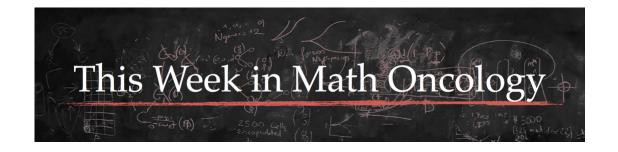


This week in MathOnco 191

1 message

This week in Mathematical Oncology <hisweekmathonco@substack.com>
To: cristian.axenie@gmail.com <cristian.axenie@gmail.com>

Today, 14:01



This week in MathOnco 191

Vasculogenesis, spatial distribution models, precision dosing, collateral sensitivity, 2021 in review, and more.

Jeffrey West, Maximilian Strobl, and Sandy Anderson Jan 6 \bigcirc \bigcirc \bigcirc

"This week in Mathematical Oncology" - Jan. 6, 2022

> <u>mathematical-oncology.org</u>

From the editor:

Happy New Year! The Mathematical Oncology team is expanding. Please welcome Saskia Haupt (Heidelberg University), who will be helping us keep the mathematical-oncology.org website up-to-date. Thanks for volunteering Saskia!

Jeffrey West

jeffrey.west@moffitt.org

PS, don't miss David Basanta's "Year in Review" blog post, providing a summary all the exciting posts from 2021.



In order to better plan out additional website features, we are asking you to complete this short survey (~5 minutes). The first section (1 page) is about the newsletter, and the second is about the website (1 page).

MathOnco Survey

We have many exciting plans for 2022, but we want to hear from you. For example - what would you rather us work on next:

- Lab pages featuring MathOnco folks around the world
- Search bar: search MathOnco web page by topic
- Calendar of seminars/abstracts/deadlines/...
- A collection of educational resources (recordings of tutorials, lectures, lecture material, ...)

Please let us know by filling out the survey, found here.

Together, we continue to build up the field of mathematical oncology!



1. A novel nonlocal partial differential equation model of endothelial progenitor cell cluster formation during the early stages of

vasculogenesis

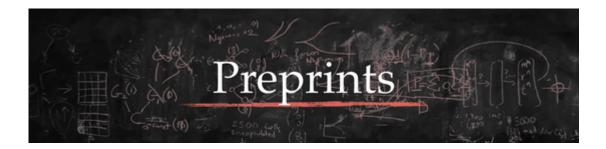
Chiara Villa, Alf Gerisch, Mark A. J. Chaplain

Extinction of Bistable Populations is Affected by the Shape of their Initial Spatial Distribution

Yifei Li, Stuart T. Johnston, Pascal R. Buenzli, Peter van Heijster, Matthew J. Simpson

- 3. Digital Evolution for Ecology Research: A Review Emily Dolson, Charles Ofria
- 4. A continued learning approach for model-informed precision dosing: updating models in clinical practice

 Corinna Majer, Jana de Willes, Niklas Hartung, Charlotte Kloft, Wilhelm.
 - Corinna Maier, Jana de Wiljes, Niklas Hartung, Charlotte Kloft, Wilhelm Huisinga
- 5. A phenotype-structured model to reproduce the avascular growth of a tumor and its interaction with the surrounding environment Giada Fiandaca, Sara Bernardi, Marco Scianna, Marcello Edoardo Delitala
- Combination of antiangiogenic treatment with chemotherapy as a multi-input optimal control problem Urszula Ledzewicz, Heinz Schättler
- 7. Dynamic Phenotypic Switching and Group Behavior Help Non-Small Cell Lung Cancer Cells Evade Chemotherapy
 Arin Nam, Atish Mohanty, Supriyo Bhattacharya, Sourabh Kotnala, ...,
 Herbert Levine, Mohit Kumar Jolly, Prakash Kulkarni, Ravi Salgia
- 8. Data-Driven Discovery of Mathematical and Physical Relations in Oncology Data Using Human-Understandable Machine Learning Daria Kurz, Carlos Salort Sánchez, Cristian Axenie



Bridging scales in a multiscale pattern-forming system
 Laeschkir Würthner, Fridtjof Brauns, Grzegorz Pawlik, Jacob Halatek,
 Jacob Kerssemakers, Cees Dekker, Erwin Frey

Interpreting dN/dS under different selective regimes in cancer evolution

Andrés Pérez-Figueroa, David Posada

- Dynamic collateral sensitivity profiles highlight challenges and opportunities for optimizing antibiotic sequences Jeff Maltas, Kevin B Wood
- 4. Mathematical modelling, selection and hierarchical inference to determine the minimal dose in IFN α therapy against Myeloproliferative Neoplasms

Gurvan Hermange, William Vainchenker, Isabelle Plo, Paul-Henry Cournède

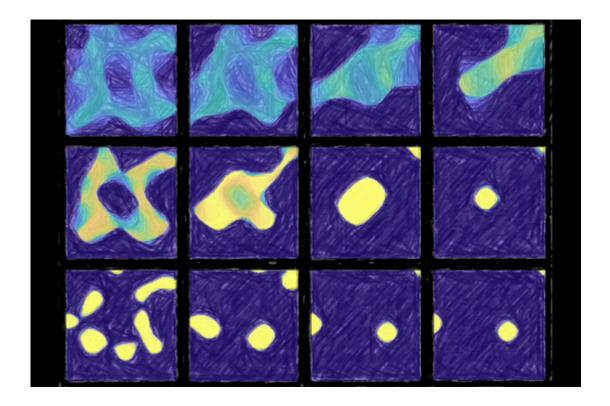


1. Biological Modeling

Phillip Compeau: A free course in modeling biological systems at multiple scales



The newsletter now has a dedicated homepage where we post the cover artwork for each issue. We encourage submissions that coincide with the release of a recent paper from your group.



Caption: We developed a novel nonlocal partial differential equation model of the early stages of cluster-based vasculogenesis in order to disentangle the mechanisms responsible for the formation and size of the endothelial progenitor cell (EPC) clusters characterising this neovascularisation process with big therapeutic potential (e.g. in tumours and ischemia). In this figure we show snapshots of numerical simulations suggesting that increasing matrix degradation (top-to-bottom), likely linked with hypoxia, might speed up cluster formation (EPC density plot, time increasing left-to-right; figures have been digitally enhanced) without affecting cluster size. For more detail, read here.

Created by: Chiara Villa



Visit the mathematical oncology page to view jobs, meetings, and special issues. We will post new additions here, but the full list can found at mathematical-oncology.org.

- 1. Jobs
- 2. Conferences / Meetings
- 3. Special issues



If you liked this post from This week in Mathematical Oncology, why not share it?



