**Education**

Ph.D. Biophysics Harvard University 2014

*The role of deleterious passenger mutations in cancer*, Advisor: [Dr. Leonid A Mirny](mailto:leonid@mit.edu)

B.S. Physics & B.S. BiochemistryUniversity of Rochester 2008

*Magna cum laude*

**Research Experience**

STANFORD UNIVERSITY, DEPARTMENT OF BIOLOGY 2014 – present

Developed a DNA barcode and CRISPR-based method to investigating the fitness landscape and growth heterogeneity of lung adenocarcinoma in vivo with [Dr. Dmitri A. Petrov](mailto:dpetrov@stanford.edu) and [Dr. Monte M. Winslow](mailto:mwinslow@stanford.edu)

HARVARD UNIVERSITY, DEPARTMENT OF BIOPHYSICS 2008 – 2014

Identified passenger mutations as deleterious to cancer cells and a barrier to carcinogenesis using evolutionary theory, genomic analyses, and experimental validation, advised by Dr. Leonid A. Mirny

UC SAN DIEGO, CENTER FOR THEORETICAL BIOLOGICAL PHYSICS 2007

Studied hydrogen-bonding in proteins using molecular dynamics with [Dr. José Onuchic](mailto:jonuchic@rice.edu)

UNIVERSITY OF ROCHESTER, BIOCHEMISTRY AND BIOPHYSICS 2006 – 2008

Determined the cytosolic regions of STE2 in yeast with [Dr. Mark E. Dumont](mailto:Mark_Dumont@urmc.rochester.edu)

WADSWORTH CENTER, MOLECULAR GENETICS 2002 – 2004

Identified the binding partners of STP3 in yeast with [Dr. Randall Morse](mailto:morse@wadsworth.org)

**Awarded Grants**

R01, NCI 2022

**PI**, Tumor-barcoding coupled with high-throughput sequencing for quantitative radiogenomics of the abscopal response in NSCLC*,* $1,000,000 (CA271540)

Case Comprehensive Cancer Center: Pilot Grant 2021

**PI**, *Cancer evolution under androgen receptor inverse agonist,* $80,000

K99/R00, NCI: PATHWAY TO INDEPENDENCE 2018

**PI**, *Quantifying the sources and dynamics of tumor growth variability using Tuba-seq,* $985,260 (CA226506)

VELOSANO, CLEVELAND CLINIC: RACING FOR THE CURE 2018

**Co-Investigator** (PI: [Omar Mian, MD, PhD](mailto:miano@ccf.org)), *Establishing the function of key differentially-expressed genes in small-cell carcinoma of the bladder*, $100,000

R01, NCI: TRANSLATIONAL CLINICAL IGR 2016

**Co-author** (PI: Monte Winslow), *A quantitative multiplexed platform for the pharmacogenomic analysis of lung cancer*, $554,000 (CA207133)

U54, NCI PS-OC: YOUNG-INVESTIGATORS AWARD 2013

**PI**, *Direct visualization of the role of horizontal gene transfer in the evolution of drug resistance in cancer*, $15,000 (CA143874)

U54, NCI PS-OC: TRANS-NETWORK GRANT 2011

**Co-Author** (PI: Leonid Mirny), *Genotypic determinants of metastatic fitness: a delicate balance of passenger and driver mutations*, $400,000 (CA143874)

**Teaching Experience**

STANFORD UNIVERSITY, DEPARTMENT OF BIOLOGY 2016

Mentored Mahi Elango on using machine learning to understand drug vulnerabilities of the Cell Line Encyclopedia, which was an Intel ISEF 2016 Regional Finalist

MASSACHUSSETTES INSTITUTE OF TECHNOLOGY, PRIMES PROGRAM 2011 – 2012

Mentored [Dash Elhauge](http://web.mit.edu/primes/t-Elhauge.shtml) and Michael Zhang, high school students who investigated: *The role cell fusion in cancer development*

HARVARD UNIVERSITY, DEPARTMENT OF CHEMISTRY 2010

Teaching Assistant for both undergraduate- and graduate-level Statistical Thermodynamics, Instructor: [Dr. Eugene Shakhnovich](mailto:eugene@belok.harvard.edu)

