

ADRIEN JOLLY – Computational biologist

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GitHub : <https://github.com/AdrienJolly>

Education

PhD in Systems Biology

University of Heidelberg, Germany

(2015-2020)

Master in molecular and cellular biology

Université Pierre et Marie Curie (UPMC) and École Normale Supérieure (ENS), Paris France

Major in immunology (Pasteur course Advanced Immunology) and systems biology (ENS)

(2013-2015)

Bachelor in life sciences

Université Pierre et Marie Curie, Paris, France

(2010-2013)

Legal education

Master in European business Law

Université Paris XI, France

(2006-2007)

Integrated Programme in European Law

University of Warwick (England), Universität des Saarlandes (Germany)

Université de Lille 2 (France)

(2004-2006)

Bachelor in French Law

Université de Lille 2 (France)

(2001-2004)

Experience

April 2022- present : Medical scientist (computational) at Universitätsklinikum Frankfurt: shared position between the research groups Machine Learning in Oncology (DKTK) and Basic Mechanisms in Stem Cell Biology (Uniklinikum Frankfurt)

September 2020- March 2022 : postdoctoral researcher at the German Cancer Research Center, Division of Theoretical Systems Biology

September 2015- September 2020 : PhD at the German Cancer Research Center, Division of Theoretical Systems Biology (80% computational / 20% experimental work)

Supervision : Prof. Thomas Höfer

Impact of the microenvironment on mouse hematopoiesis and T cell development

January-June 2015 : Master thesis at Université de Versailles St Quentin en Yvelines

Supervision : Prof. Henri-Jean Garchon,

Characterization of Human Monocyte Derived Dendritic Cells activation in Ankylosing Spondylitis from Microarray data

July-August 2014 : Internship at hospital Pitié Salpêtrière in Paris, France

Supervision : Dr. Encarnita Mariotti-Ferrandiz, and Prof. Adrien Six

High throughput sequencing and bioinformatic analysis of T Cell Receptor repertoire

As a legal adviser :

2007 to 2012 : legal adviser in the Asset Management and Investment Banking industries

Research Interests

- Mechanistic modeling of biological systems
- determination of cell population dynamics *in vivo*
- deciphering the interactions between hematopoietic/leukemic cells with their microenvironment
- Immune repertoire analysis

Programming languages and computational skills

- R (bulk and single Omics data analysis, machine learning)
- Python (Numerical computing, package development, NGS data analysis)
- Matlab (simulation of mathematical models, parameter estimation)
- Julia (simulation of mathematical models, parameter estimation)
- Linux commands (regular usage of a computing cluster)

Experimental skills

- Mammalian Cell culture, Flow Cytometry, Cell Sorting
- Animal experiments (FELASA B Certificate)
- NGS Library Preparation and sequencing (Illumina)

Fellowships

- 2022 – present : Medical scientist fellowship from the Mildred-Scheel-Nachwuchszentrum Frankfurt (4 years postdoc salary + funding for consumables and travel)

Publications

Carvajal Ibañez D, Skabkin M, Hooli J, Cerrizuela S, Göpferich M, **Jolly A**, Volk K, Zumwinkel M, Bertolini M, Figlia G, Höfer T, Kramer G, Anders S, Teleman AA, Marciniak-Czochra A, Martin-Villalba A. Interferon regulates neural stem cell function at all ages by orchestrating mTOR and cell cycle. *EMBO Mol Med*. 2023

Jolly A, Fanti AK, Kongsaysak-Lengyel C, Claudino N, Gräßer I, Becker NB, Höfer T. CycleFlow simultaneously quantifies cell-cycle phase lengths and quiescence *in vivo*. *Cell Rep Methods*. 2022

Grassmann S., Mihatsch L., Mir J., Kazeroonian A., Rahimi R., Flommersfeld S., Schober K., Hensel I., Leube J., Pachmayr L. O., Kretschmer L, Zhang Q., **Jolly A.**, Chaudhry M. Z., Schiemann M., Cicin-Sain L., Höfer T., Busch D. H., Flossdorf M. & Buchholz V.R *Nature Immunology* 2020; Early emergence of T central memory precursors programs clonal dominance during chronic viral infection

Mende N*, **Jolly A***, Percin GI*, Günther M, Rostovskaya M, Krishnan SM, Oostendorp RAJ, Dahl A, Anastassiadis K, Höfer T, Waskow C. *Blood* 2019; Prospective isolation of nonhematopoietic cells of the niche and their differential molecular interactions with HSCs.

Becker NB, Günther M, Li C, **Jolly A**, Höfer T. *Journal of Theoretical Biology* 2019 ; Stem cell homeostasis by integral feedback through the niche

Arndt K, Kranz A, Fohgrub J, **Jolly A**, Bledau AS, Di Virgilio M, Lesche M, Dahl A, Höfer T, Stewart AF, Waskow C. *Blood* 2018 ; SETD1A protects HSCs from activation-induced functional decline *in vivo*.