

# ANDREW JONES

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## EDUCATION

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| <b>Princeton University</b> • Princeton, NJ | 2019 –      |
| PhD, <i>Computer Science</i>                |             |
| Advisor: Barbara E. Engelhardt              |             |
| <b>Brown University</b> • Providence, RI    | 2016 – 2017 |
| MSc, <i>Computer Science</i>                |             |
| Advisor: Thomas Serre                       |             |
| <b>Brown University</b> • Providence, RI    | 2012 – 2016 |
| BSc, <i>Neuroscience</i>                    |             |

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## RESEARCH

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| <b>Graduate Researcher</b> – Princeton University<br>Princeton, NJ  | 2019 –      |
| <ul style="list-style-type: none"><li>• Currently developing statistical and machine learning tools to analyze high-dimensional biomedical data.</li><li>• Focus on probabilistic models for analysis and alignment of complex data types drawn from multiple modalities.</li><li>• <b>Publications:</b><ul style="list-style-type: none"><li>• CONTRASTIVE POISSON LATENT VARIABLE MODELS: <a href="https://arxiv.org/abs/2102.06731">ARXIV:2102.06731</a></li><li>• PROBABILISTIC CONTRASTIVE PRINCIPAL COMPONENT ANALYSIS: <a href="https://arxiv.org/abs/2012.07977">ARXIV:2012.07977</a></li><li>• NESTED POLICY REINFORCEMENT LEARNING: <a href="https://arxiv.org/abs/2110.02879">ARXIV:2110.02879</a></li></ul></li></ul> |             |
| <b>Associate Computational Biologist</b> – Broad Institute of MIT and Harvard<br>Cambridge, MA  | 2018 – 2019 |
| <ul style="list-style-type: none"><li>• Built statistical tools to study the transcriptional patterns of cancer cells that are targeted by small molecule therapies, resulting in a first-author manuscript and a conference presentation.</li><li>• Other projects included analyzing drug-perturbed single-cell RNAseq data and building a computational tool to align the transcriptomes of cancer cell lines and patient tumors.</li><li>• <b>Publications:</b><ul style="list-style-type: none"><li>• STATISTICAL MODELING OF DRUG RESPONSE IN CANCER CELL LINES: <a href="https://doi.org/10.1101/2020.03.04.976217">DOI:10.1101/2020.03.04.976217</a></li></ul></li></ul>  |             |
| <b>Graduate Research Assistant</b> – Brown University<br>Providence, RI   | 2016 – 2017 |
| <ul style="list-style-type: none"><li>• Developed computer vision models for analyzing the eye gaze patterns of children with Autism Spectrum Disorder, resulting in a Master's Report paper.</li></ul>   |             |
| <b>Undergraduate Research Assistant</b> – Brown University<br>Providence, RI  | 2014 – 2016 |
| <ul style="list-style-type: none"><li>• The BrainGate lab develops brain-computer interfaces (BCIs) for patients with tetraplegia, with the aim of restoring these patients' communication and mobility</li><li>• Created a tool to improve the patients' control of the speed of a computer cursor while using the BCI, and shared my findings in my undergraduate honors thesis.</li></ul>  |             |

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## TEACHING

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| <b>Teaching Assistant</b> – COS424 (Fundamentals of ML), Princeton University      | Spring 2021 |
| <b>Teaching Assistant</b> – COS126 (Intro. Computer Science), Princeton University | Fall 2020   |
| <b>Lead Teaching Assistant</b> – Computational Vision, Brown University            | Fall 2015   |

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## PUBLICATIONS, PREPRINTS, AND ABSTRACTS (\*JOINT AUTHORSHIP)

- **A Jones**, FW Townes, D Li, BE Engelhardt. "Contrastive latent variable modeling with application to case-control sequencing experiments." *The Annals of Applied Statistics* (2021).
- A Mandyam, **A Jones**, K Laudanski, BE Engelhardt. "Nested policy reinforcement learning." [arXiv:2110.02879](https://arxiv.org/abs/2110.02879) (2021).
- Y Cohen-Sharir, et al. "Selective vulnerability of aneuploid human cancer cells to inhibition of the spindle assembly checkpoint." *Nature* (2021).

- C Zirbesa, **A Jones**, K Manzel, N Denburg, and J Barrash. “Assessing the Effects of Healthy and Neuropathological Aging on Personality with the Iowa Scales of Personality Change.” *Developmental Neuropsychology*. (2021).
- D Li\*, **A Jones**\*, BE Engelhardt. “Probabilistic Contrastive Principal Component Analysis.” arXiv:2012.07977 (2020).
- **A Jones**, A Tsherniak, JM McFarland. “Post-perturbational transcriptional signatures of cancer cell line vulnerabilities.” *BioRxiv* (2020).
- JM McFarland, et al. “Multiplexed single-cell transcriptional response profiling to define cancer vulnerabilities and therapeutic mechanism of action.” *Nature Communications* 11.1 (2020): 1-15.
- A Warren, **A Jones**, T Shibue, WC Hahn, JS Boehm, F Vazquez, A Tsherniak, JM McFarland. “Global computational alignment of tumor and cell line transcriptional profiles.” *BioRxiv* (2020).
- **A Jones**, JM McFarland, M Kocak, A Tsherniak. “Predicting small molecule mechanism of action from transcriptional profiles using deep neural networks.” *Deep Learning to Accelerate Drug Discovery* (2018).
- **A Jones**, T Serre. Computational modeling of visual saliency and attention in the Smart Playroom. 2017 Computer Science Master’s Paper (2018).
- DE Warren, MJ Sutterer, J Bruss, TJ Abel, **A Jones**, H Kawasaki, M Voss, M Cassell, MA Howard, D Tranel. “Surgically disconnected temporal pole exhibits resting functional connectivity with remote brain regions.” *bioRxiv* (2017): 127571.
- **A Jones**, D Milstein, L Hochberg, B Jarosiewicz. “Inferring intended speed from curvature as a means to improve decoding in brain-computer interfaces for people with paralysis.” *Neuroscience Honors Thesis* (2016).

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#### AWARDS AND FELLOWSHIPS

Broad Institute Travel Award	2018
Neuroscience Honors, Brown University	2016
Sigma Xi Honor Research Society	2016
Undergraduate Teaching and Research Award	2015

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#### SERVICE

##### Journal reviewing

- Nature Methods
- Nature Biotechnology
- Genome Biology
- Nature Machine Intelligence
- Nature Communications

##### Workshop reviewing

- “Your Model is Wrong: Robustness and misspecification in probabilistic modeling” (NeurIPS 2021)

**Undergraduate Research Mentor** – Princeton University 2020 –

- Primary mentor for two undergraduates pursuing thesis projects.

**Contributing Writer** – Princeton Insights 2020 –

**Research Mentor** – Broad Institute Summer Scholars Program Summer 2018

**Meiklejohn Peer Advisor** – Brown University 2013–2016

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#### TALKS

- *Predicting small molecule mechanism of action from transcription* (2018). Broad Institute/Dana Farber Cancer Program Meeting.
- *TensorFlow Tutorial* (2018) Broad Institute, Cancer Data Science. I organized and led a full-day TensorFlow tutorial and workshop.

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#### OTHER WORK EXPERIENCE

<b>Data Science Intern</b> – AthenaHealth	Summer 2017
<b>Graduate Researcher</b> – Broad Institute of MIT and Harvard	Summer 2016
<b>Undergraduate Researcher</b> – University of Iowa, Dept. of Neurology	Summers 2014, 2015