

Philipp Ross, Ph.D.

HHMI Postdoctoral Scientist • Mendoza Lab • University of Chicago
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Research Summary

Biochemist with proven expertise in characterizing macromolecular structure and function relevant to human immunity including MHCs, KIRs, TCRs, antibodies, nanobodies, and Fabs. Extensive experience with recombinant protein expression, x-ray crystallography, and flow cytometry and proficiency with yeast-display mediated screening, protein engineering, and high-throughput cellular assays using functional readouts. Actively seeking positions that allow me to apply my expertise, contribute to cutting-edge projects, and work with and develop emerging technologies.

Education

University of Chicago, Ph.D. with Biochemistry Emphasis in Genetics, Genomics, & Systems Biology

University of Chicago, M.Sc. in Genetics, Genomics, & Systems Biology

Oct. 2016 - Mar. 2023

Laboratory of Erin J. Adams, Department of Biochemistry and Molecular Biology

Dissertation Title: "Structural and Cellular Properties of Archaic and Non-Classical HLA Molecules"

Committee Members: Carole Ober, Allan D. Drummond, Bana Jabri, Vincent J. Lynch

Bachelors of Science in Bioengineering

Cum laude | Binghamton University, Binghamton, NY USA

Sep. 2008 - May 2013

Research

HHMI Postdoctoral scientist

2023 - Present

Protein Engineering, University of Chicago

Advisor: Dr. Juan L. Mendoza

As a postdoctoral scientist, I use yeast display-based protein engineering methods to both study and enhance cytokine signaling pathways with the goal of developing better immunotherapeutics

Graduate student

2017 - 2023

Molecular Immunology, University of Chicago

Advisor: Dr. Erin J. Adams

As a graduate student, I used cellular, biochemical, and structural methods to understand how archaic and non-classical MHCs present peptides and interact with innate and adaptive immune cells in humans.

Computational biologist

2013 - 2016

Computational Biology, Pennsylvania State University

Advisor: Dr. Manuel Llinás

As a technician, I designed and utilized bioinformatic workflows to understand pre- and post-transcriptional regulation in the deadliest human infecting species of the parasite that causes malaria, *Plasmodium falciparum*.

Undergraduate researcher **Mathematical Modeling**, Binghamton University
2012 - 2013 **Advisor:** Dr. Hiroki Sayama

As an undergraduate student, I designed a computational simulation and graphical user interface looking at the socioeconomic consequences of the widespread adoption of 3D printers implemented in Mathematica and Python.

Funding, Fellowships, & Scholarships

2021 - 2026 **R01 Grant Award Recognition**, Contributor
PI: Dr. Erin Adams
Project Title: Molecular and functional investigation of the role of HLA-F in immune regulation

2020 - 2021 **R21 Grant Award Recognition**, Contributor
PI: Dr. Erin Adams
Project Title: Molecular characterization of the functional isoforms of HLA-F in human health and cancer

2020 **Not Awarded, F31 Predoctoral Fellowship** Score: 32 Percentile: 23
Title: Molecular mechanisms of HLA-F recognition at the maternal-fetal interface

2019 **Not Awarded, R.C. Lewontin Early Award**
Title: HLA-F, a non-classical MHC, in immunity, reproduction, and human evolution

2018 **Honorable Mention, NSF GRFP**
Title: HLA-F, a non-classical MHC, in immunity, reproduction, and human evolution

2009 - 2010 **SMART Grant**, Binghamton University

2008 **Kathleen Mallory Memorial Scholarship**, Earl L. Vandermeulen High School

Awards & Honors

2013 **Graduated Cum Laude**, Bachelors of Science in Bioengineering, Binghamton University

2013 **Member**, Tau Beta Pi Engineering Honor Society

Submitted manuscripts

Philipp Rosst[†], Hugo Hilton[†], Jane Lodwick, Tomasz Slezak, Lisbeth A. Guethlein, Curtis P. McMurtrey, Alex S. Han, Morten Nielsen, Daniel Yong, Kristof T. Nolan, Charles L. Dulberger William H. Hildebrand, Minglei Zhao, Anthony Kossiakoff, Peter Parham, Erin J. Adams “*Molecular characterization of the archaic HLA-B*73:01 allele reveals presentation of a unique peptidome and skewed engagement by KIR2DL2.*”
Under Review at JBC <https://www.biorxiv.org/content/10.1101/2024.11.25.625330v1>

Manuscripts in Preparation

Kristof Nolan[†], **Philipp Rosst**[†], Tomasz Slezak, Jane Lodwick, Jason Krawic, Curtis McMurtrey, Samuel Weng, Allen Huff, William Hildebrand, Anthony Kossiakoff, Erin J. Adams “*HLA-F is predominantly peptide-loaded on the surface of cells and may present KIR-permissive peptides.*”

Journal Publications

Lia Chappell, **Philipp Ross**, Lindsey Orchard, Timothy J Russell, Thomas D Otto, Matthew Berriman, Julian C Rayner, Manuel Llinás “*Refining the transcriptome of the human malaria parasite Plasmodium falciparum using amplification-free RNA-seq*” **BMC Genomics** 2020 <https://doi.org/10.1186/s12864-020-06787-5>

