
BIOGRAPHICAL SKETCH

NAME: Olivier Pertz

POSITION TITLE: Associate Professor for Cellular Dynamics

| INSTITUTION AND LOCATION | DEGREE | Completion Date MM/YYYY | FIELD OF STUDY |
|---|-----------------|----------------------------|----------------|
| University of Lausanne, CH | Diploma | 1995 | Biology |
| University of Basel, CH | PhD | 1999 | Biophysics |
| Scripps Research Institute, CA, USA | Postdoc | 2004 | Cell Biology |
| University of California San Diego, CA, USA | Staff scientist | 2007 | Cell Biology |
| University of Basel, CH | Assistant Prof | 2008 | Cell Biology |
| University of Bern, CH | Associate Prof | 2016 | Cell Biology |
| University of Bern, CH | Full Prof | 2024 | Cell Biology |

Lab website: <https://www.pertzlab.net/>

Google scholar ID : <https://scholar.google.ch/citations?user=dYonf8AAAAJ&hl=en&oi=ao>

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Personal Statement

My research focuses on the spatio-temporal signaling events that regulate Cell Morphogenesis and Fate Decisions. The main hypothesis relevant to our research is that these signaling events are highly dynamic, are precisely regulated in time and space, and can be highly heterogeneous within distinct cells of a population. An important current limitation in our field is that this spatio-temporal resolution of signaling is missed in classic biochemical methods that average populations of thousands of cells, or that rely on the analysis of static, steady-states. To address these challenges, we devise novel quantitative approaches to measure/manipulate signaling dynamics at biologically relevant time/length scales. This includes the development of multiplexed biosensor systems to image signaling dynamics at the single cell level, the development of optogenetic and microfluidic actuators to manipulate single cells with unprecedented spatio-temporal resolution, computer vision approaches to automate image analysis of large datasets, statistical analysis and mathematical modelling to make sense of the large datasets we produce. Our research provides new insights to target oncogenic signaling during cancer, as well as aberrant morphogenetic processes such as cell migration and invasion, and neuronal development and regeneration.

Employment history

2024- : Full Professor

2016-2023 : Associate Professor, Institute of Cell Biology, University of Bern, Switzerland

2014-2015 : Assistant Professor, Department of Biomedicine, University of Basel, Switzerland

2008-2013 : SNF Professor, Department of Biomedicine, University of Basel, Switzerland

2004-2007 : Staff scientist in the group of Richard Klemke, University of California San Diego, USA

2001-2004 : Postdoc in the group of Klaus Hahn, The Scripps Research Institute, La Jolla, CA, USA

2000-2001 : Postdoc in the group of Glen Nemerow, The Scripps Research Institute, La Jolla, CA, USA

1995-1999 : PhD thesis in the group of Juergen Engel, Biocenter of the University Basel, Basel, CH

Honors and awards

12/2015: Best publication of the year award, Dept. of Biomedicine, University of Basel

03/2005: Special symposium award at Roche Symposium for leading bioscientists

07/2004-06/2006: Philip Morris External Research Program Postdoctoral Fellowship

07/2002-06/2004: Swiss National Science Foundation Advanced Postdoctoral Fellowship

07/2001-06/2002: Roche Research Foundation and Novartis Postdoctoral Fellowships

01/2000-06/2001: Swiss National Science Foundation Postdoctoral Fellowship

Research projects as principal investigator (last 5 years)