JOHNS HOPKINS UNIVERSITY, CENTER FOR TALENTED YOUTH 2008

Teaching Assistant for Nuclear Science (Instructor: [Dr. Yuliya Kuznetsova](mailto:kuznetso@physics.ucsd.edu)) and Fast-Paced High School Physics (Instructor: [Dr. William M. Kallfelz](mailto:kallfelz@cwu.edu))

UNIVERSITY OF ROCHESTER, DEPARTMENT OF BIOLOGY 2008

Teaching Assistant for Introduction to Biochemistry, Instructor: [Dr. Terry Platt](mailto:tpla@mail.rochester.edu)

**Scholarships & Awards**

Scholarship to Physicists working on Cancer, Weizmann Institute of Science 2018

CSBS Postdoctoral Fellowship, Stanford University 2015

CEHG Postdoctoral Fellowship, Stanford University 2014

Scholarship to Emergent Order in Biology, IAS, Cargèse, France 2012

Best Poster Award, NCI Physical Sciences in Oncology Investigator’s Meeting 2011

Phi Beta Kappa, University of Rochester 2008

Martin Tiernan Scholarship, University of Rochester 2005

**Invited Talks**

Max Planck for Evolutionary Biology, Plön Germany 2020

Integrative Genetics and Genomics Graduate Group, UC Davis 2019

Center for Theoretical Evolutionary Genomics Seminar, UC Berkeley 2018

Molecular Biology & Genetics Seminar, Johns Hopkins University 2018

THOR Seminar, Clevland Clinic, Case Western Reserve 2017

**Contributed Talks**

Systems Approaches to Cancer Biology 2018

Stanford Biology Retreat 2018

Gordon Research Seminar, Molecular Mechanisms in Evolution 2017

Stanford Systems Biology of Cancer Annual Symposium 2017

Stanford Cancer Biology Retreat 2015 & 2016

Forecasting Evolution? 2015

CEHG Stanford Symposium 2015

9th European Conference on Mathematical and Theoretical Biology 2014

American Physical Society, March Meeting 2012

Workshop on Mathematical Oncology IV: Integrative Cancer Biology 2012

NCI Physical Sciences in Oncology, Network Investigator's Meeting 2012

Journal of Biomolecular Structure and Dynamics 2011

**Patents**

Winslow, MM; Petrov, DA; **McFarland CD**; Rogers ZN; Winters IP. 2017. *Compositions and Methods for Multiplexed Quantitative Analysis of Cell Lineages*. US Patent Application 62481067, filed April 2017. Patent Pending.

**Other**

[Verified Referee](https://publons.com/author/1458576/christopher-mcfarland) for *Nature Ecology & Evolution, PLoS Computational Biology, PLoS One*, *Genome Biology*, *Genome Biology & Evolution,* *Cancer Research, Physical Biology*, *Journal of Theoretical Biology* & *Journal of Statistical Computation and Simulation*

Developed and maintain several software packages:

[fast\_prng](https://bitbucket.org/cdmcfarland/fast_prng) – Fastest exponential & normal pseudorandom number generator in C

[tuba-seq](https://github.com/petrov-lab/tuba-seq) – Processing & interpretation of ultra-deep DNA barcode sequencing

**Publications**

C Li, WY Lin, H Rizvi, H Cai, **CD McFarland**, ZN Rogers, M Yousefi, IP Winters, CM Rudin, DA Petrov, MM Winslow. (2021) Quantitative in vivo analyses reveal a complex pharmacogenomic landscape in lung adenocarcinoma. *Cancer Res,* [**8:**5472](https://cancerres.aacrjournals.org/content/early/2021/07/02/0008-5472.CAN-21-0716).

H Cai, SK Chew, C Li, MK Tsai, L Andrejka, CW Murray, NW Hughes, EG Shuldiner, EL Ashkin, R Tang, KL Hung, LC Chen, SC Lee, M Yousefi, WY Lin, CA Kunder, L Cong, **CD McFarland**, DA Petrov, C Swanton, MM Winslow. (2021) A Functional Taxonomy of Tumor Suppression in Oncogenic KRAS–Driven Lung Cancer. *Cancer Discov,* [**20:**1325](https://cancerdiscovery.aacrjournals.org/content/11/7/1754.abstract).

