Joshua T. Vogelstein

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Demographic & Personal Information

Current Appointments

09/19 – now	**Joint Appointment**, *Department of Biostatistics*, Johns Hopkins University (JHU).
08/15 – now	**Joint Appointment**, *Department of Applied Mathematics and Statistics*.
08/14 – now	**Assistant Professor**, *Department of Biomedical Engineering*, Johns Hopkins University (JHU).
08/14 – now	**Core Faculty**, *Institute for Computational Medicine (ICM)*.
08/14 – now	**Core Faculty**, *Center for Imaging Science (CIS)*.
08/14 – now	**Joint Appointment**, *Department of Neuroscience*.
08/14 – now	**Joint Appointment**, *Department of Computer Science*.
08/14 – now	**Assistant Research Faculty**, *Human Language Technology Center of Excellence*.
10/12 – now	**Affiliated Faculty**, *Institute for Data Intensive Engineering and Sciences*.

Education & Training

2009	**Ph.D in Neuroscience**, *Johns Hopkins School of Medicine*, Advisor: Eric Young, **Thesis:** OOPSI: a family of optical spike inference algorithms for inferring neural connectivity from population calcium imaging .
2009	**M.S. in Applied Mathematics & Statistics**, *Johns Hopkins University*.
2002	**B.A. in Biomedical Engineering**, *Washington University, St. Louis*.

Academic Experience

08/18 – now	**Director of Biomedical Data Science Focus Area**.
05/16 – now	**Visiting Scientist**, *Howard Hughes Medical Institute*, Janelia Research Campus.
10/12 – 08/14	**Endeavor Scientist**, *Child Mind Institute*.
08/12 – 08/14	**Affiliated Faculty**, *Kenan Institute for Ethics*. Duke University
08/12 – 08/14	**Adjunct Faculty**, *Department of Computer Science*.
12/09 – 01/11	**Post-Doctoral Fellow**, *Department of Applied Mathematics and Statistics*, Supervised by Carey E. Priebe, Johns Hopkins University. **Research** Statistics of populations of networks.
06/01 – 09/01	**Research Assistant**, *Prof. Randy O'Reilly, Dept. of Psychology*. University of Colorado
06/00 – 09/00	**Clinical Engineer**, *Johns Hopkins Hospital*.
06/99 – 08/99	**Research Assistant under Dr. Jeffrey Williams**, *Dept. of Neurosurgery, Johns Hopkins Hospital*.
06/98 – 08/98	**Research Assistant under Professor Kathy Cho**, *Dept. of Pathology, Johns Hopkins School of Medicine*.

▬▬ Publications

Published Peer-Reviewed Research Articles

Note: CV author in bold; Trainees in italics,
(55 papers; top 10 cited 3,128 times; H-index 30) as of 2019/12/19

[55]   *Shangsi Wang, Jesús Arroyo*, **Joshua T Vogelstein**, and Carey E Priebe. "Joint Embedding of Graphs". In: Transactions on Pattern Analysis and Machine Intelligence in press (Oct. 2019). URL: http://arxiv.org/abs/1703.03862.

[54]   *Youjin Lee, Cencheng Shen*, Carey E Priebe, and **Joshua T Vogelstein**. "Network dependence testing via diffusion maps and distance-based correlations". In: Biometrika (Sept. 2019). ISSN: 0006-3444. DOI: 10.1093/biomet/asz045. arXiv: 1703.10136. URL: https://doi.org/10.1093/biomet/asz045.

[53]   *Jaewon Chung, Benjamin D. Pedigo, Eric W. Bridgeford, Bijan K. Varjavand*, and **Joshua T. Vogelstein**. "GraSPy: Graph Statistics in Python". In: Journal of Machine Learning Research 20.158 (Apr. 2019), pp. 1–7. eprint: https://arxiv.org/abs/1904.05329. URL: http://jmlr.org/papers/v20/19-490.html.

[52]   **Joshua T. Vogelstein**, *Eric W. Bridgeford, Benjamin D. Pedigo, Jaewon Chung*, Keith Levin, Brett Mensh, and Carey E. Priebe. "Connectal Coding: Discovering the Structures Linking Cognitive Phenotypes to Individual Histories". In: Current Opinion in Neurobiology 55 (Apr. 2019), pp. 199–212. ISSN: 18736882. DOI: 10.1016/j.conb.2019.04.005. URL: https://doi.org/10.1016/j.conb.2019.04.005.

[51]   Jake J. Son, Jon C. Clucas, Curt White, Anirudh Krishnakumar, **Joshua T. Vogelstein**, Michael P. Milham, and Arno Klein. "Thermal sensors improve wrist-worn position tracking". In: npj digital medicine 2.1 (Feb. 2019). ISSN: 2398-6352. DOI: 10.1038/s41746-019-0092-2. URL: https://doi.org/10.1038/s41746-019-0092-2.