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| 2016-2018 | ca 200 k\$. | Systems X.ch. Deciphering a prototypical receptor tyrosine kinase signaling network at the single cell level using a genetically-encodable optogenetic circuit. |
| 2016-2018 | ca 600 k\$ | Swiss National Science Foundation. Mapping Rho GTPase signaling networks through acute, dynamic perturbation of spatio-temporal signaling fluxes. |
| 2016-2018 | ca 250 k\$ | Swiss National Science Foundation joint. Understanding single cell-level MAPK activation dynamics for manipulation of neuronal stem cell self-renewal and differentiation fates. |
| 2016-2018 | ca 360 k\$ | Swiss cancer league. A tumor on a chip approach to understand signaling networks mediating melanoma drug resistance at the single cell level. |
| 2017-2019 | ca 250 k\$ | Swiss National Science Foundation. Dissecting a Rho GTPase spatio-temporal signaling network regulating growth cone motility, neurite and axonal outgrowth. |
| 2018 | ca 350 k\$ | Swiss National Science Foundation R'equip. A high content confocal microscope for fast 3D imaging of living samples. |
| 2018-2019 | ca 135 k\$. | UniBE interdisciplinary grant. An integrated microscopy/computer vision platform for high content 3D imaging of aberrant proliferative signalling in cancer organoids. |
| 2019-2023 | ca 759 k\$ | Swiss National Science Foundation. Decoding and Re-encoding Receptor Tyrosine Kinase Fate decision signalling. |
| 2019-2024 | ca 656 k\$ | Swiss National Science Foundation. Real time Exploration of GTPase-Cytoskeletal feedback underlying Contractile Actomyosin Systems. |
| 2019-2023 | 350 k\$ | Human Frontier Science Program. A spatiotemporal map of signaling processes controlling human stem cell renewal and differentiation. |
| 2020 | ca 60 k\$ | Novartis Foundation. Realtime Control of Fate Decision Signaling to understand and manipulate Epithelial Homeostasis. |
| 2021-2024 | ca 227 k\$ | Uniscientia Foundation. Real time feedback microscopy to identify emergent properties of an epithelial ecosystem. |
| 2022 | ca 25 k\$ | CZI. Democratizing Image Analysis with an Easy-to-Train Classifier. |
| 2023-2026 | ca 860 k\$ | Swiss National Science Foundation. Multiscale mapping of fate decision receptor Tyrosine kinase signaling. |

Memberships in panels and scientific reviewing activities

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| 2007 - | Peer reviewer of more than 70 original manuscripts (including <i>Science</i> , <i>PNAS</i> , <i>Elife</i> , <i>Journal of Cell Biology</i> , <i>Developmental Cell</i> , <i>Nature Communications</i> , ...) and more than 50 project grants (including Swiss, French, German, ... National Science Foundation and Cancer Leagues). |
| 2016 - | "Molecular Biology & Biochemistry" Expert Committee, Graduate School for Cellular and Biomedical Sciences, University of Bern |
| 2017 – 2022 | Science faculty representative at the board of the Microscopy Imaging Center |
| 2017 – 2022 | Science faculty representative at the board of the IT services |
| 2021 - | Member of the Precision Medicine Center and Bern Data Science Initiative at University of Bern |
| 2016 - | Member of multiple hiring committees |

Supervision of junior researchers

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| 2008-2015 | Department of Biomedicine, University of Basel: 7 postdocs, 4 PhD students, 1 master student |
| 2016- | Institute of Cell Biology, University of Bern: 5 postdocs, 9 PhD students, 8 master students, 15 bachelor students |

Invited speaker at international conferences (selected, last 5 years)

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| 2023 | European Light Microscopy Initiative meeting. Leiden, Netherlands. |
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2022 At the interface of cell fate and tissue dynamics. Muenster, Germany.
 3rd Franco-Japanese Developmental meeting. Strasbourg, France
 Inserm Optobiology workshop. Online.
 AIChE 3rd optogenetics technologies and applications. Heidelberg, Germany.
 Emerging Systems Medecine. Stockholm, Sweden
 2021 Spatio-temporal encoding and decoding in cell signaling. Online.
 2020 Conferences canceled due to COVID19
 2019 French optogenetics club meeting. Paris, France
 2018 European Cytoskeletal meeting. Prague, Czech Republic.
 Unito-Polito conference series in cancer. Turin, Italy.

