

Paul L. Maurizio

CONTACT INFORMATION	The University of Chicago Section in Genetic Medicine Knapp Center for Biomedical Discovery 900 E 57th Street, Room 9144 Chicago, Illinois 60637-1428	https://mauriziopaul.github.io/landing maurizio@uchicago.edu (914) 610-3984
EDUCATION	<input type="checkbox"/> Ph.D. , University of North Carolina at Chapel Hill (UNC-CH) Bioinformatics & Computational Biology, Department of Genetics, School of Medicine	05/2018
	<input type="checkbox"/> Sc.M. , Johns Hopkins University Molecular Microbiology & Immunology, Bloomberg School of Public Health	08/2011
	<input type="checkbox"/> B.A. , Swarthmore College Major in Biochemistry; Major in Religion	05/2005
ACADEMIC POSITIONS HELD	<input type="checkbox"/> Postdoctoral Scholar , University of Chicago Department of Medicine, Section in Genetic Medicine Supervisor: Luis B. Barreiro, Ph.D. Research Areas: single-cell RNA-seq; immunology; social status effects; functional genomics	08/2018–Present
	<input type="checkbox"/> Visiting Scholar , Johns Hopkins University Bloomberg School of Public Health, Department of Molecular Microbiology & Immunology Supervisor: Fidel Zavala, M.D. Research Areas: molecular parasitology transgenesis; preclinical vaccine/adjuvant testing	07/2011–06/2012
GRADUATE RESEARCH EXPERIENCES	<input type="checkbox"/> Graduate Research Assistant , UNC-CH • Advisors: Mark T. Heise, Ph.D. & William Valdar, Ph.D. • Committee: Terrence S. Furey, Ph.D. (chair); Fernando Pardo-Manuel de Villena, Ph.D.; Ralph S. Baric, Ph.D.; Jeremy E. Purvis, Ph.D. • Research Areas: Bayesian statistical modeling; virology; heritability; QTL mapping; RNA-seq	05/2013–05/2018
	<input type="checkbox"/> Graduate Research Assistant (Rotations), UNC-CH • Advisors: David M. Margolis, M.D.; Aravinda M. de Silva, Ph.D.; Kristina De Paris, Ph.D. • Research Areas: HIV-1 latency; dengue virus; antimalarial treatment response in infants	07/2012–05/2013
	<input type="checkbox"/> Graduate Research Assistant , Johns Hopkins University • Advisor: Douglas E. Norris, Ph.D. • Research Areas: population genetics; vector ecology; flavivirus infection in <i>Culex</i> mosquitoes	11/2009–05/2011
GRANTS, FELLOWSHIPS & SCHOLARSHIPS	<input type="checkbox"/> PI, UChicago Pilot Grant , for collaborative work within the Department of Medicine Proposal: “Modeling the Effects of Social Stress on Cell-to-Cell Variation in the Immune Response to Influenza Vaccination”	2019
	<input type="checkbox"/> Fellow, NIH T32 Fellowship , Virology Training Grant, UNC-CH	2015–2016
	<input type="checkbox"/> Scholar, Master’s Tuition Scholarship , JHU	2010–2011
	<input type="checkbox"/> Fellow, Joshua Lippincott Fellowship , Swarthmore College	2009–2010
	<input type="checkbox"/> Fellow, NSF Summer REU in Prokaryotic Biology , University of Georgia Advisor: Juergen Wiegand, Ph.D., Department of Microbiology	2004
	<input type="checkbox"/> Fellow, NASA Astrobiology Summer Program , The Pennsylvania State University Advisor: Hiroshi Ohmoto, Ph.D., Department of Geosciences	2003
HONORS & AWARDS	<input type="checkbox"/> Award, Travel , 2 nd Annual Science Policy Symposium at the Rockefeller University The National Science Policy Network, NYC, NY	11/2018
	<input type="checkbox"/> Award, Outstanding Oral Presentation , 15 th Annual CTC Meeting Complex Trait Community, Memphis, TN	06/2017

- ❑ **Award, Travel**, 2nd Penn Symposium on Mathematical & Computational Biology 05/2017
(*declined, unable to attend*), Philadelphia, PA
- ❑ **Award, Notable Poster**, 1st Annual Research Computing Symposium, UNC-CH 05/2014
- ❑ **Award, Student Membership**, Tropical Medicine Dinner Club of Baltimore 2010 & 2011
- ❑ **Award, Poster**, 3rd Place, 82nd Annual Eastern Branch Meeting 03/2011
Entomological Society of America, Harrisburg, PA
- ❑ **Award, Blue Ribbon Poster**, Johns Hopkins Global Health Day, JHU 02/2011
- ❑ **Award, Global Health Field Research**, JHU Center for Global Health 04/2010
- ❑ **Award, Simpson Student Fund**, Tropical Medicine Dinner Club of Baltimore 03/2010
- ❑ **Deans' Award**, Swarthmore College 05/2005

