SINGLE SCALE AND MULTI-SCALE MODELS OF VIRAL INFECTIONS AND ANTI-VIRAL IMMUNE RESPONSES: APPLICATIONS TO INFECTIOUS AND NON-INFECTIOUS DISEASES

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Workshop: "NEW TRENDS IN BIOMATHEMATICS: Applications in Oncology and Immunology"



Non-infectious diseases:

oncogenic viruses: we want to control & eliminate them... **C Infectious diseases:** we

want to control the infections & eliminate viruses causing them...



Oncolytic viruses



Non-infectious

diseases: viruses can be used as treatments for these diseases (control & increase replication)

(e.g., oncolytic viruses in cancer treatment)



http://oncolyticvirus.files.wordpress.com/2010/12/oncolytic-virus-in-action

VIRUS SPREAD ACROSS MULTIPLE SPATIAL & TEMPORAL SCALES:

1. Environmental transmission of viruses: nosocomial infections: hospital wards/ bays/operating rooms/facilities/...
=> infected patient



3. Cell-to-cell transmission: anti-

viral immune responses that control the infection...



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Here: focus on infectious diseases

Here: focus on nominfectious diseases (cancer)

Environmental & human-to-human transmission of viruses: nosocomial infections

• Nosocomial infections in the context of COVID-19 (collaboration with Dr. B. Parcell, Ninewells Hospital, UK)





https://link.springer.com/article/10.1007/s00449-022-02733-9

Environmental & human-to-human transmission of viruses: nosocomial infections



 Nosocomial infections in the context of COVID-19 (collaboration with Dr. B. Parcell, Ninewells Hospital, UK)
 Ward: four 4-bed bays + six single-bed rooms
 Ward: four 6-bed bay

Question (asked in summer 2020):

How to distribute the hospitalized patients in wards with

- 4-bed bays + single-bed rooms vs.
- 6-bed bays + single-bed rooms to reduce SARS-CoV-2 spread across the hospital?
 - Infected patients moved to single rooms (if available)
 - Infected rooms were closed for cleaning... no new patients admitted to hospital

D. Moreno Martos, B. Parcell, RE (2020). Modelling the transmission of infectious diseases inside hospital bays: implications for Covid-19 . Math. Biosci. Eng., 17(6) , 8084-8104



Environmental & human-to-human transmission of viruses: nosocomial infections

 Nosocomial infections in the context of COVID-19 (collaboration with Dr. B. Parcell, Ninewells Hospital, UK)

Agent-based network model: each node is a patient (in a bed) with specific characteristics: $I_{n,t} = [C_{n,1,t}, C_{n,2,t}, ..., C_{n,m,t}]$

- 1. Epidemiological status (susceptible, exposed, infected, recovered)
- 2. Bed # in which the patient is placed ...
- 3. Start of the incubation period
- 4. Duration of incubation period
- 5. Time since individual has become infectious
 - -> viral transmission possible
- 1. Recovery time



Model can account for stochastic fluctuations between interconnected individuals inside the bay





B. Parcell

(a) 4–bed bays: Average number of individuals per time Susceptible 0 Exposed 5 40 60 20 20 0 time (days) Recovered Infected 5 60 20 40 20 time (days) Bed 2 Bed 1 1.00 1.00



(c) 6–bed bays:



B. Parcell



time (days)

time (days)

40

40

t (days)

t=0

t=10

t=20

t=30

t=40

t=50

t=60

60

20

20

R

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Hypotheses tested & confirmed in silico: Having only 4-bed bays is better to reduce the spread of the infection

D. Moreno Maros, B. Parcell, RE (2020). Modelling the transmission of infectious diseases inside hospital bays: implications for Covid-19. Math. Biosci. Eng., 17(6), 8084-8104







Hypotheses tested in silico: the role of asymptomatic individuals

- More often RT-PCR tests (every day vs. every 3 days) are better at detecting exposed/infected patients (in particular for 6-bed bays)
- Impact of mask wearing by medical staff... and infected staff infecting patients ...

https://link.springer.com/article/10.1007/s00449-022-02733-9

D. Moreno Maros, B. Parcell, RE (2020). Modelling the transmission of infectious diseases inside hospital bays: implications for Covid-19 . Math. Biosci. Eng., 17(6) , 8084-8104

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Models, even if not perfectly calibrated (scarce data), can be used to make administrative (& public policy) decisions... especially in exceptional times...

• Open the hospital for elective surgery procedures..



Average number of individuals per time

100



D. Moreno Maros, S. Folley, B. Parcell, D. Trucu, RE (**2022**). A computational investigation of COVID-19 transmission inside hospital wards and associated costs. Math. Biosci. Eng., 17(6), 8084-8104

Changes in the incubation period for human-to-human transmission...





B. Parcell





Changes in the incubation period for human-to-human transmission...