List of publications (past five years)

Lu L., Guyomar T., Vagne Q., Berthoz R., Torres-Sánchez A., Lieb M., Martin-Lemaitre C., Van Unen K., Honigmann A., **Pertz O.**, Salbreux G., Riveline D. Polarity-driven three-dimensional spontaneous rotation of a cell doublet. In press at Nature physics.

<https://www.biorxiv.org/content/10.1101/2022.12.21.521355v1>

Spontaneous rotations in epithelia as an interplay between cell polarity and RhoA activity at boundaries. Lo Vecchio S., **Pertz O.**, Szopos M., Navoret L., Riveline D. In press at Nature Physics.

<https://www.biorxiv.org/content/10.1101/2021.11.11.468187v1>

Heydasch M., van Unen J., Hinderling L., Dobrzynski M., **Pertz O.** GTPase activating protein DLC1 spatio-temporally regulates Rho signaling. Reviewed preprint at Elife.

<https://elifesciences.org/reviewed-preprints/90305v1>

Pulfer A., Pizzagalli D.U., Gagliardi P.A., Hinderling L., Lopez P., Zayats P., Carrillo-Barbera P., Antonello P., Palomino-Segura M., Giusti A., Thelen M., Gambardella L.M., Murooka T.T., **Pertz O.**, Krause R., Fernandez Gonzalez S. Transformer-based spatial-temporal detection of apoptotic cell death in live-cell imaging. Reviewed Preprint at Elife.

<https://elifesciences.org/reviewed-preprints/90502>

Gagliardi P.A., Graedel B., Jacques M-A., Hinderling L., Pertz O., Dobrzynski M. ARCOS, Automatic Recognition of Collective Signaling Events. J Cell Biol. 2023 Oct 2;222(10):e202207048.

Preprint highlight: <https://www.molbiolcell.org/doi/full/10.1091/mbc.P22-08-1005>

Nałęcz-Jawecki P., Gagliardi P.A., Kochańczyk M., Dessauges C., **Pertz O.**, Lipniacki T. The MAPK/ERK channel capacity exceeds 6 bit/hour. PLoS Comput Biol 19(5): e1011155. <https://doi.org/10.1371/journal.pcbi.1011155>.

Duerr L., Reinhardt J., Dobrzyński M., Hell T., Smiesko M., **Pertz O.**, Hamburger M., Garo E. Dimerosesquiterpene and sesquiterpene lactones from Artemisia argyi inhibiting oncogenic PI3K/AKT signaling in melanoma cells. Journal of Nat. Prod. **2022** Nov 9.

Hell T., Dobrzyński M., Gröflina F., Reinhardt J., Duerr L., **Pertz O.**, Hamburger M., Wolfender J.L., Garo E. Flavonoids from Ericameria nauseosa inhibiting PI3K/AKT pathway in human melanoma cells. Biomedicine and Pharmacotherapy. Biomed Pharmacother. **2022** Oct 17;156:113754.

Ender P., Gagliardi P.A., Dobrzyński M., Dessauges C., Höhener T., Jacques M-A, Cohen A.R., **Pertz O.** Spatio-temporal Control of ERK Pulse Frequency Coordinates Fate Decisions during Mammary Acinar Morphogenesis. Dev Cell. **2022** Sep 7:S1534-5807(22)00594-9.

Preview in Dev Cell: [https://www.cell.com/developmental-cell/fulltext/S1534-5807\(22\)00631-1](https://www.cell.com/developmental-cell/fulltext/S1534-5807(22)00631-1)

Höhener T., Landolt A, Dessauges C, Gagliardi P.A, **Pertz O.** LITOS - a versatile LED illumination tool for optogenetic stimulation. Sci Rep. **2022** Jul 30;12(1):13139.

Dessauges C., Mikelson J., Dobrzyński M., Jacques M.A., Frismantien A., Gagliardi P.A., Khammash M., **Pertz O.** Optogenetic actuator - ERK biosensor circuits identify MAPK network nodes that shape ERK dynamics. Mol Syst Biol. **2022** Jun;18(6):e10670.

