CHIARA VILLA

Sorbonne Université & Postdoc & Mathematical Biology

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Website https://chiaravilla.github.io/website/index.html

Languages Italian, English, French

Software MATLAB, Python, LaTeX, Fortran90, COMSOL, Maple, R, HTML5, MS Office Memberships European Society for Mathematical and Theoretical Biology, Society for Mathematical

Biology, Société de Mathématiques Appliquées et Industrielles

ACADEMIC APPOINTMENTS

01/23 - today	PRFP postdoc, Laboratoire Jacques-Louis Lions, Sorbonne Université, Paris (FR)
	Laureate of the Paris Region Fellowship Programme (PR and EU MSCA funding)
	Project "Mechanistic modelling of cell migration and cancer invasion"
04/22 - 12/22	Postdoc, Laboratoire Jacques-Louis Lions, Sorbonne Université, Paris (FR)
,	Postdoctoral researcher in the group of Prof Benoît Perthame (ERC ADORA funding)

EDUCATION & RESEARCH EXPERIENCE

09/18 - 03/22	PhD, Mathematics, University of St Andrews, St Andrews (UK)
	Supervisors: Prof Mark Chaplain, Dr Tommaso Lorenzi
	Thesis title: 'Partial differential equation modelling in cancer and development'
2014 - 2018	MMaths, Applied Mathematics, University of St Andrews, St Andrews (UK)
	Fast Track, First Class Honours awarded.
Summer 2017	Undergraduate Summer Research Internship, University of St Andrews
Summer 2016	Complex Systems Biology Research Internship, Università degli Studi di Torino

FUNDING, GRANTS AND PRIZES AWARDED

2023	BOUM SMAI funding for the organisation to the workshop "Mathematical challenges in modelling cancer dynamics" in Paris in October 2023 (€1000).
2022	
2023	PEPS JCJC funding for the project "Conservative numerical schemes for novel struc-
	tured PDE models of cancer invasion" with Alexandre Poulain (€4900).
2023	UFR funding for the organisation to the workshop "Mathematical challenges in mod-
	elling cancer dynamics" in Paris in October 2023 (€1500).
2022	Junior Fellowship for the participation to the workshop "Parabolic and kinetic mod-
	els in population dynamics" in Toulouse in September 2022.
2022	Paris Region Fellowship Programme laureate with the proposed project on
	"Mechanistic models of cell migration and cancer invasion" (€257760).
2021	IHP financial support awarded by the Institute Henri Poincaré for the participation
	to the "Mathematical modeling of organization in living matter" thematic program in
	Paris during $10/01-01/04\ 2022\ (\text{€4500})$.
2020	SMBdevBio Poster Prize 1 awarded by the Society for Mathematical Biology,
_0_0	Developmental Biology subgroup, at the online SMB2020 meeting (\$250)
2020	LMS ECR Travel Grant awarded by the London Mathematical Society to attend
2020	the 12th European Conference on Mathematical and Theoretical Biology (£500)
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2018	PhD funding awarded by the School of Mathematics and Statistics, UoStA
	$(\pounds 49124.25)$
2018	The Principal's Scholarship for Academic Excellence, prize awarded to the top
	50 academically performing students in their final year at the $UoStA(£1000)$
2014 - 2018	The Deans' list, annual award for academic excellence by the Deans of the UoStA
2017	Research scholarship awarded by the UoStA to participate in the Undergraduate
	Summer Research Internship (£1684.29)
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PROFESSIONAL RESPONSIBILITIES AND OUTREACH

