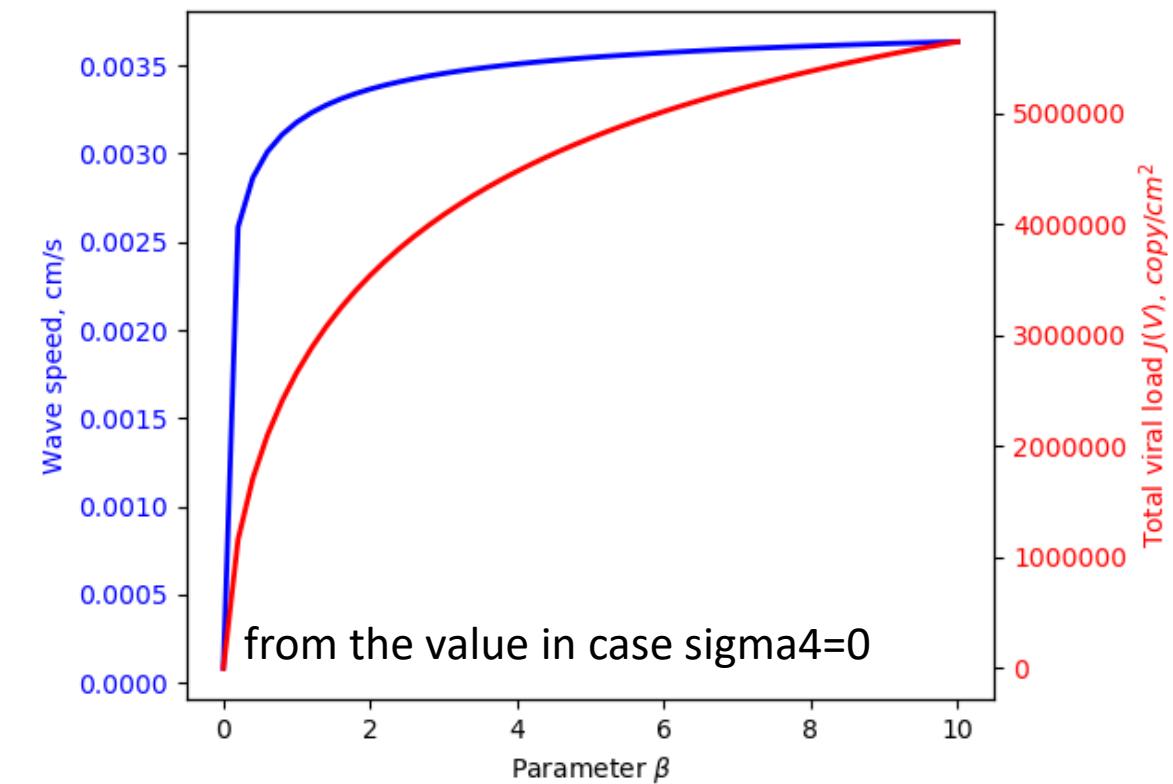
$v_{el}=0.0037683636560039402$

$j_v=7513497.885188183$

$\overline{\text{alpha}} = 0$



$\overline{\text{alpha}} = 0.1$



Effective cytokine clearance increases both the wave speed and total viral load from the value of the no-clearance case but not higher than the value of UIV case

Conclusions

- The inflammation decreases both the total viral load and the wave speed
- In the swell (edema) presence, the total viral load and the wave speed are reduced by inflammation (effective cell death induced by inflammation)
- The increased clearance results in the increasing of the total viral load and the wave speed up to values which they have in the case without inflammation (in the infinity)

The results show that the swell can be a positive mechanism for revealing the virus infection. The last conclusion requires further investigations

Thank you for attention!