PREPRINTS AND
WORKING PAPERS

Maurizio PL[†], Fuseini H, Tegha G, Hosseinipour M and De Paris K. Signatures of divergent antimalarial treatment responses in peripheral blood from infants and adults in Malawi. <https://doi.org/10.1101/564757>. (*submitted*, † = corresponding author)

Shorter JR*, **Maurizio PL***, Bell TA, Shaw GD, Miller DR, Gooch TJ, Spence JS, McMillan L, Valdar W and Pardo-Manuel de Villena F. A diallel of the mouse Collaborative Cross founders reveals strong strain-specific maternal effects on litter size. doi: <https://doi.org/10.1101/458877> (*revised and resubmitted*, * = co-first authors) *1 citation (Google Scholar)*

Sanz J, **Maurizio PL**, Snyder-Mackler N, Simons ND, Voyles T, Kohn J, Michopoulos V, Wilson M, Tung J and Barreiro LB. Social history and exposure to pathogen signals modulate social status effects on gene regulation in rhesus macaques. <https://doi.org/10.1101/552356>. (*submitted*)

Keele GR, **Maurizio PL**, Oreper D and Valdar W. Bayesian decision theoretic design of two-founder experimental crosses given diallel data. doi: <https://doi.org/10.1101/489682>. (*working paper*)

PEER-REVIEWED
PUBLICATIONS
(PUBLISHED)

Maurizio PL, Ferris MT, Keele GR, Miller DR, Shaw GD, Whitmore AC, West A, Morrison CR, Noll KE, Plante KS, Cockrell AS, Threadgill DW, Pardo-Manuel de Villena F, Baric RS, Heise MT and Valdar W. **2018**. Bayesian diallel analysis reveals *Mx1*-dependent and *Mx1*-independent effects on response to influenza A virus in mice. *G3: Genes, Genomes, Genetics*. 8(2): 427-445. doi: <https://doi.org/10.1534/g3.117.300438>. PMID:29187420. *5 citations*

Turner SD, **Maurizio PL**, Valdar W, Yandell BS and Simon PW. Dissecting the genetic architecture of shoot growth in carrot (*Daucus carota* L.) using a diallel mating design. **2018**. *G3: Genes, Genomes, Genetics*. 8(2): 411-426. doi: <https://doi.org/10.1534/g3.117.300235>. PMID:29187419. *6 citations*

Espinosa DA, Yadava A, Angov E, **Maurizio PL**, Ockenhouse CF and Zavala F. **2013**. Development of a chimeric *Plasmodium berghei* strain expressing the repeat region of the *P. vivax* circumsporozoite protein for in vivo evaluation of vaccine efficacy. *Infection and Immunity*. 81(8):2882-2887. doi: <https://dx.doi.org/10.1128/IAI.00461-13>. PMID:23716612. *36 citations*

Walsh MC, Kim GK, **Maurizio PL**, Molnar EE and Choi Y. **2008**. TRAF6 auto-ubiquitination-independent activation of the NF κ B and MAPK pathways in response to IL-1 and RANKL. *PLoS One*. 3(12):e4064. doi: <https://dx.doi.org/10.1371/journal.pone.0004064>. PMID:19112497. *145 citations*

MANUSCRIPTS IN
PREPARATION

Maurizio PL, Keele GR, Cai Y, Ferris MT, Miller DR, Whitmore AC, West A, Morrison CR,

Noll KE, Plante KS, Cockrell AS, Pardo-Manuel de Villena F, Baric RS, Heise MT and Valdar W. Influenza lung immunopathology is driven by lymphocyte QTL on Chromosome 2 in Collaborative Cross F1 intercross mice. (*in preparation*)

Maurizio PL, Ferris MT, Linnertz C, Morrison CR, Plante KS, Pardo-Manuel de Villena F, Valdar W and Heise MT. Host parent-of-origin effects modulate influenza virus severity and post-infection gene expression in the lung in F1 reciprocal cross mice. (*in preparation*)

Schaefer A, **Maurizio PL**, Ferris MT, Whitmore AC, Pardo Manuel de Villena F, Threadgill DW, McWeeney S, Heise MT and Baric RS. Host genetic regulation of SARS-CoV infectivity in *in vivo* pathogenesis. (*in preparation*)

ADDITIONAL
PUBLICATIONS

Maurizio PL. 2018. Modeling the Host Genetic Determinants of Influenza Virus Pathogenesis in Mice. Doctor of Philosophy (Ph.D.) Dissertation. University of North Carolina at Chapel Hill. 270 pp. (dissertation, accepted 04/2018)

Maurizio PL and Ferris MT. **2017.** “The Collaborative Cross Resource for Systems Genetics Research of Infectious Diseases.” *Methods in Molecular Biology: Systems Genetics - Methods and Protocols*. Springer/Humana Press: New York, NY. Editors: Klaus Schughart, Robert Williams. eBook ISBN: 978-1-4939-6427-7, Hardcover ISBN: 978-1-4939-6425-3. doi: https://dx.doi.org/10.1007/978-1-4939-6427-7_28. PMID:27933545. (chapter) [4 citations](#)