B. Parcell



Bed 1 Bed 2 2 Bed 4 Bed 4 bed 5 Bed 4 bed 5 Bed 4

Buying Time—The Immune System Determinants of the Incubation Period to Respiratory Viruses

Tamar Hermesh^{1,†}, Bruno Moltedo^{1,†}, Carolina B. López^{1,2} and Thomas M. Moran^{1,*}

Infection Onset of Activation of i		Symptoms innate immunity	Effector T-Cells in	nfiltrate the lungs	Virus cleared	
Dag	y0 Da	Day 2		y 6	Day 10	
	INCUBATION PERIOD	INFLAMMATORY	PHASE	RESOLUTION OF INFECT	ION	
	STEALTH PHASE	INITIATION OF INNATE AND ADA	APTIVE IMMUNITY	EFFECTOR T-CELLS CLEAR VIRUS-INFECT	TED CELLS	
	Virus	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	** *			
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Also viral transmission probably depends on the immune system (& anti-viral immune response) of the infected individuals ...



What is the in-host dynamics of SARS-CoV-2

virus? A challenge within a multiscale vision of living systems

N. Bellomo * R. Eftimie [†] G. Forni [‡]





What is the in-host dynamics of SARS-CoV-2

virus?

systems

A challenge within a multiscale vision of living







Qualitative investigation (& exploration) of model dynamics





What is the in-host dynamics of SARS-CoV-2 virus? A challenge within a multiscale vision of living systems

N. Bellomo * R. Eftimie ^{\dagger} G. Forni ^{\ddagger}



Qualitative investigation of model dynamics



Days (from symptoms onset)

What is the in-host dynamics of SARS-CoV-2 virus? A challenge within a multiscale vision of living systems

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T cell numbers (ELISPOT) vs. Days









Timeline of mathematical models

Virus enters the body & infection starts

Timeline of (many) clinical datasets



There are datasets to validate the models, but maybe not in the form modellers want them => adapt the models Viruses that cause infectious diseases...

- SARS-CoV-2: mild to severe infections
- Adenoviruses: mild respiratory infections (common cold)
 - ✓ infect mammalian species & birds
- Vesicular stomatitis virus: vesicles develop on the tong, excess salivation,...
 - ✓ Infects cattle, pigs, horses
- ...many other viruses...

Viruses used to treat cancers:

• Vaccine viruses: trigger tumourspecific immunity that eradicate tumours & maintain immunological memory



• Oncolytic viruses: genetically modified to selectively infect, replicate in and kill tumour cells

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SARS-CoV-2 replicates and displays oncolytic properties in clear cell and papillary renal cell carcinoma

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Oi Kuan Choong<sup>1,2</sup>, Rasmus Jakobsson <sup>1,3</sup>, Anna Grenabo Bergdahl<sup>4,5</sup>, Sofia Brunet<sup>2,6</sup>, Ambjörn Kärmander <sup>2,6</sup>, Jesper Waldenström<sup>2,6</sup>, Yvonne Arvidsson<sup>1,7</sup>, Gülay Altiparmak<sup>1,7</sup>, Jonas A. Nilsson<sup>2,8,9</sup>, Joakim Karlsson <sup>2,8,9</sup>, Kristina Nyström<sup>2,6</sup>*, Martin E. Johansson <sup>1,2,7</sup>*
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Repurposing the oncolytic virus VSV Δ 51M as a COVID-19 vaccine

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Almohanad A. Alkayyal^{1,2*}, Manar Darwish², Reham Ajina^{2,3}, S. sh Y. Alabbas², Mohammed A. Alotaibi², Abeer Alsofyani^{4,5}, Maha L. Khamseen², Maumonah Hakami², Omar A. Albaradie^{2,6}, Abdulaziz M. ¹-aqlan^{2,6}, Sharif Hala^{5,7}, Abdullah Faisal Alsahafi^{5,7}, Samer Zakri^{5,7}, Adnan Almuzaini⁸, Khamis Alsharari⁸, Feras Kaboha⁸, Mustafa Y. Taher⁹, Haggag S. Zein^{21,7}, Fayhan Alroqi^{2,11,12} and Ahmad Bakur Mahmoud^{9,13,14*}

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Oncolytic viruses: genetically modified to selectively infect, replicate in and kill tumour cells

Basic reproductive number (epidemiological & in-host levels)

- As small as possible for controlling the spread of an infectious disease (spread of a virus) through a
 population (of humans, of cells, ...)
- As large as possible to ensure the spread of an oncolytic virus through a cell population
 - Not only how many viral particles are released from 1 infected cell, but also how many of these particles infect other cells due to various physical and immune barriers



https://x.com/roinnslainte

ONCOLYTIC VIRAL THERAPIES: THE FINE BALANCE BETWEEN THE ANTI-VIRAL IMMUNE RESPONSES AND ANTI-TUMOUR IMMUNE RESPONSES



B.W. Bridle et al., Molecular Therapy, 2010, 1430–1439.







