Alison F. Feder

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Pronouns: she/her

Appointments

2021- Assistant Professor, Department of Genome Sciences, *University of Washington*
2018-2021 Miller Fellow, Department of Integrative Biology, *University of California, Berkeley*
Hosts: Oskar Hallatschek & Monty Slatkin

Education

2013-2018 PhD, Biology, Stanford University, Stanford, CA
Advisor: Dmitri Petrov
2012-2013 MSc (by Research), Statistics, University of Oxford, Oxford, UK
Advisor: Gil McVean
2008-2012 BA, Mathematics, *summa cum laude*, University of Pennsylvania, Philadelphia, PA
Advisor: Joshua Plotkin

Research Funding

2018-2021 Miller Fellowship [Website]
2017-2018 Stanford Center for Computational, Evolutionary & Human Genomics Fellowship [Website]
2016-2017 Gerald J. Lieberman Fellowship [Website]
Awarded yearly to twelve Stanford graduate students whose teaching, research and university service demonstrate potential for academic leadership.
2014 Center for Computational, Evolutionary and Human Genomics Trainee Grant
2012-2017 National Science Foundation Graduate Research Fellowship [Website]
2012-2013 Thouron Award [Website]

Awards

2018 Milner Prize in Evolutionary Biology
2018 Samuel Karlin Prize in Mathematical Biology
Awarded to one graduate student in the Stanford Department of Biology per year whose dissertation reaches the highest standard of mathematical biology
2018 James F. Crow Early Career Researcher Finalist (Genetics Society of America)
2017 Gil Omenn Prize for the best article in evolutionary medicine published in the previous year
Awarded for 'More effective drugs lead to harder selective sweeps in the evolution of drug resistance in HIV-1'
2015 Excellence in Teaching Award (Department of Biology, Stanford)
2012 Penn Genome Frontiers Institute Excellence in Genomics Undergraduate Award
2012 Phi Beta Kappa (University of Pennsylvania)

Peer-Reviewed Publications (* denotes equal contributions)

10. **A. F. Feder**, P. S. Pennings, D. A. Petrov (2021). The clarifying role of time series data in the population genetics of HIV. *PLOS Genetics* 17(1): e1009050. [Link]

9. **A. F. Feder**, P. S. Pennings, J. Hermisson*, D. A. Petrov* (2019). Evolutionary dynamics in structured populations under strong population genetic forces. *(G3: GENES, GENOMES, GENETICS)* 9(10):3395-3407. [Link]

   Selected for 2019 G3 Spotlight issue

8. R. S. Mehta, **A. F. Feder**, S. M. Boca, N. A. Rosenberg (2019). The relationship between haplotype-based F_{ST} and haplotype length. *Genetics* 213(1):281-295. [Link]

7. K. Theys*, **A. F. Feder***, M. Gelbart*, M. Hartl, A. Stern, and P. S. Pennings (2018). Within-patient HIV mutation frequencies reveal fitness costs of CpG dinucleotides, drastic amino acid changes and G $\rightarrow$ A mutations. *PLoS Genetics* 14(6): e1007420. [Link]

6. **A. F. Feder**, C. Kline, P. Polacino, M. Cottrell, A. D. Kashuba, B. F. Keele, S.-L. Hu, D. A. Petrov, P. S. Pennings*, and Z. Ambrose* (2017). A spatio-temporal assessment of simian/human immunodeficiency virus (SHIV) evolution reveals a highly dynamic process within the host. *PLoS Pathogens*, 13(5): e1006358. [Link]

5. B. A. Wilson*, N. R. Garud*, **A. F. Feder***, Z. J. Assaf*, and P. S. Pennings (2016). The population genetics of drug resistance evolution in natural populations of viral, bacterial and eukaryotic pathogens. *Molecular Ecology*, 25(1):42–66. [Link]