Maurizio PL. 2011. Detection and vertical transmission of *Culex* flavivirus in *Culex quinquefasciatus* (Diptera: Culicidae) mosquitoes from Zambia, Africa. Master of Science (Sc.M.) thesis. Johns Hopkins University. 127 pp. https://catalog.library.jhu.edu/catalog/bib_4040612. (thesis)

ADDITIONAL
PROFESSIONAL
EXPERIENCE

- Staff Research Associate**, University of California, Los Angeles 10/2007–07/2009
Microbiology, Immunology & Molecular Genetics
Supervisor: M. Carrie Miceli, Ph.D.
Research Areas: drug validation; mouse and tissue culture models of muscular dystrophy
- Research Specialist**, University of Pennsylvania 10/2005–09/2007
Pathology & Laboratory Medicine
Supervisor: Yongwon Choi, Ph.D.
Research Areas: innate immune signaling; ubiquitination and TRAF6 signaling
- Field Assistant**, Ecology, Grand Canyon Trust 05/2005–07/2005
Supervisor: Ethan Aumack, Sc.M.

PROFESSIONAL
DEVELOPMENT

- Participant, GENETICS Peer Review Training Program**, accepted 06/2018–Present
- Participant**, Scientific Writing from the Reader’s Perspective Workshop, UNC-CH 06/2017
- Participant**, Rigor & Reproducibility Workshop, UNC-CH 05/2016
- Student**, Short Course on Systems Genetics, The Jackson Laboratory 09/2014–10/2014
Bar Harbor, ME
- Participant**, Next Generation Sequencing Workshop, UNC-CH 06/2014
- Vaccine Science & Policy Certificate**, Dept. of International Health, JHU 2010

LEADERSHIP &
SERVICE

- Champions Program Mentor** 11/2018–Present
First-Generation, Low-Income, and Immigrant Network (FLI Network), University of Chicago
Service: One-on-one mentorship of undergraduate; helping with career, professional development and goal achievement
- Director, Board of Directors**, elected 10/2015–Present
Universities Allied for Essential Medicines, North America, 501(c)(3)

Service: Corporate Secretary (10/2015–02/2018); Fundraising Committee (2015–2017); Human Resources Committee (2017–Present); promoted responsible management of non-profit; supported national and global university student activities, including access to medicines and human rights initiatives

- Panelist, Carolina Grad Student Firsts**, UNC-CH 01/2018–04/2018
 Service: Volunteered on three speaker panels, representing my experience as a first-generation (1st-gen) undergraduate and graduate student; promoted graduate and doctoral education to UNC-CH and Duke University 1st-gen undergraduates
- Session Chair**, UNC-Chapel Hill Virology Colloquium, Chapel Hill, NC 10/2015
- Session Chair**, Evolution 2014 Conference, Raleigh, NC 06/2014
- Peer Mentor**, First-Year Group 09/2013–12/2013
 Biological & Biomedical Sciences Program, UNC-CH
- Guest Blogger**, 12th Annual World Vaccine Congress & Expo 04/2012
 National Harbor, MD
- HIV Tester & Counselor**, Institute for Human Virology 07/2010–01/2012
 University of Maryland School of Medicine, Baltimore, MD
- Tutor**, Health Professions Recruitment and Exposure Program, JHU 01/2010–03/2010

- TEACHING EXPERIENCE
 Coding Instructor, Introduction to R 06/2016–08/2016
 How to Learn to Code, Small Group, UNC-CH
 Website: <http://mauriziopaul.github.io/intro-to-R/overview/>
- Coding Helper**, Software Carpentry Workshop (Git, SQL), UNC-CH 04/2016
- Teaching Assistant**, Foundations in Population Genomics, BCB 722 03/2014–05/2014
 Instructor: Praveen Sethupathy, Ph.D.; UNC-CH
- Teaching Assistant**, Biological Chemistry Laboratory 01/2004–05/2004
 CHEM 038, Swarthmore College

- SKILLS & TRAVEL
 Programming, Computing & Statistics: R, Python, Mathematica, Matlab, STATA, JAGS, Stan, SQL, LaTeX, bash, git
- Graduate Courses Taken in Quantitative Methods:** Bayesian Statistics, Databases, Mathematical Modeling, Sequence Analysis, Infectious Disease Dynamics, Introduction to Statistical Modeling, Statistical Methods in Public Health, Structural Bioinformatics, Topics in Computer Science: Computational Genetics, Topics in Population Genetics
- Extended Professional Travel:** Macha, Zambia (2010); Hangzhou, China (2007); Australia (2003)

- PROFESSIONAL AFFILIATIONS
 The Genetics Society of America (GSA), Member 2018–Present
- Sigma Xi**, The Scientific Research Society 2004–Present
- AAAS**, Science Program for Excellence in Science, Sponsored Membership 2014–2017