[50]   Carey E. Priebe, Youngser Park, **Joshua T. Vogelstein**, John M. Conroy, Vince Lyzinski, Minh Tang, Avanti Athreya, Joshua Cape, and *Eric Bridgeford*. "On a two-truths phenomenon in spectral graph clustering". In: Proceedings of the National Academy of Sciences of the United States of America 116.13 (Feb. 2019), pp. 5995–6000. ISSN: 10916490. DOI: 10.1073/pnas.1814462116. arXiv: 1808.07801. URL: https://www.pnas.org/content/early/2019/03/07/1814462116.short.

[49]   **Joshua T. Vogelstein**, *Eric W. Bridgeford*, Qing Wang, Carey E. Priebe, Mauro Maggioni, and *Cencheng Shen*. "Discovering and deciphering relationships across disparate data modalities". In: eLife 8 (Jan. 2019). ISSN: 2050084X. DOI: 10.7554/eLife.41690. arXiv: 1609.05148. URL: https://elifesciences.org/articles/41690.

[48]   Runze Tang, Michael Ketcha, Alexandra Badea, Evan D Calabrese, Daniel S Margulies, **Joshua T Vogelstein**, Carey E Priebe, and Daniel L Sussman. "Connectome Smoothing via Low-rank Approximations". In: Transactions in Medical Imaging (Dec. 2018). URL: https://ieeexplore.ieee.org/document/8570772.

[47]   *Cencheng Shen*, Carey E Priebe, and **Joshua T Vogelstein**. "From Distance Correlation to Multiscale Graph Correlation". In: Journal of the American Statistical Association (Oct. 2018). URL: https://www.tandfonline.com/doi/full/10.1080/01621459.2018.1543125.

[46]   **Joshua T. Vogelstein**, Eric Perlman, Benjamin Falk, *Alex Baden*, William Gray Roncal, *Vikram Chandrashekhar*, Forrest Collman, Sharmishtaa Seshamani, Jesse L. Patsolic, *Kunal Lillaney*, Michael Kazhdan, Robert Hider, Derek Pryor, Jordan Matelsky, Timothy Gion, *Priya Manavalan*, Brock Wester, Mark Chevillet, Eric T. Trautman, Khaled Khairy, *Eric Bridgeford*, Dean M. Kleissas, Daniel J. Tward, Ailey K. Crow, Brian Hsueh, Matthew A. Wright, Michael I. Miller, Stephen J. Smith, R. Jacob Vogelstein, Karl Deisseroth, and Randal Burns. "A Community-Developed Open-Source Computational Ecosystem for Big Neuro Data". In: Nature Methods 15.11 (Oct. 2018), pp. 846–847. ISSN: 15487105. DOI: 10.1038/s41592-018-0181-1. arXiv: 1804.02835. URL: https://www.nature.com/articles/s41592-018-0181-1.

[45]   Avanti Athreya, Donniell E. Fishkind, Minh Tang, Carey E. Priebe, Youngser Park, **Joshua T. Vogelstein**, Keith Levin, Vince Lyzinski, Yichen Qin, and Daniel L Sussman. "Statistical Inference on Random Dot Product Graphs: a Survey". In: Journal of Machine Learning Research 18 (May 2018), pp. 1–92. ISSN: 15337928. arXiv: 1709.05454. URL: http://jmlr.org/papers/v18/17-448.html.

[44]   Joshua D. Cohen, Lu Li, Yuxuan Wang, Christopher Thoburn, Bahman Afsari, Ludmila Danilova, Christopher Douville, Ammar A. Javed, Fay Wong, Austin Mattox, Ralph H. Hruban, Christopher L. Wolfgang, Michael G. Goggins, Marco Dal Molin, Tian Li Wang, Richard Roden, Alison P. Klein, Janine Ptak, Lisa Dobbyn, Joy Schaefer, Natalie Silliman, Maria Popoli, **Joshua T. Vogelstein**, *James D. Browne*, Robert E. Schoen, Randall E. Brand, Jeanne Tie, Peter Gibbs, Hui Li Wong, Aaron S. Mansfield, Jin Jen, Samir M. Hanash, Massimo Falconi, Peter J. Allen, Shibin Zhou, Chetan Bettegowda, Luis A. Diaz, Cristian Tomasetti, Kenneth W. Kinzler, Bert Vogelstein, Anne Marie Lennon, and Nickolas Papadopoulos. "Detection and localization of surgically resectable cancers with a multi-analyte blood test". In: Science 359.6378 (Feb. 2018), pp. 926–930. ISSN: 10959203. DOI: 10.1126/science.aar3247. URL: https://doi.org/10.1126/science.aar3247.

[43]   Daniele Durante, David B. Dunson, and **Joshua T. Vogelstein**. "Rejoinder: Nonparametric Bayes Modeling of Populations of Networks". In: Journal of the American Statistical Association 112 (Oct. 2017). ISSN: 0162-1459. DOI: 10.1080/01621459.2017.1395643. URL: https://doi.org/10.1080/01621459.2017.1395643.