*School of M	Iathematics	and	Statistics.	University	of	St	Andrews
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01/23 - today	Member of 'Comité Parité', Laboratoire Jacques-Louis Lions, Sorbonne Université
10/22 - today	Postdoctoral Research Rep, Laboratoire Jacques-Louis Lions, Sorbonne Université
03/21 - today	Journal Peer Reviewer, Bulletin of Mathematical Biology, iScience, Mathemati-
	cal Biosciences , International Journal of Non-Linear Mechanics, European Control
	Conference 2022, Frontiers in Ecology and Evolution (Special issue: From Ecology to
	Cancer Biology and Back Again)
10/23	Mathematical challenges in modelling cancer dynamics workshop organiser,
	Laboratoire Jacques-Louis Lions, Sorbonne Université, https://mc2d.sciencesconf.org/
09/20 - 12/21	StAMBio seminar organiser*, Weekly talks by internal and guest speakers, online
11/20	Piscopia Society*, PhD testimonial to encourage female/non-binary students who
	are considering a PhD in mathematics, promoting equality and diversity in STEM
01/20	${\bf Postgraduate\ Interdisciplinary\ Mathematics\ Symposium\ organiser}^*, Edzell$
09/18 - 09/19	Postgraduate Research Rep & Postgraduate Research Executive Rep*
09/18 - 09/19	University of St Andrews Student Rep, SMSTC
11/18	Outreach event*, Organiser and speaker at the event 'MT234 Research and Party'

MENTORING, TEACHING AND MARKING

All activities of 2017-2022 undertaken with the School of Mathematics and Statistics, University of St Andrews. Teaching activities undertaken with groups of 50 (demonstrating) or 11 (tutoring) students. Feedback on Explanation (E), Organisation (O) and Availability (A) on a scale of 1 (excellent) to 5 (poor).

01/23 - 08/23	Master thesis supervision of Federica Padovano (EPFL), at LJLL (SU)
09/18 - 06/22	Mentor in Peer Mentoring scheme of 4 Undergraduate, 3 Master, 2 PhD students
Autumn 2020	MT2000 Computing Workshop, Demonstrator of computing in Python
Autumn 2019	MT2000 Computing Workshop, Demonstrator of computing in Python
Autumn 2019	MT2501 Linear Mathematics, Tutor of 2 groups (E=1.44, O=1.33, A=1.33)
Spring 2019	MT2507 Mathematical Modelling, Tutor of 2 groups (E=1.45, O=1.85, A=1.45),
	Demonstrator of 3 groups
Autumn 2018	MT2503 Multivariate Calculus, Tutor of 2 groups (E=1.17, O=1.5, A=1.17)
Autumn 2018	MT2504 Combinatorics and Probability, Marking of 100 computing projects
Autumn 2017	UK Undergraduate Ambassadors Scheme, weekly teaching assistance and activ-
	ities with secondary school pupils (S1, S3, Advanced Higher Maths), UoSA module
	ID4001 - Communication and Teaching in Science, Waid Academy, Anstruther (UK)

SELECTED SCIENTIFIC MEETINGS

Scientific meetings where I was invited to present my research (full list of meetings attended on my website)

May 2024	Mathematical Biology Seminar
	University of Leeds, Leeds
April 2024	Mathematical and numerical tools for Oncology Workshop
	Oncolille Institut, Lille
April 2024	Seminar of the Puissant Lab
	St Louis Research Institute, Saint-Louis Medical Center, Paris
Nov 2023	Mechanistic models for continuous phenotypic adaptation Workshop
	University of Leeds, Leeds
Aug 2023	SIMAI 2023
	Minisymposium, SIMAI, Matera
Jun 2023	Mathematical Biology: Analysis and Application Workshop
	Technische Universität Dresden, Dresden

May 2023	The Evolution Seminar
	Bielefeld University (Evolutionary Biology group), Bielefeld
Apr 2023	Seminar 'Analyse Numérique et Équations aux Dérivées Partielles'
	Université de Lille (Laboratoire Paul Painlevé), Lille
Feb 2023	Multiscale analysis and methods for PDEs Workshop
	Institute for Mathematical Sciences, Singapore
Nov 2022	Synthsys Seminar
	Centre for Synthetic and Systems Biology, Edinburgh
Oct 2022	Modelling cell and tissue biomechanics Workshop
	Sorbonne University (LJLL), Paris
Sep 2022	12th European Conference for Mathematical and Theoretical Biology
	Minisymposium, ESMTB, Heidelberg
Jun 2021	SoftMech Workshop
	University of St Andrews, Online
May 2021	Mathematical Biology on the Mediterranean Coast
	Sorbonne University (LJLL), Online
Jun 2020	Interplay between Oncology, Mathematics and Numerics
	Sorbonne University (LJLL), Inserm, University of Poitiers, Online conference
Dec 2019	Scottish Mathematical Biology Forum
	Maxwell Institute for Mathematical Sciences, Edinburgh

