Benjamin D. Pedigo

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I am a PhD Candidate and NSF Graduate Research Fellow in the Department of Biomedical Engineering at Johns Hopkins University. My research is in the NeuroData lab where I am advised by Dr. Joshua T. Vogelstein and co-advised by Dr. Carey E. Priebe. My work focuses on using statistical and computational techniques to help understand nanoscale connectomes. Currently, I am collaborating with Dr. Marta Zlatic and Dr. Albert Cardona's groups to analyze the first nanoscale connectome of the *Drosophila* larva brain. I also collaborate with Microsoft Research on the development of graspologic, a Python package for statistical analyses of networks.

Education & Training

- 08/18 now **PhD Student**, *Department of Biomedical Engineering*, Johns Hopkins University. **Highlighted courses:** Neuro Data Design, Matrix Theory, Sparse Representations in Machine Learning, Neuroscience and Cognition, Probability and Statistics.
- 09/14 06/18 **Undergraduate Student**, *Department of Bioengineering*, University of Washington. **Highlighted courses:** Neural Coding and Computation, Neural Engineering, Neural Tech Studio, Computational Methods of Data Analysis, Data Structures and Algorithms, High Performance Scientific Computing.

Positions Held Current Position

08/18 – now **PhD Candidate**, *Department of Biomedical Engineering*, Supervised by Dr. Joshua T. Vogelstein and co-supervised by Dr. Carey E. Priebe, Johns Hopkins University. **Research:** Analysis of nanoscale connectomes, network statistics, Python network data science.

Previous Positions

- 05/21 08-20 **Research Intern**, *Societal Resilience*, Supervised by Jonathan Larson, Microsoft Research. **Research**: Organizational communication networks and their relation to reporting structures.
- 06/20 08-20 **Research Intern**, *Team Essex*, Supervised by Weiwei Yang and Dr. Chris White, Microsoft Research.

Research: Network embedding and visualization techniques, network traversal embeddings.

06/17 – 09/17 **Computational Neuroanatomy Intern**, *Neural Coding Group*, Supervised by Dr. Nuno da Costa, Allen Institute for Brain Science.

Research: Nanoscale connectomes, quality control for image alignment, Python development.

- 07/16 06/18 **Undergraduate Researcher**, *Center for Sensorimotor Neural Engineering*, Supervised by Dr. Chet Moritz and Dr. Sarah Mondello, University of Washington. **Research:** Optogenetic spinal cord stimulation after spinal cord injury.
- 07/15 07/16 **Undergraduate Researcher**, *Department of Biology*, Supervised by Dr. Emily Carrington and Dr. Matthew N. George, University of Washington. **Research:** Biomechanical properties of marine mussel attachments.

Awards & Honors

- 2022 Trainee Highlight Award, BRAIN Initiative Meeting.
- 2020 NSF Graduate Research Fellowship.
- 2018 **Summa Cum Laude**, University of Washington. Top 0.5% of graduating class.
- 2017 Levinson Emerging Scholars Award, University of Washington.
- 2017 UW Institute for Neuroengineering Undergraduate Fellowship, University of Washington.

- 2016 **Center for Sensorimotor Neural Engineering Undergraduate Fellowship**, University of Washington.
- 2016 Mary Gates Research Scholarship, University of Washington.
- 2015 Mary Gates Research Scholarship, University of Washington.
- 2014 2018 **Dean's List**, University of Washington.