4. **A. F. Feder**, S.-Y. Rhee, S. P. Holmes, R. W. Shafer, D. A. Petrov*, and P. S. Pennings* (2016). More effective drugs lead to harder selective sweeps in the evolution of drug resistance in HIV-1. *eLife*, 5:e10670. [Link]

3. **A. F. Feder***, S. Kryazhimskiy*, and J. B. Plotkin (2014). Identifying signatures of selection in genetic time series. *Genetics*, 196(2):509–522. [Link]

2. **A. F. Feder**, D. A. Petrov, and A. O. Bergland (2012). LDx: estimation of linkage disequilibrium from high-throughput pooled resequencing data. *PLoS One*, 7(11):e48588. [Link]

1. K. E. Lohmueller, A. Albrechtsen, Y. Li, S. Y. Kim, T. Korneliussen, N. Vinckenbosch, G. Tian, E. Huerta-Sanchez, **A. F. Feder**, N. Grarup, T. Jørgensen, T. Jiang, D. R. Witte, A. Sandbæk, I. Hellmann, T. Lauritzen, T. Hansen, O. Pedersen, J. Wang, R. Nielsen (2011). Natural selection affects multiple aspects of genetic variation at putatively neutral sites across the human genome. *PLoS Genetics*, 7(10):e1002326. [Link]

Pre-prints (* denotes equal contributions)

A. F. Feder, K. Harper, P. S. Pennings, Understanding patterns of HIV multi-drug resistance through models of temporal and spatial drug heterogeneity. *bioRxiv*, 807560. [Link]

Invited Presentations

^{v} *virtually*

2021^{v}	[Scheduled] NIH Laboratory of Viral Diseases, Bethesda, USA
2021^{v}	Temporal Genomics Working Group
2021^{v}	Miller Institute for Basic Research in Science, UC Berkeley, Berkeley, USA
2021^{v}	Quantitative Evolution, Phylogeny and Ecology: from models to data and back, IHP Workshop, Paris, France
2021 ^{v}	Institute of Ecology & Evolution, University of Oregeon, Eugene, USA
2020 ^{v}	Ecology & Evolution Seminar, University of California, Davis, USA
2020	Department of Genome Sciences, University of Washington, Seattle, USA
2019	Department of Ecology & Evolutionary Biology, University of Chicago, Chicago, USA
2019	Department of Computational Biology, Cornell University, Ithaca, USA
2019	Science & Mathematics Seminar, University of Puget Sound, Tacoma, USA
2019	European Society of Evolutionary Biology, Turku, Finland
2019	Society of Molecular Biology & Evolution, Manchester, UK
2019	Trainee Invited Speaker Series, Arjun Raj Lab at Penn, Philadelphia, USA
2019	Science & Technology Seminar, Joint Genome Institute, Walnut Creek, USA
2019	Departmental seminar, University of San Francisco, San Francisco, USA
2018	Palo Alto Research Center, Palo Alto, USA
2018	Milner Prize Lecture, University of Bath, Bath, UK
2018	Systems Biology Seminar, Cancer Research UK Cambridge Institute, UK
2018	Ad hoc seminar, University of California, Davis, USA
2018	Institute for Disease Modeling Annual Symposium, Seattle, USA
2017	Center for Theoretical Evolutionary Genomics, University of California, Berkeley, USA
2017	Institute for Disease Modeling, Bellevue, USA
2017	Center for Inference and Dynamics of Infectious Disease, Fred Hutchinson Cancer Research Institute, Seattle, USA
2017	Omenn Prize talk at the International Society of Evolution, Medicine and Public Health, Groningen, Netherlands
2017	Program for Evolutionary Dynamics, Harvard University, Cambridge, USA
2016	"Darwin's Weekly" Seminar, University of Chicago, Chicago, USA