[42]   *Gregory Kiar*, Krzysztof J. Gorgolewski, Dean Kleissas, William Gray Roncal, Brian Litt, Brian Wandell, Russel A. Poldrack, Martin Wiener, R. Jacob Vogelstein, Randal Burns, and **Joshua T. Vogelstein**. "Science in the cloud (SIC): A use case in MRI connectomics". In: GigaScience 6.5 (May 2017), pp. 1–10. ISSN: 2047-217X. DOI: 10.1093/gigascience/gix013. arXiv: 1610.08484. URL: https://academic.oup.com/gigascience/article-lookup/doi/10.1093/gigascience/gix013.

[41]   *Shaojie Chen*, Kai Liu, Yuguang Yang, Yuting Xu, Seonjoo Lee, Martin Lindquist, Brian S. Caffo, and **Joshua T. Vogelstein**. "An M-estimator for reduced-rank system identification". In: Pattern Recognition Letters 86 (Jan. 2017), pp. 76–81. ISSN: 0167-8655. DOI: 10.1016/J.PATREC.2016.12.012. URL: https://www.sciencedirect.com/science/article/pii/S0167865516303671.

[40]   Anish K. Simhal, Cecilia Aguerrebere, Forrest Collman, **Joshua T. Vogelstein**, Kristina D. Micheva, Richard J. Weinberg, Stephen J. Smith, and Guillermo Sapiro. "Probabilistic fluorescence-based synapse detection". In: PLoS Computational Biology 13.4 (2017). DOI: 10.1371/journal.pcbi.1005493. URL: https://doi.org/10.1371/journal.pcbi.1005493.

[39]   *Da Zheng, Disa Mhembere*, Vince Lyzinski, **Joshua T. Vogelstein**, Carey E. Priebe, and Randal Burns. "Semi-external memory sparse matrix multiplication for billion-node graphs". In: IEEE Transactions on Parallel and Distributed Systems 28.5 (2017), pp. 1470–1483. ISSN: 10459219. DOI: 10.1109/TPDS.2016.2618791. arXiv: 1602.02864. URL: https://ieeexplore.ieee.org/abstract/document/7593270.

[38]   *Cencheng Shen*, **Joshua T. Vogelstein**, and Carey E. Priebe. "Manifold matching using shortest-path distance and joint neighborhood selection". In: Pattern Recognition Letters 92 (2017), pp. 41–48. ISSN: 01678655. DOI: 10.1016/j.patrec.2017.04.005. arXiv: 1412.4098. URL: http://www.sciencedirect.com/science/article/pii/S016786551730106X.

[37]   Norbert Binkiewicz, **Joshua T. Vogelstein**, and Karl Rohe. "Covariate-assisted spectral clustering". In: Biometrika 104.2 (2017), pp. 361–377. ISSN: 14643510. DOI: 10.1093/biomet/asx008. arXiv: 1411.2158. URL: https://doi.org/10.1093/biomet/asx008.

[36]   Daniele Durante, David B. Dunson, and **Joshua T. Vogelstein**. "Nonparametric Bayes Modeling of Populations of Networks". In: Journal of the American Statistical Association 112.520 (2017), pp. 1516–1530. ISSN: 1537274X. DOI: 10.1080/01621459.2016.1219260. arXiv: 1406.7851. URL: https://doi.org/10.1080/01621459.2016.1219260.

[35]   Qing Wang, Ming Zhang, *Tyler Tomita*, **Joshua T. Vogelstein**, Shibin Zhou, Nickolas Papadopoulos, Kenneth W. Kinzler, and Bert Vogelstein. "Selected reaction monitoring approach for validating peptide biomarkers". In: Proceedings of the National Academy of Sciences of the United States of America 114.51 (2017), pp. 13519–13524. ISSN: 10916490. DOI: 10.1073/pnas.1712731114. URL: http://www.pnas.org/content/114/51/13519.short.

[34]   David Grant Colburn Hildebrand, Marcelo Cicconet, Russel Miguel Torres, Woohyuk Choi, Tran Minh Quan, Jungmin Moon, Arthur Willis Wetzel, Andrew Scott Champion, Brett Jesse Graham, Owen Randlett, George Scott Plummer, Ruben Portugues, Isaac Henry Bianco, Stephan Saalfeld, *Alexander David Baden*, *Kunal Lillaney*, Randal Burns, **Joshua Tzvi Vogelstein**, Alexander Franz Schier, Wei Chung Allen Lee, Won Ki Jeong, Jeff William Lichtman, and Florian Engert. "Whole-brain serial-section electron microscopy in larval zebrafish". In: Nature 545.7654 (2017), pp. 345–349. ISSN: 14764687. DOI: 10.1038/nature22356. URL: https://doi.org/10.1038/nature22356.

[33]   Danai Koutra, Neil Shah, **Joshua T. Vogelstein**, Brian Gallagher, and Christos Faloutsos. "DeltaCon: Principled Massive-Graph Similarity Function with Attribution". In: ACM Transactions on Knowledge Discovery from Data 10.3 (Feb. 2016). ISSN: 1556-4681. DOI: 10.1145/2824443. URL: http://doi.acm.org/10.1145/2824443.