Publications and Preprints

- † denotes equal contribution
- [1] A. Saad-Eldin, **B. D. Pedigo**, C. E. Priebe, and J. T. Vogelstein. "Graph Matching via Optimal Transport". In: *arXiv preprint arXiv:2111.05366* (2021). URL: https://arxiv.org/abs/2111.05366.
- [2] H. S. Helm, M. Abdin, **B. D. Pedigo**, S. Mahajan, V. Lyzinski, Y. Park, A. Basu, C. M. White, W. Yang, C. E. Priebe, et al. "Leveraging semantically similar queries for ranking via combining representations". In: *arXiv preprint arXiv:2106.12621* (2021). URL: https://arxiv.org/abs/2106.12621.
- [3] J. Chung, E. Bridgeford, J. Arroyo, **B. D. Pedigo**, A. Saad-Eldin, V. Gopalakrishnan, L. Xiang, C. E. Priebe, and J. T. Vogelstein. "Statistical Connectomics". In: *Annual Review of Statistics and Its Application* 8 (2021), pp. 463–492. URL: https://www.annualreviews.org/doi/abs/10.1146/annurev-statistics-042720-023234.
- [4] V. Gopalakrishnan, J. Chung, E. Bridgeford, **B. D. Pedigo**, J. Arroyo, L. Upchurch, G. A. Johnson, N. Wang, Y. Park, C. E. Priebe, and J. T. Vogelstein. *Multiscale Comparative Connectomics*. 2020. URL: https://arxiv.org/abs/2011.14990.
- [5] A. S. Charles, B. Falk, N. Turner, T. D. Pereira, D. Tward, **B. D. Pedigo**, J. Chung, R. Burns, S. S. Ghosh, J. M. Kebschull, et al. "Toward Community-Driven Big Open Brain Science: Open Big Data and Tools for Structure, Function, and Genetics". In: *Annual Review of Neuroscience* 43 (2020). URL: https://www.annualreviews.org/doi/abs/10.1146/annurev-neuro-100119-110036.
- [6] T. L. Athey, T. Liu, **B. D. Pedigo**, and J. T. Vogelstein. "AutoGMM: Automatic and Hierarchical Gaussian Mixture Modeling in Python". In: *arXiv preprint arXiv:1909.02688* (2019). URL: https://arxiv.org/abs/1909.02688.
- [7] J. Chung[†], **B. D. Pedigo**[†], E. W. Bridgeford, B. K. Varjavand, H. S. Helm, and J. T. Vogelstein. "GraSPy: Graph Statistics in Python." In: *Journal of Machine Learning Research* 20.158 (2019), pp. 1–7. URL: https://arxiv.org/abs/1904.05329.
- [8] J. T. Vogelstein, E. W. Bridgeford, **B. D. Pedigo**, J. Chung, K. Levin, B. Mensh, and C. E. Priebe. "Connectal coding: discovering the structures linking cognitive phenotypes to individual histories". In: *Current opinion in neurobiology* 55 (2019), pp. 199–212. URL: https://www.sciencedirect.com/science/article/pii/S0959438818301430.
- [9] M. N. George, **B. D. Pedigo**, and E. Carrington. "Hypoxia weakens mussel attachment by interrupting DOPA cross-linking during adhesive plaque curing". In: *Journal of The Royal Society Interface* 15.147 (2018), p. 20180489. URL: https://royalsocietypublishing.org/doi/full/10.1098/rsif.2018.0489.

Talks

- [1] **B. D. Pedigo**. "Hypothesis testing for connectome comparisons: a statistical analysis of bilateral symmetry in an insect brain connectome". In: Drexel University Biomedical Engineering and Neuroscience, Invited Seminar, Mar. 2022.
- [2] **B. D. Pedigo**, M. Winding, M. Zlatic, A. Cardona, C. E. Priebe, and J. T. Vogelstein. "Maggot brain, mirror image? A statistical analysis of bilateral symmetry in an insect brain connectome". In: Neuromatch 4.0, Selected Talk, Dec. 2021.
- [3] **B. D. Pedigo** and J. T. Vogelstein. "graspologic: A python package for rigorous statistical analysis of populations of attributed connectomes". In: BRAIN Informatics Webinar, Oct. 2021.

- [4] **B. D. Pedigo.** "Network data science for bilateral brains: Applications in the larval Drosophila connectome". In: Brain Connectivity Workshop 5, Invited Talk, Mar. 2021.
- [5] **B. D. Pedigo**, M. Winding, A. Saad-Eldin, T. Liu, A. Cardona, M. Zlatic, C. E. Priebe, and J. T. Vogelstein. "Statistical tools for nanoscale connectomics: clustering neurons in the Drosophila larva brain and other applications". In: Neuromatch 3.0, Oct. 2020.
- [6] A. Saad-Eldin, B. D. Pedigo, Y. Park, C. E. Priebe, and J. T. Vogelstein. "NeuroGraphMatch". In: Neuromatch 3.0, Oct. 2020.
- [7] T. Liu, **B. D. Pedigo**, T. L. Athey, and J. T. Vogelstein. "Hierarchical stochastic block modeling in the Drosophila connectome". In: Neuromatch 3.0, Oct. 2020.
- [8] V. Gopalakrishnan, J. Chung, E. Bridgeford, J. Arroyo, **B. D. Pedigo**, C. E. Priebe, and J. T. Vogelstein. "Statistical Methods for Multiscale Comparative Connectomics". In: Neuromatch 3.0, Oct. 2020.