S Tilk, C Curtis, DA Petrov, **CD McFarland**. (2019) Most cancers carry a substantial deleterious load due to Hill-Robertson interference. *bioRxiv,* [764340.](https://www.biorxiv.org/content/10.1101/764340v2)

ZN Rogers\*, **CD McFarland**\*, IP Winters, JA Seoane, JJ Brady, S Yoon, C Curtis, DA Petrov, MM Winslow. (2018) Mapping the in vivo fitness landscape of lung adenocarcinoma tumor suppression in mice. *Nat Genetics*, [**50**:483-6](https://www.nature.com/articles/s41588-018-0083-2).

IP Winters, SH Chiou, NK Paulk, **CD McFarland**, PV Lalgudi, RK Ma, L Lisowski, AJ

Connolly, DA Petrov, MA Kay, MM Winslow. (2017) Multiplexed in vivo homology-directed repair and tumor barcoding enables parallel quantification of Kras variant oncogenicity. *Nat Commun,* [**8**:2053-69](https://www.nature.com/articles/s41467-017-01519-y)*.*

ZN Rogers\*, **CD McFarland**\*, IP Winters\*, S Naranjo, CH Chuang, DA Petrov, MM

Winslow. (2017) A quantitative and multiplexed approach to uncover the fitness landscape of tumor suppression in vivo. *Nat Methods,* [**14**:737-42](https://www.nature.com/nmeth/journal/vaop/ncurrent/full/nmeth.4297.html).

“Research Highlight” at *Nat Rev Genetics*, [**18**:456](https://petrov.stanford.edu/pdfs/TubaseqNGcommentary.pdf).

**CD McFarland**\*, JA Yaglom\*, JW Wojtkowiak\*, JG Scott, DL Morse, MY Sherman, LA

Mirny. (2017) The damaging effect of passenger mutations on cancer progression. *Cancer Res*, [**77**:4763-72](http://cancerres.aacrjournals.org/content/early/2017/05/23/0008-5472.CAN-15-3283-T).

BM Grüner, CJ Schulze, D Yang, D Ogasawara, MM Dix, ZN Rogers, C Chuang, **CD**

**McFarland**, S Chiou, JM Brown, BF Cravatt, M Bogyo, MM Winslow. (2016) An in vivo multiplexed small-molecule screening platform. *Nat Methods,* [**13**:883–9](http://www.nature.com/nmeth/journal/v13/n10/full/nmeth.3992.html).

**CD McFarland**. (2015) A modified ziggurat algorithm for generating exponentially and

normally distributed pseudorandom numbers. *J Statist Comput Simulations,* [**86**:1281-94](http://www.tandfonline.com/eprint/VQzbEjcgIJtWcdEdhbam/full).

**CD McFarland**, LA Mirny, KS Korolev. (2014) A tug-of-war between driver and passenger

mutations in cancer and other adaptive processes. *PNAS,* [**111**:15138-43](http://www.pnas.org/content/111/42/15138.full).

“Editor’s Choice” at *Science,* [**306**:597](http://www.sciencemag.org/content/346/6209/597.7.short).

JA Yaglom, **CD McFarland**, LA Mirny, MY Sherman. (2014) Oncogene-triggered

suppression of DNA repair leads to DNA instability in cancer. *Oncotarget*, [**5**:8367-78](http://www.ncbi.nlm.nih.gov/pmc/articles/PMC4226689/).

**CD McFarland**, KS Korolev, GV Kryukov, SR Sunyaev, LA Mirny (2013). Impact of

deleterious passenger mutations on cancer progression. *PNAS*, [**110**:2910-2915](http://www.pnas.org/content/early/2013/02/05/1213968110).

“Research Highlight” at *Nat Rev Cancer*, [**13**:219](http://www.nature.com/nrc/journal/v13/n4/full/nrc3488.html).

\*Contributed Equally