[32]   Vince Lyzinski, Donniell E. Fishkind, Marcelo Fiori, **Joshua T. Vogelstein**, Carey E. Priebe, and Guillermo Sapiro. "Graph Matching: Relax at Your Own Risk". In: IEEE Transactions on Pattern Analysis and Machine Intelligence 38.1 (Jan. 2016), pp. 60–73. ISSN: 01628828. DOI: 10.1109/TPAMI.2015.2424894. arXiv: 1405.3133. URL: http://doi.org/10.1109/TPAMI.2015.2424894.

[31]   Eva L. Dyer, *William Gray Roncal*, Hugo L. Fernandes, Doga Gürsoy, Vincent De Andrade, Rafael Vescovi, Kamel Fezzaa, Xianghui Xiao, **Joshua T. Vogelstein**, Chris Jacobsen, Konrad P. Körding, and Narayanan Kasthuri. "Quantifying Mesoscale Neuroanatomy Using X-Ray Microtomography". In: eNeuro 4 (2016). ISSN: 2373-2822. DOI: 10.1523/ENEURO.0195-17.2017. eprint: 1604.03629. URL: https://doi.org/10.1523/ENEURO.0195-17.2017.

[30]   Raag D. Airan, **Joshua T. Vogelstein**, Jay J. Pillai, Brian Caffo, James J. Pekar, and Haris I. Sair. "Factors affecting characterization and localization of interindividual differences in functional connectivity using MRI". In: Human Brain Mapping 37.5 (2016), pp. 1986–1997. ISSN: 10970193. DOI: 10.1002/hbm.23150. URL: http://dx.doi.org/10.1002/hbm.23150.

[29]   Li Chen, Cencheng Shen, **Joshua T. Vogelstein**, and Carey E. Priebe. "Robust Vertex Classification". In: IEEE Transactions on Pattern Analysis and Machine Intelligence 38.3 (2016), pp. 578–590. ISSN: 01628828. DOI: 10.1109/TPAMI.2015.2456913. URL: http://dx.doi.org/10.1109/TPAMI.2015.2456913.

[28]   Carey E. Priebe, Daniel L. Sussman, Minh Tang, and **Joshua T. Vogelstein**. "Statistical Inference on Errorfully Observed Graphs". In: Journal of Computational and Graphical Statistics 24.4 (Oct. 2015), pp. 930–953. ISSN: 15372715. DOI: 10.1080/10618600.2014.951049. arXiv: 1211.3601. URL: https://doi.org/10.1080/10618600.2014.951049.

[27]   Kristen M. Harris, Josef Spacek, Maria Elizabeth Bell, Patrick H. Parker, Laurence F. Lindsey, Alexander D. Baden, **Joshua T. Vogelstein**, and Randal Burns. "A resource from 3D electron microscopy of hippocampal neuropil for user training and tool development". In: Scientific Data 2 (2015). ISSN: 20524463. DOI: 10.1038/sdata.2015.46. URL: https://doi.org/10.1038/sdata.2015.46.

[26]   Li Chen, **Joshua T Vogelstein**, Vince Lyzinski, and Carey E Priebe. "A Joint Graph Inference Case Study: the C.elegans Chemical and Electrical Connectomes". In: Worm 5 (2015). ISSN: 2162-4054. DOI: 10.1080/21624054.2016.1142041. eprint: 1507.08376. URL: http://arxiv.org/abs/1507.08376.

[25]   *William R Gray Roncal*, Dean M Kleissas, **Joshua T Vogelstein**, *Priya Manavalan*, *Kunal Lillaney*, Michael Pekala, Randal Burns, R Jacob Vogelstein, Carey E Priebe, Mark A Chevillet, and Gregory D Hager. "An automated images-to-graphs framework for high resolution connectomics". In: Frontiers in Neuroinformatics 9 (2015). ISSN: 1662-5196. DOI: 10.3389/fninf.2015.00020. URL: http://journal.frontiersin.org/article/10.3389/fninf.2015.00020.

[24]   **Joshua T. Vogelstein**, John M. Conroy, Vince Lyzinski, Louis J. Podrazik, Steven G. Kratzer, Eric T. Harley, Donniell E. Fishkind, R. Jacob Vogelstein, and Carey E. Priebe. "Fast Approximate Quadratic programming for graph matching". In: <u>PLoS ONE</u> 10.4 (2015). ISSN: 19326203. DOI: 10.1371/journal.pone.0121002. URL: http://dx.doi.org/10.1371/journal.pone.0121002.

[23]   **Joshua T. Vogelstein** and Carey E. Priebe. "Shuffled Graph Classification: Theory and Connectome Applications". In: <u>Journal of Classification</u> 32.1 (2015), pp. 3–20. ISSN: 14321343. DOI: 10.1007/s00357-015-9170-6. arXiv: 1112.5506. URL: https://doi.org/10.1007/s00357-015-9170-6.

[22]   Vince Lyzinski, Daniel L. Sussman, Donniell E. Fishkind, Henry Pao, Li Chen, **Joshua T. Vogelstein**, Youngser Park, and Carey E. Priebe. "Spectral clustering for divide-and-conquer graph matching". In: <u>Parallel Computing</u> 47 (2015), pp. 70–87. ISSN: 01678191. DOI: 10.1016/j.parco.2015.03.004. arXiv: 1310.1297. URL: https://doi.org/10.1016/j.parco.2015.03.004.

