

# Matthew G. Jones

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## Education

<sup>†</sup> Indicates expected

2017–2021 <sup>†</sup> Ph.D., Bioinformatics, University of California, San Francisco

Thesis Title: Computational tools for single-cell lineage tracing & applications to cancer biology

Research Advisors: Jonathan Weissman and Nir Yosef

2013–2017 B.A., Computer Science, University of California, Berkeley

Research Advisors: Rasmus Nielsen, Russ Corbett-Detig, & Nir Yosef

## Publications

### Journal Articles

- (1) Gong W\*, Granados A\*, Hu J\*, **Jones MG\***, Raz O\*, Salvador-Martinez I\*, Zhang H\*, ..., Meyer P. Benchmarked approaches for cell lineage reconstructions of in vitro dividing cells and in silico models of *Caenorhabditis elegans* and *Mus musculus* developmental trees. *Cell Systems* (In press). 2021.
- (2) Quinn JJQ\*, **Jones MG\***, Okimoto RA, Nanjo S, Chan MM, Yosef N, Bivona TG, Weissman JS. [Single-cell lineages reveal the rates, routes, and drivers of metastasis in cancer xenografts.](#) *Science*. 2021
- (3) **Jones MG\***, Khodaverdian A\*, Quinn JJ\*, Chan MM, Hussmann JA, Wang R, Xu C, Weissman JS, Yosef N. [Inference of Single Cell Phylogenies from Lineage Tracing Data with Cassiopeia.](#) *Genome Biology*. 2020
- (4) Newberry RW, ..., **Jones MG**, ..., DeGrado WF, Kampmann M. [Robust Sequence Determinants of  \$\alpha\$ -Synuclein Toxicity in Yeast Implicate Membrane Binding.](#) *ACS Chem. Biol.*. 2020
- (5) DeTomaso D\*, **Jones MG\***, Subramaniam M, Ashuach T, Ye JC, Yosef N. [Functional Interpretation of Single-Cell Similarity Maps.](#) *Nature Communications*. 2019
- (6) Chan MM\*, Smith ZD\*, Grosswendt S, Kretzmer H, Norman T, Adamson B, Jost M, Quinn JJ, Yang D, **Jones MG**, Khodaverdian A, Yosef N, Meissner A, Weissman JS. [Molecular recording of mammalian embryogenesis.](#) *Nature*. 2019

- (7) Corbett-Detig, R. and **Jones, M.** [SELAM: Simulation of Epistasis and Local adaptation during Admixture with Mate choice](#). *Bioinformatics*. 2016

**\* denotes equal contribution**

## Refereed Workshop Papers

- (1) Ouardini K, Lopez R, **Jones MG**, Prillo S, Zhang R, Jordan MI, Yosef N. [Reconstructing unobserved cellular states from paired single-cell lineage tracing and transcriptomics data](#). ICML Workshop in Computational Biology. 2021.

## Teaching Assistant

2020	CS176: Algorithms for Computational Biology	University of California, Berkeley
2020	BP205B: Dynamical Systems Modeling	University of California, San Francisco
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## Funding, Awards & Honors

2019-2021	UCSF Discovery Fellowship
2020	Allen Institute Cell Lineage Reconstruction DREAM Challenge Best Performer
2019	Quantitative Biology Consortium Retreat Best Poster
2018	Quantitative Biology Consortium Retreat Best Poster

## Presentations

### Invited Talks

2021	Weinberg Lab @ MIT, journal club
2021	Royer Group @ Biohub & Wagner Group @ UCSF, group meeting
2021	AI/ML @ Genentech, seminar
2021	NIH Single Cell Users Group, seminar
2021	Yale Center for Biomedical Data Science, seminar
2020	UC Berkeley Computational Biology Skills Seminar, seminar
2019	UC Berkeley Computational Biology Skills Seminar, seminar

## **Contributed Talks**

- 2021 The Cancer Target Discovery and Development Alliance NIH Site Visit
- 2021 CZI Biohub Seed Networks Computational Biology Meeting
- 2020 Hindsight 2020 - The Allen Institute Developmental Recording Symposium
- 2020 CZI Seed Network 2020 Annual Meeting
- 2020 Center for Genomic Editing and Recording (CGER) Retreat
- 2019 UC Berkeley Computational and Genomic Biology Retreat
- 2019 Quantitative Biology Consortium Retreat

## **Posters**

- 2019 Quantitative Biology Consortium Retreat
- 2019 Chan-Zuckerberg Biohub Confab
- 2019 Next Generation Genomics
- 2018 Quantitative Biology Consortium Retreat

## **Industry Experience**

- 2020 Google Health Genomics, research intern
- 2016 United States Medical Affairs @ Genentech, intern

## **Appointments**

- 2020 Weizmann Institute - Rehovot, Israel (Hosts: Ido Amit & Nir Yosef)

## Professional Activities

2020-2021	UC Berkeley Computational Biology Skills Seminar, coordinator
2019	UC Berkeley Computational Biology Skills Seminar, participant
2018-2019	Northern California Computational Biology Student Symposium, coordinator
2019	UCSF Integrative Program in Quantitative Biology (IPQB) Bootcamp, coordinator
2018	UCSF Integrative Program in Quantitative Biology (IPQB) Bootcamp, Bioinformatics Module Leader

## Supervision

2019-	Richard Zhang, Research Engineer	UC Berkeley
2020-2021	Khalil Ouardini, MSc Student	ENS Cachan, MVA
2019-2021	Yanay Rosen, Undergraduate	UC Berkeley
2019-2021	Robert Wang, Undergraduate	UC Berkeley

## Software

<a href="#">Cassiopeia</a>	Analysis tools for single-cell lineage tracing data.
<a href="#">VISION</a>	Functional annotation of scRNA-seq data.
<a href="#">SELAM</a>	Simulation framework for large-scale population admixture.

## References

Available on request.