

# CHIARA VILLA

Sorbonne Université  $\diamond$  Postdoc  $\diamond$  Mathematical Biology

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<b>Website</b>	<a href="https://chiaravilla.github.io/website/index.html">https://chiaravilla.github.io/website/index.html</a>
<b>Languages</b>	Italian, English, French
<b>Software</b>	MATLAB, Python, LaTeX, Fortran90, COMSOL, Maple, R, HTML5, MS Office
<b>Memberships</b>	European Society for Mathematical and Theoretical Biology, Society for Mathematical Biology, Société de Mathématiques Appliquées et Industrielles

## ACADEMIC APPOINTMENTS

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01/23 - today	<b>PRFP postdoc</b> , <i>Laboratoire Jacques-Louis Lions, Sorbonne Université</i> , 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	<b>Postdoc</b> , <i>Laboratoire Jacques-Louis Lions, Sorbonne Université</i> , Paris (FR) Postdoctoral researcher in the group of Prof Benoît Perthame (ERC ADORA funding)

## EDUCATION & RESEARCH EXPERIENCE

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

## FUNDING, GRANTS AND PRIZES AWARDED

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2023	<b>BOUM SMAI funding</b> for the organisation to the workshop “Mathematical challenges in modelling cancer dynamics” in Paris in October 2023 ( <b>€1000</b> ).
2023	<b>PEPS JCJC</b> funding for the project “Conservative numerical schemes for novel structured PDE models of cancer invasion” with Alexandre Poulain ( <b>€4900</b> ).
2023	<b>UFR funding</b> for the organisation to the workshop “Mathematical challenges in modelling cancer dynamics” in Paris in October 2023 ( <b>€1500</b> ).
2022	<b>Junior Fellowship</b> for the participation to the workshop “Parabolic and kinetic models in population dynamics” in Toulouse in September 2022.
2022	<b>Paris Region Fellowship Programme</b> laureate with the proposed project on “Mechanistic models of cell migration and cancer invasion” ( <b>€257760</b> ).
2021	<b>IHP financial support</b> 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 ( <b>€4500</b> ).
2020	<b>SMBdevBio Poster Prize 1</b> awarded by the Society for Mathematical Biology, Developmental Biology subgroup, at the online SMB2020 meeting ( <b>\$250</b> )
2020	<b>LMS ECR Travel Grant</b> awarded by the London Mathematical Society to attend the 12th European Conference on Mathematical and Theoretical Biology ( <b>£500</b> )
2018	<b>PhD funding</b> awarded by the School of Mathematics and Statistics, UoStA ( <b>£49124.25</b> )
2018	<b>The Principal’s Scholarship for Academic Excellence</b> , prize awarded to the top 50 academically performing students in their final year at the UoStA ( <b>£1000</b> )
2014 - 2018	<b>The Deans’ list</b> , annual award for academic excellence by the Deans of the UoStA
2017	<b>Research scholarship</b> awarded by the UoStA to participate in the Undergraduate Summer Research Internship ( <b>£1684.29</b> )

## PROFESSIONAL RESPONSIBILITIES AND OUTREACH

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*\*School of Mathematics and Statistics, University of St Andrews*

07/24	<b>Minisymposium organiser</b> at <i>ECMTB2024</i> on ‘Recent advances in modelling cancer invasion’
01/23 - today	<b>Member of ‘Comité Parité’</b> , <i>Laboratoire Jacques-Louis Lions, Sorbonne Université</i>
10/22 - today	<b>Postdoctoral Research Rep</b> , <i>Laboratoire Jacques-Louis Lions, Sorbonne Université</i>
03/21 - today	<b>Journal Peer Reviewer</b> , <i>Bulletin of Mathematical Biology, iScience, Mathematical Biosciences, International Journal of Non-Linear Mechanics, European Control Conference 2022, Frontiers in Ecology and Evolution</i> (Special issue: From Ecology to Cancer Biology and Back Again)
10/23	<b>Mathematical challenges in modelling cancer dynamics workshop organiser</b> , <i>Laboratoire Jacques-Louis Lions, Sorbonne Université</i> , <a href="https://mc2d.sciencesconf.org/">https://mc2d.sciencesconf.org/</a>
09/20 - 12/21	<b>StAMBio seminar organiser*</b> , Weekly talks by internal and guest speakers, online
11/20	<b>Piscopia Society*</b> , PhD testimonial to encourage female/non-binary students who are considering a PhD in mathematics, promoting equality and diversity in STEM
01/20	<b>Postgraduate Interdisciplinary Mathematics Symposium organiser*</b> , <i>Edzell</i>
09/18 - 09/19	<b>Postgraduate Research Rep &amp; Postgraduate Research Executive Rep*</b>
09/18 - 09/19	<b>University of St Andrews Student Rep</b> , <i>SMSTC</i>
11/18	<b>Outreach event*</b> , Organiser and speaker at the event ‘MT234 Research and Party’