[21]   Narayanan Kasthuri, Kenneth Jeffrey Hayworth, Daniel Raimund Berger, Richard Lee Schalek, José Angel Conchello, Seymour Knowles-Barley, Dongil Lee, Amelio Vázquez-Reina, Verena Kaynig, Thouis Raymond Jones, Mike Roberts, Josh Lyskowski Morgan, Juan Carlos Tapia, H. Sebastian Seung, William Gray Roncal, **Joshua Tzvi Vogelstein**, Randal Burns, Daniel Lewis Sussman, Carey Eldin Priebe, Hanspeter Pfister, and Jeff William Lichtman. "Saturated Reconstruction of a Volume of Neocortex". In: <u>Cell</u> 162.3 (2015), pp. 648–661. ISSN: 10974172. DOI: 10.1016/j.cell.2015.06.054. URL: https://doi.org/10.1016/j.cell.2015.06.054.

[20]   David E. Carlson, **Joshua T. Vogelstein**, Qisong Wu, Wenzhao Lian, Mingyuan Zhou, Colin R. Stoetzner, Daryl Kipke, Douglas Weber, David B. Dunson, and Lawrence Carin. "Multichannel Electrophysiological Spike Sorting via Joint Dictionary Learning and Mixture Modeling". In: <u>IEEE Transactions on Biomedical Engineering</u> 61.1 (Jan. 2014), pp. 41–54. ISSN: 0018-9294. DOI: 10.1109/TBME.2013.2275751. arXiv: 1304.0542. URL: http://ieeexplore.ieee.org/document/6571240/.

[19]   Nicholas C. Weiler, Forrest Collman, **Joshua T. Vogelstein**, Randal Burns, and Stephen J. Smith. "Synaptic molecular imaging in spared and deprived columns of mouse barrel cortex with array tomography". In: <u>Scientific Data</u> 1 (2014). ISSN: 20524463. DOI: 10.1038/sdata.2014.46. URL: http://www.nature.com/articles/sdata201446.

[18]   Elizabeth M. Sweeney, **Joshua T. Vogelstein**, Jennifer L. Cuzzocreo, Peter A. Calabresi, Daniel S. Reich, Ciprian M. Crainiceanu, and Russell T. Shinohara. "A comparison of supervised machine learning algorithms and feature vectors for MS lesion segmentation using multimodal structural MRI". In: <u>PLoS ONE</u> 9.4 (2014). ISSN: 19326203. DOI: 10.1371/journal.pone.0095753. URL: https://doi.org/10.1371/journal.pone.0095753.

[17]   **Joshua T. Vogelstein**, Youngser Park, Tomoko Ohyama, Rex A. Kerr, James W. Truman, Carey E. Priebe, and Marta Zlatic. "Discovery of brainwide neural-behavioral maps via multiscale unsupervised structure learning". In: <u>Science</u> 344.6182 (2014), pp. 386–392. ISSN: 10959203. DOI: 10.1126/science.1250298. URL: https://science.sciencemag.org/content/344/6182/386.

[16]   R. Cameron Craddock, Saad Jbabdi, Chao Gan Yan, **Joshua T. Vogelstein**, F. Xavier Castellanos, Adriana Di Martino, Clare Kelly, Keith Heberlein, Stan Colcombe, and Michael P. Milham. "Imaging human connectomes at the macroscale". In: <u>Nature Methods</u> 10.6 (2013), pp. 524–539. ISSN: 15487091. DOI: 10.1038/nmeth.2482. eprint: NIHMS150003. URL: https://doi.org/10.1038/nmeth.2482.

[15]   Dai Dai, Huiguang He, **Joshua T. Vogelstein**, and Zengguang Hou. "Accurate prediction of AD patients using cortical thickness networks". In: <u>Machine Vision and Applications</u> 24.7 (2013), pp. 1445–1457. ISSN: 09328092. DOI: 10.1007/s00138-012-0462-0. URL: https://doi.org/10.1007/s00138-012-0462-0.

[14]   **Joshua T. Vogelstein**, *William Gray Roncal*, R. Jacob Vogelstein, and Carey E. Priebe. "Graph classification using signal-subgraphs: Applications in statistical connectomics". In: <u>IEEE Transactions on Pattern Analysis and Machine Intelligence</u> 35.7 (2013), pp. 1539–1551. ISSN: 01628828. DOI: 10.1109/TPAMI.2012.235. arXiv: 1108.1427. URL: https://doi.org/10.1109/TPAMI.2012.235.

[13]   Carey E. Priebe, **Joshua Vogelstein**, and Davi Bock. "Optimizing the quantity/quality trade-off in connectome inference". In: Communications in Statistics - Theory and Methods 42.19 (2013), pp. 3455–3462. ISSN: 03610926. DOI: 10.1080/03610926.2011.630768. arXiv: 1108.6271. URL: https://doi.org/10.1080/03610926.2011.630768.