Hell T., Rutz A., Duerr L., Dobrzyński M., Reinhardt J., Lehner T., Keller M., John A., Gupta M.P., **Pertz O.**, Hamburger M., Wolfender J.L., Garo E. Combining HPLC-Based Activity Profiling with Advanced UHPLC-HRMS/MS Annotation to Accelerate the Discovery of Natural Products Targeting Oncogenic Signaling in Melanoma. J. Nat. Prod. **2022** May 31.

Mining of Single-Cell Signaling Time-Series for Dynamic Phenotypes with Clustering. Dobrzyński M, Jacques MA, **Pertz O**. *Methods Mol Biol*. **2022**;2488:183-206.

Dürr L., Hell T., Dobrzynski M., Mattei A., John A., Augsburg N., Bradanini G., Reinhardt J., Rossberg F., Drobnjakovic M., Gupta M.P., Hamburger M., **Pertz O**, Garo E. High Content Screening Pipeline for Natural Products Targeting Oncogenic Signaling in Melanoma. *J. Nat. Prod*. **2022** Apr 22;85(4):1006-1017.

Loewith R, Roux A., **Pertz O**. Chemical-Biology-derived in vivo Sensors: Past, Present, and Future. *Chimia (Aarau)*. **2021** Dec 22;75(12):1017-1021.

Gagliardi P.A., Dobrzyński M., Jacques M-A., Dessauges C., Ender P., Blum Y., Hughes R.M., Cohen A.R., **Pertz O**. Collective ERK/Akt activity waves orchestrate epithelial homeostasis by driving apoptosis-induced survival. *Developmental Cell*. **2021** May 28;S1534-5807(21)00436-6.

Highlighted in Science Signaling: <https://www.science.org/doi/10.1126/scisignal.abk1364>

Jacques M-A, Dobrzyński M., Gagliardi P.A., Sznitman R. and **Pertz O**. CODEX, a neural network approach to explore signaling dynamics landscapes. *Mol Syst Biol* (**2021**)17:e10026

Müller P.M., Rademacher J., Bagshaw R.D., Wortmann C., Barth C., van Unen J., Alp K.M., Giudice G., Eccles R.L., Heinrich L.E., Pascual-Vargas P., Sanchez-Castro M., Brandenburg L., Mbamalu G., Tucholska M., Spatt L., Czajkowski M.T., Welke R.W., Zhang S., Nguyen V., Rustemi T., Trnka P., Freitag K., Larsen B., Popp O., Mertins P., Gingras A.C., Roth F.P., Colwill K., Bakal C., **Pertz O**, Pawson T., Petsalaki E., Rocks O. Systems analysis of RhoGEF and RhoGAP regulatory proteins reveals spatially organized RAC1 signalling from integrin adhesions. *Nat Cell Biol*. **2020** Apr;22(4):498-511.

Dobrzynski M., Jacques M.A., **Pertz O**. Mining single-cell time-series datasets with Time Course Inspector. *Bioinformatics*, Volume 36, Issue 6, 15 March **2020**, Pages 1968–1969.

Blum Y.*, Mikelson J.*, Dobrzynski M.*, Ryu H., Jacques M.A., Jeon N.L., Khammash M., **Pertz O**. Temporal perturbation of ERK dynamics reveals network architecture of FGF2/MAPK signaling. *Mol Syst Biol*. **2019** Nov;15(11):e8947. Authors contributed equally.

Gagliardi P., **Pertz O**. Developmental ERK Signaling Illuminated. *Developmental Cell*. **2019** Feb 11;48(3):289-290.

Bagonis M., Fusco L., **Pertz O***, Danuser G.* Automated profiling of growth cone heterogeneity defines relations between morphology and motility. *J Cell Biol*. **2019** Jan 7;218(1):350-379. * Both authors contributed equally.

Olivier Pertz, 23-12-2023

Olivier Pertz