MAJOR RESEARCH OUTPUTS

Preprints

- [10] F. Padovano, L. Almeida, C. Villa, The development of drug resistance in metastatic tumours under chemotherapy: an evolutionary perspective, 2024. hal-04595087
- [9] L. Almeida, A. Poulain, A. Pourtier, C. Villa, Mathematical modelling of the contribution of senescent fibroblasts to basement membrane digestion during carcinoma invasion, 2024. hal-04574340

Papers published in peer-reviewed journals

- [8] C. Villa, M.A.J. Chaplain, T. Lorenzi, Modelling phenotypic heterogeneity in vascularised tumours, SIAM Journal on Applied Mathematics, 81, 434–453, 2021. DOI: 10.1137/19M1293971, hal-04415631.
- [7] C. Villa, M.A.J. Chaplain, T. Lorenzi, Evolutionary dynamics in vascularised tumours under chemotherapy: Mathematical modelling, asymptotic analysis and numerical simulations, *Vietnam Journal of Mathematics*, 49, 143–167, 2021. DOI: 10.1007/s10013-020-00445-9, hal-04415601.
- [6] C. Villa, M.A.J. Chaplain, A. Gerisch, T. Lorenzi, Mechanical models of pattern and form in biological tissues: the role of stress-strain constitutive equations, *Bulletin of Mathematical Biology*, 83:80, 2021. DOI: 10.1007/s11538-021-00912-5, hal-04415645.
- [5] F. Mottes, C. Villa, M. Osella, M. Caselle, The impact of whole genome duplications on the human gene regulatory networks, *PLOS Computational Biology*, 17(12):e1009638, 2021. DOI: 10.1371/journal.pcbi.1009638 hal-04415666.
- [4] C. Villa, A. Gerisch, M.A.J. Chaplain, A novel nonlocal partial differential equation model of endothelial progenitor cell cluster formation during the early stages of vasculogenesis, *Journal of Theoretical Biology*, 534(1):110963, 2022. DOI: 10.1016/j.jtbi.2021.110963, hal-04415625.
- [3] L. Almeida, J.A. Denis, N. Ferrrand, T. Lorenzi, M. Sabbah, C. Villa, Evolutionary dynamics of glucose-deprived cancer cells: insights from experimentally-informed mathematical modelling, *Journal of the Royal Society Interface*, 21(210):20230587, 2024. DOI: 10.1098/rsif.2023.0587, hal-03947209v2.

Conference proceedings

[2] T. Lorenzi, F.R. Macfarlane, C. Villa, Discrete and continuum models for the evolutionary and spatial dynamics of cancer: a very short introduction through two case studies, (pp. 359-380) in *Trends in Biomathematics: Modeling Cells, Flows, Epidemics, and the Environment*, Ed. R. Mondaini, Springer, Cham, 2019. DOI: 10.1007/978-3-030-46306-9_22, hal-04415585.

Doctoral thesis

[1] C. Villa, Partial differential equation modelling in cancer and development, PhD thesis, University of St Andrews, St Andrews, 2022. HAL Id: tel-04442733.

Open access code

- [C3] A. Poulain and C. Villa. TumInvasion-BM: Simulation of the rupture of the basement membrane by the effect of tumor cells, Matlab code. https://github.com/alexandrepoulain/TumInvasion-BM, 2024. BSD 2-Clause License.
- [C2] C. Villa and A. Gerisch. Villaetal2021bullmathbio: simulate mechanical models of pattern formation (Matlab code). https://git-ce.rwth-aachen.de/alf.gerisch/VillaEtAl2021BullMathBiol, 2021. GNU General Public License.
- [C1] C. Villa. Almeidaetal2023evolutionary: calibrate phenotype-structured equation models with experimental data (Matlab code). https://github.com/ChiaraVilla/AlmeidaEtAl2023Evolutionary, 2023. GNU General Public License.