[12]   *William R. Gray*, John A. Bogovic, **Joshua T. Vogelstein**, Bennett A. Landman, Jerry L. Prince, and R. Jacob Vogelstein. "Magnetic Resonance Connectome Automated Pipeline: An Overview". In: IEEE Pulse 3.2 (Mar. 2012), pp. 42–48. ISSN: 21542287. DOI: 10.1109/MPUL.2011.2181023. URL: http://ieeexplore.ieee.org/document/6173097/.

[11]   Nicholas J Roberts, **Joshua T Vogelstein**, Giovanni Parmigiani, Kenneth W Kinzler, Bert Vogelstein, and Victor E Velculescu. "The predictive capacity of personal genome sequencing". In: Science Translational Medicine 4 (2012). ISSN: 19466234. DOI: 10.1126/scitranslmed.3003380. URL: https://doi.org/10.1126/scitranslmed.3003380.

[10]   Donniell E. Fishkind, Daniel L. Sussman, Minh Tang, **Joshua T. Vogelstein**, and Carey E. Priebe. "Consistent adjacency-spectral partitioning for the stochastic block model when the model parameters are unknown". In: SIAM Journal on Matrix Analysis and Applications 34.1 (2012), pp. 23–39. ISSN: 0895-4798. DOI: 10.1137/120875600. arXiv: 1205.0309. URL: http://arxiv.org/abs/1205.0309.

[9]   **Joshua T. Vogelstein**, R. Jacob Vogelstein, and Carey E. Priebe. "Are mental properties supervenient on brain properties?" In: Scientific Reports 1 (2011). ISSN: 20452322. DOI: 10.1038/srep00100. URL: https://doi.org/10.1038/srep00100.

[8]   Yuriy Mishchencko, **Joshua T Vogelstein**, and Liam Paninski. "A Bayesian approach for inferring neuronal conectivity from calcium fluorescent imaging data". In: The annals of applied statistics 5 (2011). ISSN: 19326157. DOI: 10.1214/09-AOAS303. URL: https://doi.org/10.1214/09-AOAS303.

[7]   Sonja B. Hofer, Ho Ko, Bruno Pichler, **Joshua Vogelstein**, Hana Ros, Hongkui Zeng, Ed Lein, Nicholas A. Lesica, and Thomas D. Mrsic-Flogel. "Differential connectivity and response dynamics of excitatory and inhibitory neurons in visual cortex". In: Nature Neuroscience 14.8 (2011), pp. 1045–1052. ISSN: 10976256. DOI: 10.1038/nn.2876. URL: https://doi.org/10.1038/nn.2876.

[6]   Liam Paninski, Yashar Ahmadian, Daniel Gil Ferreira, Shinsuke Koyama, Kamiar Rahnama Rad, Michael Vidne, **Joshua Vogelstein**, and Wei Wu. "A new look at state-space models for neural data". In: Journal of Computational Neuroscience 29.1-2 (2010), pp. 107–126. ISSN: 09295313. DOI: 10.1007/s10827-009-0179-x. URL: https://doi.org/10.1007/s10827-009-0179-x.

[5]   **Joshua T. Vogelstein**, Brendon O. Watson, Adam M. Packer, Rafael Yuste, Bruno Jedynak, and Liam Paninski. "Spike inference from calcium imaging using sequential Monte Carlo methods". In: Biophysical Journal 97.2 (2009), pp. 636–655. ISSN: 15420086. DOI: 10.1016/j.bpj.2008.08.005. URL: https://doi.org/10.1016/j.bpj.2008.08.005.

[4]   **Joshua T Vogelstein**, Adam M Packer, Tim A Machado, Tanya Sippy, Baktash Babadi, Rafael Yuste, and Liam Paninski. "Fast non-negative deconvolution for spike train inference from population calcium imaging". In: Journal of Neurophysiology 104 (2009). ISSN: 0022-3077. DOI: 10.1152/jn.01073.2009. arXiv: 0912.1637. URL: https://doi.org/10.1152/jn.01073.2009.

[3]   R. Jacob Vogelstein, Udayan Mallik, **Joshua T. Vogelstein**, and Gert Cauwenberghs. "Dynamically reconfigurable silicon array of spiking neurons with conductance-based synapses". In: IEEE Transactions on Neural Networks 18.1 (2007), pp. 253–265. ISSN: 10459227. DOI: 10.1109/TNN.2006.883007. URL: https://doi.org/10.1109/TNN.2006.883007.

[2]   **Joshua T. Vogelstein**, Lawrence H. Snyder, and Dora E. Angelaki. "Accuracy of saccades to remembered targets as a function of body orientation in space". In: Journal of Neurophysiology 90.1 (2003), pp. 521–524. ISSN: 00223077. DOI: 10.1152/jn.00141.2003. URL: https://doi.org/10.1152/jn.00141.2003.

