**Education**

Ph.D. Biophysics Harvard University 2014

*The role of deleterious passenger mutations in cancer*

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

*Magna cum laude*

**Affiliations**

CASE WESTERN RESERVE UNIVERSITY 2021 – *present*

Assistant Professor of Genetics & Genome Sciences

Case Comprehensive Cancer Center, Member

Systems Biology & Bioinformatics, Trainer

PAST SCIENTIFIC EMPLOYERS

Dmitri A Petrov, *Postdoctoral Scholar*Monte M Winslow, *Postdoctoral Scholar*Leonid A Mirny, *PhD Student*Mark E Dumont, *Undergraduate Reseracher*Randall Morse, *High School Student*

**Awarded Grants**

R01, NCI 2022

**PI**, Tumor-barcoding coupled with high-throughput sequencing for quantitative radiogenomics of the abscopal response in NSCLC*,* $1,396,660 (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**

CASE WESTERN RESERVE UNIVERSITY 2022 – *present*

GENE 505: Genetics Journal Club, Course Director  
GENE 503: Readings and Discussions in Genetics, Instructor

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**

Grand Rounds, University of Cincinnati 2024

Department of Genetics, University of Georgia 2023

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

**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://www.webofscience.com/wos/author/record/364336) 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*

Guest editor for *Fronteirs in Genetics* Structural Biology of Genetic Mutations

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**

J Maltas, DS Tadele, A Durmaz, **CD McFarland**, M Hinczewski, JG Scott. (2024) Frequency-dependent ecological interactions increase the prevalence, and shape the distribution, of pre-existing drug resistance. *PRX Life,* [In Press](https://journals.aps.org/prxlife/accepted/87071I5dZ501ae05d03d1379847c369a1a911f685).

J Sax, **CD McFarland**, B Carroll. (2024) Limitations of the Commercially Available Gene Expression Test in Predicting Cutaneous Squamous Cell Carcinoma Metastasis and Clinical Outcomes. *J Am Acad Derm,* [**10:**1016](https://www.jaad.org/article/S0190-9622(24)00746-1/abstract).

Y Wang, A Khalil, A Kamar, M Du, T Dinh, **CD McFarland**, Zhenghe Wang. (2023) Unveiling immune checkpoint regulation: exploring the power of in vivo CRISPR screenings in cancer immunotherapy. *Front in Genetics,* [**14:**1304425](https://www.frontiersin.org/journals/genetics/articles/10.3389/fgene.2023.1304425/full).

T Dinh, M Rahm, Z Wang, **CD McFarland**, Athar Khalil. (2023) Exploring the molecular landscape of NNK-induced transformation: A comprehensive genome-wide CRISPR/Cas9 screening. *Genes & Diseases,* [**11:**101131](https://www.sciencedirect.com/science/article/pii/S2352304223004142).

J Tian, **CD McFarland**, J Woodard. (2023) Structural understanding of the functional consequences of missense mutation. *Front in Genetics*, [**14:**1325326](https://www.frontiersin.org/journals/genetics/articles/10.3389/fgene.2023.1325326/full).

S Tilk, S Tkachenko, C Curtis, DA Petrov, **CD McFarland.** (2022) Most cancers carry a substantial deleterious load due to Hill-Robertson interference. *eLife,* [67790](https://elifesciences.org/articles/67790).

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).

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