Munir Akkaya, Abhisheka Bansal, Patrick W Sheehan, Mirna Pena, Alvaro Molina-Cruz, Lindsey M Orchard, Clare K Cimperman, Chen-Feng Qi, **Philipp Ross**, Takele Yazew, Daniel Sturdevant, Sarah L Anzick, Girija Thiruvengadam, Thomas Dan Otto, Oliver Billker, Manuel Llinás, Louis H Miller, Susan K Pierce “*A single-nucleotide polymorphism in a Plasmodium berghei ApiAP2 transcription factor alters the development of host immunity*” **Science Advances** 2020 <https://doi.org/10.1126/sciadv.aaw6957>

Joana Mendonca Santos, **Philipp Ross***, Gabrielle Josling*, Preeti Joshi, Lindsey Orchard, Tracey Campbell, Ariel Schieler, Ileana M Cristea, Manuel Llinás “*Red blood cell invasion by the malaria parasite is coordinated by the PfAP2-I transcription factor*” **Cell Host & Microbe** 2017 <https://doi.org/10.1016/j.chom.2017.05.006>

Shiri Eshar, Lindsey Altenhofen, Alona Rabner, **Philipp Ross**, Yair Fastman, Yael Mandel-Gutfreund, Rotem Karni, Manuel Llinás, Ron Dzikowski “*PfSR1 controls alternative splicing and steady-state RNA levels in Plasmodium falciparum through preferential recognition of specific RNA motifs*” **Molecular microbiology** 2015 <https://doi.org/10.1111/mmi.13007>

Amber Ferger, Wai Lau, **Philipp Ross**, Wyman Zhao, Hiroki Sayama, Steen Rasmussen “*Impact of Personal Fabrication Technology on Social Structure and Wealth Distribution: An Agent-Based Simulation Study*” **MIT Press** 2013

Conferences & Seminars

Poster “*Correlating Binding Affinity and Signaling Strength in the JAK/STAT Pathway*” HHMI Science Meeting (Feb. 2025)

Virtual Talk, “*Molecular mechanisms of HLA-F recognition at the maternal-fetal interface*” Maternal/Fetal Interface Seminar Series (Oct. 2020)

Poster “*The unique peptidome and structure of the archaic HLA-B allele, HLA-B*73:01*” UChicago Molecular Biosciences Retreat (Oct. 2019)

Talk “*HLA-F, a non-classical MHC, in immunity, reproduction, and human evolution*” UChicago Molecular Biosciences Retreat (Oct. 2019)

Service

2024 - Present	Career Mentor , Nucleate Dojo - supporting undergraduate biotech talent
2023	Judge (Remote) , International Genetically Engineered Machine (iGEM)
2023 - Present	Primary Wet Lab Supervisor , UChicago iGEM (GeneHackers)
2020 - 2022	Graduate Student Advisor , UChicago iGEM (GeneHackers)
2019 - 2021	Graduate Program Student Representative , Deans Council, University of Chicago
2017 - 2019	Science Connections Volunteer , Museum of Science and Industry
2017 - Present	Resident Scientist , Skype a Scientist (http://www.skypeascientist.com)
2017	Judge , Spring Symposium & Student Research Conference in STEM
2010 - 2011	President , Binghamton Bioengineering Club, Binghamton University

Teaching & Mentorship

2023 - Present	Postdoctoral Supervisor Mentees: Zhijie Chen and Shima Shabani As a postdoctoral associate in the Mendoza lab, I mentor the graduate students on a daily basis and provide feedback of various kinds including experimental, professional, and more broadly scientific.
2023	Primary Wet Lab Supervisor Mentees: Sneha Agarwal, Tommy Walsh, and Clara Deimling I directly mentored three undergraduates with minimal wet lab experience during the summer of 2023 in order to generate data for the wet lab component of our 2023 iGEM project, Green Levodopa Optimised with Transaminases (Glow), for which we received a Gold Medal.
2019 - 2021	Undergraduate mentor Mentee: Daniel Yong I mentored and worked with Daniel to structurally characterize a rare MHC molecule known as HLA-B*73:01, a project which will lead to a co-author publication.
2019	T.A. - Evolution of Biological Molecules Instructors: Dr. Joe Thornton & Dr. Allan Drummond University of Chicago
2017	T.A. - Genetic Analysis of Model Organisms Instructors: Dr. Doug Bishop, Dr. Jocelyn Malamy, & Dr. Edwin (Chip) Ferguson University of Chicago

Preprint Peer Reviews

Aging represses lung tumorigenesis and alters tumor suppression

Reviewed by: **Philipp Ross** and Juan L. Mendoza

[10.5281/zenodo.12747023](https://zenodo.org/record/12747023)

Indels allow antiviral proteins to evolve functional novelty inaccessible by missense mutations

Reviewed by: **Philipp Ross**, William Grubbe, and Juan L. Mendoza

[10.5281/zenodo.11644732](https://zenodo.org/record/11644732)

Cryo-EM structure and biochemical analysis of human chemokine receptor CCR8

Reviewed by: **Philipp Ross**, William Grubbe, and Juan L. Mendoza

[10.5281/zenodo.11644389](https://zenodo.org/record/11644389)

Virion morphology and on-virus spike protein structures of diverse SARS-CoV-2 variants

Reviewed by: James Fraser, Luisa Vasconcelos, Liyi Cheng, Samantha Lish, S. Chan Baek, Lang Ding, Alexandra Probst, Naiya Phillips, William Grubbe, Youchen Guan, and **Philipp Ross**

[10.5281/zenodo.10779310](https://zenodo.org/record/10779310)