## MENTORING, TEACHING AND MARKING

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

## SELECTED SCIENTIFIC MEETINGS

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

July 2024	<b>13th European Conference for Mathematical and Theoretical Biology</b> Minisymposium, <i>ESMTB &amp; University of Castilla La Mancha</i> , Toledo
July 2024	<b>European Congress of Mathematics</b> Minisymposium, <i>ECM &amp; Universidad de Sevilla</i> , Sevilla
May 2024	<b>Mathematical Biology Seminar</b> <i>University of Leeds</i> , Leeds
April 2024	<b>Mathematical and numerical tools for Oncology Workshop</b> <i>Oncolille Institut</i> , Lille
April 2024	<b>Seminar of the Puissant Lab</b> <i>St Louis Research Institute</i> , Saint-Louis Medical Center, Paris

Nov 2023	<b>Mechanistic models for continuous phenotypic adaptation Workshop</b> <i>University of Leeds, Leeds</i>
Aug 2023	<b>SIMAI 2023</b> Minisymposium, <i>SIMAI</i> , Matera
Jun 2023	<b>Mathematical Biology: Analysis and Application Workshop</b> <i>Technische Universität Dresden, Dresden</i>
May 2023	<b>The Evolution Seminar</b> <i>Bielefeld University (Evolutionary Biology group), Bielefeld</i>
Apr 2023	<b>Seminar ‘Analyse Numérique et Équations aux Dérivées Partielles’</b> <i>Université de Lille (Laboratoire Paul Painlevé), Lille</i>
Feb 2023	<b>Multiscale analysis and methods for PDEs Workshop</b> <i>Institute for Mathematical Sciences, Singapore</i>
Nov 2022	<b>Synthsys Seminar</b> <i>Centre for Synthetic and Systems Biology, Edinburgh</i>
Oct 2022	<b>Modelling cell and tissue biomechanics Workshop</b> <i>Sorbonne University (LJLL), Paris</i>
Sep 2022	<b>12th European Conference for Mathematical and Theoretical Biology</b> Minisymposium, <i>ESMTB</i> , Heidelberg
Jun 2021	<b>SoftMech Workshop</b> <i>University of St Andrews, Online</i>
May 2021	<b>Mathematical Biology on the Mediterranean Coast</b> <i>Sorbonne University (LJLL), Online</i>
Jun 2020	<b>Interplay between Oncology, Mathematics and Numerics</b> <i>Sorbonne University (LJLL), Inserm, University of Poitiers, Online conference</i>

## MAJOR RESEARCH OUTPUTS

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### Preprints

- [13] A.P. Browning, R. Crossley, C. Villa, P. K. Maini, A.L. Jenner, T. Cassidy and S. Hamis, Identifiability of heterogeneous phenotype adaptation from low-cell-count experiments and a stochastic model, 2024. **bioRxiv 2024.08.19.608540**
- [12] S. Hamis, A.P. Browning, A.L. Jenner, C. Villa, P. K. Maini and T. Cassidy, Growth rate-driven modelling reveals how phenotypic adaptation drives drug resistance in BRAFV600E-mutant melanoma 2024. **bioRxiv 2024.08.14.607616**
- [11] C. Villa, P. K. Maini, A.P. Browning, A.L. Jenner, S. Hamis and T. Cassidy, Reducing phenotype-structured PDE models of cancer evolution to systems of ODEs: a generalised moment dynamics approach, 2024. **hal-04599519**
- [10] F. Padovano, 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/jour-

nal.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.