[1]   David L. Greenspan, Denise C. Connolly, Rong Wu, Rachel Y. Lei, **Joshua T.C. Vogelstein**, Young Tak Kim, Jung Eun Mok, Nubia Muñoz, F. Xavier Bosch, Keerti Shah, and Kathleen R. Cho. "Loss of FHIT expression in cervical carcinoma cell lines and primary tumors". In: Cancer Research 57 (1997). ISSN: 00085472. URL: http://cancerres.aacrjournals.org/content/57/21/4692.

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Teaching

Ongoing Courses

Fall '19 **NeuroData Design I**, *EN.580.237/437/637*, Course Director, enrollment 46.

Spring '19 **NeuroData Design II**, *EN.580.438/638*, Course Director, enrollment 18.

Fall '18 **NeuroData Design I**, *EN.580.237/437/637*, Course Director, enrollment 22.

Spring '17 **NeuroData Design II**, *EN.580.238/438/638*, Course Director, enrollment 14.

Winter '17 **BME Research Intersession**, *EN.580.574*, Course Director, enrollment 6.

Fall '17 **NeuroData Design I**, *EN.580.247/437/637*, Course Director, enrollment 15.

Spring '16 **The Art of Data Science**, *EN.580.468*, Course Director, enrollment 24.

Fall '16 **NeuroData Design I**, *EN.580.437*, Course Director, enrollment 16.

Fall '15 **Introduction to Computational Medicine**, Co-Teaching, Course Co-Director.

Spring '15 **Statistical Connectomics**, *EN.580.694*, Course Director, enrollment 26.

One-Time Courses

Spring '19 **Systems Bioengineering II**, *EN.580.422*, 2 Lectures.

Spring '19 **Computational Neuroscience**, *AS.080.321*, 2 Lectures.

Spring '18 **Systems Bioengineering II**, *EN.580.422*, 2 Lectures.

Spring '18 **Computational Neuroscience**, *AS.080.321*, 2 Lectures.

Spring '17 **Systems Bioengineering II**, *EN.580.422*, 2 Lectures.

Spring '16 **Systems Bioengineering II**, *EN.580.422*, 2 Lectures.

Winter '16 **Introduction to Connectomics**, *EN.600.221*, 1 Lecture.

Fall '16 **BME Modeling and Design**, *EN.580.111*, 1 Lecture.

Educational Workshops

Summer '19   **DiPy Workshop**, Bloomington, Indiana, 1 day lecture on statistical connectomics.

Fall '18   **Society for Neuroscience Annual Meeting**, *Educational Workshop*, San Diego, CA, 1 day lecture on statistical connectomics.

Fall '17   **Society for Neuroscience Annual Meeting**, *Educational Workshop*, San Diego, CA, 1 day lecture on statistical connectomics.

Summer '16   **CRCNS Course on Mining and Modeling of Neuroscience Data**, *Redwood Center for Theoretical Neuroscience*, University of California, Berkeley, 2 day lecture on statistical connectomics.

Mentorship

Research Track Faculty Mentorship

02/19 – now   **Hayden Helm, MSE**, *Assistant Research Faculty*, BME, JHU.
Leading research efforts developing theory and methods for lifelong learning.

08/16 – 8/18   **Eric Perlman, PhD**, *Assistant Research Scientist*, BME, JHU.
Lead Scientist developing storage, transfer, and visualization solutions for large data in our cloud infrastructure.

03/16 – now   **Jesse Patsolic, MA**, *Assistant Research Faculty*, BME, JHU.
Lead developer converting our extensions to decision forests to be merged into sklearn.

Postdoctoral Fellows and Staff Research Scientists

10/18 – now   **Alex Loftus**, *Research Assistant*, BME, JHU.
Current lead developer of NDMG, transitioning from a stand-alone package to be integrated with DiPy.

09/19 – now   **Ross Lawrence, BS**, *Research Assistant*, BME, JHU.
Responsible for documenting and bug fixing NDMG.

07/19 – now   **Ronak Mehta, MSE**, *Research Assistant*, BME, JHU.
Finalizing three manuscripts on (1) uncertainty forests, (2) time-series dependence quantification, and (3) lifelong learning forests.

07/19 – now   **Celine Drieu, PhD**, *Post-doctoral Fellow*, Kavli NDI, JHU.
Co-Advised by Assitant Prof. Kuchibhotla, Department of Psychological and Brain Sciences. Working on understanding learning and memory using two-photon calcium imaging.

07/19 – now   **Austin Grave, PhD**, *Post-doctoral Fellow*, Kavli NDI, JHU.
Co-Advised by Prof. Richard Huganir, Department of Neuroscience. Working on understanding whole brain synaptic plasticity using genetic engineering and light microscopy imaging.

06/19 – now   **Devin Crowley**, *Research Assistant*, BME, JHU.
Lead developer of our scalable Python implementaiton of LDDMM.

08/18 – now   **Jesús Arroyo, PhD**, *Post-doctoral Fellow*, CIS, JHU.
Working on graph matching and joint graph embedding.

06/18 – now   **Benjamin Falk, PhD**, *Research Engineer*, BME, JHU.
Lead software engineer, oversees all development projects, solely responsible for all cloud infrastructure.