Poster Presentations

- [1] **B. D. Pedigo**, M. Powell, E. W. Bridgeford, M. Winding, C. E. Priebe, and J. T. Vogelstein. "Generative network modeling reveals a first quantitative definition of bilateral symmetry exhibited by a whole insect brain connectome". In: From Neuroscience to Artificially Intelligent Systems (NAISys), Cold Spring Harbor Laboratory, NY, USA, Mar. 2022.
- [2] **B. D. Pedigo**, M. Winding, M. Zlatic, A. Cardona, C. E. Priebe, and J. T. Vogelstein. "A quantitative comparison of a complete connectome to artificial intelligence architectures". In: From Neuroscience to Artificially Intelligent Systems (NAISys), Cold Spring Harbor Laboratory, NY, USA, Nov. 2020.
- [3] **B. D. Pedigo**, J. Chung, E. W. Bridgeford, B. Varjavand, C. E. Priebe, and J. T. Vogelstein. "GraSPy: an Open Source Python Package for Statistical Connectomics". In: Max Planck/HHMI Connectomics Meeting, Berlin, Germany, Apr. 2019.
- [4] **B. D. Pedigo**, S. E. Mondello, A. E. Fischedick, and C. T. Moritz. "Optimization of optogenetic stimulation for spinal cord injury rehabilitation". In: UW Undergraduate Research Symposium, Seattle, WA, USA, May 2018.
- [5] **B. D. Pedigo**, S. E. Mondello, A. E. Fischedick, and C. T. Moritz. "Investigation of optogenetic-induced damage to the rat spinal cord". In: Center for Sensorimotor Neural Engineering Summer Symposium, Seattle, WA, USA, Aug. 2017.
- [6] **B. D. Pedigo**, S. E. Mondello, A. E. Fischedick, and C. T. Moritz. "Optimization of optogenetic spinal cord stimulation". In: UW Undergraduate Research Symposium, Seattle, WA, USA, May 2017.
- [7] **B. D. Pedigo**, M. N. George, and E. Carrington. "Effects of environmental factors on Mytilus mussel adhesion". In: UW Undergraduate Research Symposium, Seattle, WA, USA, May 2016.

Software

graspologic, github.com/microsoft/graspologic, Co-lead developer/maintainer.

A Python package for statistical analysis of network data, co-developed by NeuroData lab and Microsoft Research. Used in production at Microsoft and almost 100k downloads. Formerly known as GraSPy.

Mentoring

- 06/21 now **Diane Lee**, *Undergraduate Researcher*, BME, JHU.
 - Led development of new visualization tools into graspologic. Currently researching methods for quantitative network comparison.
- 08/21 now **Kareef Ullah**, *Undergraduate Researcher*, BME, JHU.

 Ensured statistical tests in graspologic were reproducible. Currently implementing network configuration models in graspologic.
- 01/20 12/21 **Ali Saad-Eldin**, *Master's Researcher*, BME, JHU.

 Led integration of graph matching tools into SciPy. Developed a new graph matching technique and wrote a manuscript on it. Currently at Amazon.
- 06/19 08/19 Kareef Ullah, High School Summer Intern, BME, JHU.

06/19 - 07/19 Kiki Zhang, High School Summer Intern, BME, JHU.

Teaching

Intersession Network Data Science, EN.580.129(13), Instructor and course creator.

2022 Johns Hopkins University

Fall/Spring NeuroData Design I & II, EN.580.237/437/637, Team Lead.

2020/2021 Johns Hopkins University

Fall/Spring NeuroData Design I & II, EN.580.237/437/637, TA.

2019/2020 Johns Hopkins University

Spring 2018 Biomedical Signals and Sensors, BIOEN 316, TA.

University of Washington

Spring 2017 Biomedical Signals and Sensors, BIOEN 316, TA.

University of Washington

Service and Organizations

2022 - now Life Sciences Advisor, A-Level Capital.

2020 - now **Mentor**, *BME Application Assistance Program*, Johns Hopkins University. Reviewed application materials for applicants from underrepresented backgrounds

 $2017 - 2018 \quad \textbf{President and Founder}, \textit{Synaptech}, \textbf{University of Washington}.$

Student organization for undergraduates in neural engineering

2017 - 2018 **Undergraduate Representative**, *Center for Sensorimotor Neural Engineering*, University of Washington.

2017 - 2018 **President**, Bird Club, University of Washington.

2016 - 2017 **Treasurer**, Bird Club, University of Washington.

2017 - 2018 **Mentor**, *BioExpo*, Northwest Association for Biomedical Research.

Languages and Tools

Proficient English, Python (NumPy, Pandas, Scikit-learn, SciPy, Seaborn, Matplotlib, NetworkX, Jupyter Notebooks, JupyterBook), Git, MATLAB, MTpX.

Inproficient R, C++, Java, Blender, HTML.