Contributed/selected presentations

* *talk* † *poster*

2018	[*] Society for Molecular Biology & Evolution, Yokohama, Japan
2018	[*] James F. Crow Award finalist session at PEQG, Madison, USA
2018	[*] HIV Dynamics & Evolution, Leavenworth, USA
2017	[†] Gordon Research Conference: Microbial Population Biology, Andover, USA
2017	[*] Gordon Research Seminar: Microbial Population Biology, Andover, USA
2017	[*] Society for Molecular Biology & Evolution Annual Meeting, Austin, USA
2016	[*] International Society of Evolution, Medicine and Public Health, Raleigh, USA
2016	[*] International HIV Drug Resistance Workshop, Boston, USA
2016	[† †] Conference on Retroviruses and Opportunistic Infections (CROI), Boston, USA
2015	[†] Bio-X Interdisciplinary Initiatives Symposium, Stanford, USA
2015	[*] Society for Molecular Biology & Evolution Annual Meeting, Vienna, Austria
2015	[†]"Forecasting Evolution?" SFB 680 Conference, Lisbon, Portugul

Contributed/Selected Presentations (continued) ** talk* † *poster*

2015 [*] Biomedical Computation at Stanford (BCATS), Stanford, USA
2011 [*] NIMBioS Undergraduate Research Conference at the Interface of Biology and Mathematics, Knoxville, USA
2011 [††] Society for Molecular Biology & Evolution Annual Meeting, Kyoto, Japan

Research Supervision

2021- Elena Romero, Genome Sciences PhD student, *U. Washington*
2020- Will Hannon, Molecular & Cellular Biology PhD student, *Fred Hutch* (J. Bloom lab)
2020- Maya Lewinsohn, MSTP student (Genome Sciences), *U. Washington* (T. Bedford lab)
2020 Helen Sakharova, Comp. Biology PhD rotation student, *UC Berkeley* (O. Hallatschek lab)
2016 Michael Herschl, undergraduate student, *Stanford University* (D. Petrov lab)

Teaching

University:

Fall 2015 Co-teacher for BioCore Exploration (3 hour course), 'Are we still evolving?' with L. Uricchio
Spring 2015 TA for Bio 143, *Evolution*
Spring 2014 TA for Bio 43, *Evolution, Ecology & Plant Biology*

High School:

2016 Guest lecturer, *Evolutionary genomics theory, application and you!*
Stanford Pre-Collegiate Institute
2014-2016 Stanford Splash! Teacher
Taught 6 one-session mini-courses to high school students (two each on mathematical/logical thinking, population genetics and statistics/probability).

Public Outreach

2019 Invited speaker at Nerd Nite East Bay, a general audience seminar series
2017 Finalist in Evolution Film Festival for "Intra-patient Simian-HIV drug resistance evolution: does blood tell the whole story?"
2016 Finalist in Evolution Film Festival for "Better drugs lead to harder sweeps in HIV-1"

Competitive travel support

2018 Young Investigator Travel Award from SMBE (Yokohama, Japan)
2016 International Society for Evolutionary Medicine and Public Health Travel Award (Durham, USA)
2016 CROI Young Investigator Scholarship (Boston, USA)
2015 Wellcome Trust Travel Award (for "Forecasting Evolution?" meeting, Lisbon, Portugal)
2013 Cargese Summer School in Quantitative Genetics Grant (Cargese, France)
2011 NiMBioS Undergraduate Conference Grant (Knoxville, USA)

Academic, Community & University Service

2020-2021	Miller Institute DEI Working Group
2019-2021	Miller Symposium Planning Committee
2018	Co-organizer of SMBE 2018 symposium on 'Intra-host evolutionary dynamics' with K. Xue
2017	Co-organizer of 'Petrino' joint lab retreat between D. Petrov and R. Andino (UCSF) labs
2016-2017	Department of Biology TA Mentorship Program mentor and program organizer
2014-2017	Stanford Bioscience Students Association new student Mentor
2014-2015	Mentored student writing NSF Graduate Research Fellowship application

Referee for American Society of Naturalists, Evolution, Genetics, Journal of Theoretical Biology, Molecular Biology and Evolution, Nature Ecology & Evolution, PCI Evolutionary Biology, PLOS Computational Biology, PLOS Genetics, PNAS