J. Bramson

B. Brydle

D.J.D. Earn

Double immunization (prime-boost) experimental protocol for murine melanoma (at McMaster University, Canada)

- Vaccine Adenovirus (Ad) expressing a human tumour antigen, to prime the antitumour immune response;
- Injection of oncolytic Vesicular Stomatitis Virus (VSV) carrying the same tumour antigen => virus replication & tumour elimination

But tumour relapses (median survival 54 days)

ONCOLYTIC VIRAL THERAPIES: THE FINE BALANCE **RESPONSES** BETWEEN ANTI-VIRAL IMMUNE ТНЕ AND **RESPONSES** ANTI-TUMOUR IMMUNE



Delaying the 2nd injection with VSV(in tumour-free mice) leads nbr. of IFN-CD8 T cells



Conceptual description of immune response (for tumour-bearing mice)



Oncolytic vesicular stomatitis virus quantitatively and qualitatively improves primary CD8⁺ T-cell responses to anticancer vaccines

> Byram W Bridle¹, Derek Clouthier², Liang Zhang², Jonathan Pol², Lan Chen², Brian D Lichty², Jonathan L Bramson², and Yonghong Wan^{2,*}

Hypothesis proposed: increasing the delay between the injection of the two viruses should allow for a higher secondary immune response (with better anti-tumour effects) => cannot test it on mice for ethical reasons...

& experiments (to increase the device of the day at a time) are expensive of the day at a time) are expensive of the day at a time of t Immuno rochonco

ONCOLYTIC VIRAL THERAPIES: THE FINE BALANCE **ONSES** RESP ΝΕ Ε Ε AND B **ONSES** RESP R MU ΝΕ I M





Conceptual description of immune response (for tumour-bearing mice)



sing the delay between the injection of the igher secondary immune response (with





ONCOLYSIS + SECRETION OF PROTEASES

viruses spread:

...

FACILITATE DEEP TUMOR PENETRATION



Multiscale modelling of cancer response to oncolytic viral therapy

 Talal Alzahrani, Raluca Eftimie, Dumitru Trucu*

 Division of Mathematics, University of Dundee, Dundee DD1 4HN United Kingdom



Viral spread into a 2D domain...with ECM degradation by tumour cells $K_2(\mu_s, \cdot, \cdot)$ at it at ive exploration of outcomes

 $\begin{array}{l} \text{Macroscal} \quad \text{Macroscal}$

A 2nd layer of complexity: Microscale dynamics

$$\frac{\partial a}{\partial \tau} = \underbrace{D_a \Delta h_1}_{\text{diffusion}} - \underbrace{\psi_{11} ap}_{uPA/PAI-1} + \left(\underbrace{\psi_{12}}_{\text{production}} - \underbrace{\psi_{13} u}_{uPA/uPAR}\right) f_{uPA}^{\epsilon Y}(y, \tau)$$

$$\frac{\partial p}{\partial \tau} = \underbrace{D_p \Delta p}_{\text{diffusion}} - \underbrace{\psi_{21} ap}_{uPA/PAI-1} - \underbrace{\psi_{22} p f_{PAI-1}^{\epsilon Y}(y, \tau)}_{PAI-1/ECM} + \underbrace{\psi_{23} m}_{\text{production}}.$$

$$\frac{\partial m}{\partial \tau} = \underbrace{D_m \Delta m}_{\text{diffusion}} + \underbrace{\psi_{31} a f_{uPA}^{\epsilon Y}(y, \tau)}_{uPA/uPAR} + \underbrace{\psi_{32} p f_{PAI-1}^{\epsilon Y}(y, \tau)}_{PAI-1/ECM} - \underbrace{\psi_{33} m}_{\text{decay}}.$$

Viral spread into a 3D domain...with ECM degradation by tumour cells ... but NO virus yet...

Mathematical Modelling of Glioblastomas Invasion within the Brain: A 3D Multi-Scale Moving-Boundary Approach

Szabolcs Suveges ¹, Kismet Hossain-Ibrahim ^{2,3}, J. Douglas Steele ⁴, Raluca Eftimie ⁵ and Dumitru Trucu



brain & tumour



...where Diffusion Tensor Imaging (DTI) scans were used to estimate the anisotropic cell diffusion term ; T1 weighted images => image segmentation => white/gray matter densities

Effective Treatment of Glioblastoma Multiforme With Oncolytic Data time t1 **Virotherapy: A Case-Series** prediction Data time t2

Benjamin Gesundheit^{1*}, Eliel Ben-David², Yehudit Posen¹, Ronald Ellie¹ Guido Wollmann^{3,4}, E. Marion Schneider⁵, Karl Aigner⁶, Lars Brauns⁷, Thomas Nesselhut⁸, Ingrid Ackva⁹, Christine Weisslein^{*} and Arno Thaller⁹





Mar 2013 Jan 2012



Jul 2012



Sept 2014

^{0.39} For the future...

- Macroscale-level spatial and spatio-temporal data is available
- For model validation & quantitative predictions we need also **microscale-level** data

Summary:

- Viral dynamics involves different multi-scale aspects that can be incorporated into mathematical/computational models
 - But multiscale (spatial & spatio-temporal) data not always available... (or if available: very few data points => no ML)
 - Technical aspects associated with parameter identification in these multi-scale models...
 - Open problems associated with the modelling of multi-scale within-host/betweenhost dynamics ...
- Many (complex) mathematical models are used only for qualitative exploration of possible model dynamics
 - Quantitative predictions require more data (& new computational approaches) to estimate parameters/functions