07/18 – now   **Audrey Branch, PhD**, *Post-doctoral Fellow*, Kavli NDI, JHU.
Co-Advised by Prof Michela Gallagher, extending brain clearing experimental technology from mice to rats. Currently with a manuscript on biorxiv.

09/16 – 08/18   **Cencheng Shen, PhD**, *Post-Doctoral Fellow*, CIS, JHU.
Developed Multiscale Graph Correlation, which is currently the premiere hypothesis testing framework, and about to be integrated into SciPy, by far the world's leading scientific computing package. Currently an Assistent Professor in Department of Statistics at University of Delaware, and still an actice collaborator and grantee.

05/16 – 06/17   **Leo Duan, PhD**, *Post-doctoral Fellow*, CIS, JHU.
Went on to do a second postdoc with Leo Dunson (who I did my second postdoc with). Currently an Assistant Professor at University of Florida.

06/16 – 07/17   **Guilherme Franca, PhD**, *Post-doctoral Fellow*, CIS, JHU.
Worked on non-parametric clustering, with an article about to be accepted in PAMI, the leading machine learning journal. Currently a postdoc for Rene Vidal.

Doctoral Student Supervision

08/19 – now   **Michael Powell, MSE**, *PhD advisee*, BME, JHU.
Dissertation will focus on explainable artificial intelligence, spearheads collaboration with Andreas Muller, Co-Director of scikit-learn, the world's leading machine learning package.

06/19 – now   **Jaewon Chung, MSE**, *PhD advisee*, BME, JHU.
Dissertation will focus on statistics of populations of human networks. Already co-first author and middle author on multiple manuscripts.

08/19 – now   **Tommy Athey, BSE**, *PhD advisee*, BME, JHU.
Dissertation will focus on MouseLight project, spearheads collaborations with Prof. Jeremias Sulam and Michael I. Miller.

08/19 – now   **Eric Bridgeford, BSE**, *PhD advisee*, Department of Biostatistics, JHU.
Dissertation will focus on statistics of human connectomes and mitigating batch effects. Already first author on several manuscripts under review, and spearheads collaboration with Prof Brian Caffo at Biostatistics.

08/18 – now   **Benjamin Pedigo, BSE**, *PhD advisee*, BME, JHU.
Dissertation will focus on analysis and modeling of the world's first whole animal connectome, in collaboration with Marta Zlatic and Albert Cardona (formerly of Janelia Research Campus). Already co-first author and middle author on multiple manuscripts.

08/18 – now   **Meghana Madyastha, BSE**, *PhD Co-advisee*, CS, JHU.
Dissertation will focus on computational aspects of accelerationg learning and inference using decision forests.

08/16 – now   **Vikram Chandrashekhar, BSE**, *PhD advisee*, BME, JHU.
Dissertation has focused on extending LDDMM to whole cleared brain datasets, spearheads collaboration with Prof. Karl Deisseroth's lab at Stanford, one of the world's leading neuroscientists.

08/14 – 01/18   **Tyler Tomita, PhD**, BME, JHU.
Developed Sparse Projection Oblique Randomer Forest in his dissertation, currently the best performing machine learning algorithm on a standard suite of over 100 benchmark problems. Currenly a postdoc with Assistant Prof. Chris Honey of Psychology and Brain Sciences.

Master's Student Supervision

06/19 – now   **Bijan Varjavand**, *MS advisee*, BME, JHU.
Submitted manuscript to PAMI on advancing statistics on populations of networks.

06/19 – now   **Sambit Panda**, *MS advisee*, BME, JHU.
Led development of Python implementation of MGC, to be integrated into SciPy.

06/19 – now   **Varun Kotharkar**, *MS advisee*, AMS, JHU.
Investigating theoretical advantages of oblique, as compared to axis-aligned, decision trees.

06/18 – now   **Drishti Mannan**, *MS advisee*, BME, JHU.
Preparing manuscript introducing novel specification for large attributed networks.

06/18 – 05/19   **Jaewon Chung**, *MSE advisee*, BME, JHU.
Co-first author of manuscript and co-lead developer of Python package for statistical analysis of networks. Currently a BME PhD student in my lab.

08/14 – 06/17   **Greg Kiar, MSE**, BME, JHU.
Lead deveoper of NDMG, the only existing "soup to nuts" pipeline for both functional and diffusion pipelines; co-first author of manuscript under review. Currently a PhD student at McGill University.

Undergraduate Student Supervision

06/19 – now   **Vivek Gopalakrishnan**, *BSE*, BME, JHU.
Winner of Pistritto Fellowship.

06/19 – now   **Richard Guo**, *BSE*, BME, JHU.

06/19 – now   **Ronan Perry**, *BSE*, BME, JHU.

08/14 – 08/18 **Eric Bridgeford, BSE**, BME, JHU.
Currently a PhD student in Biostatistics at JHSPH in my lab.

08/15 – 08/16 **Albert Lee,BSE**, BME, JHU.

06/15 – 12/15 **Ron Boger, BSE**, BME, JHU.
Currently working at a computational medicine start-up in Silicon Valley.

05/15 – 05/16 **Jordan Matelsky, BSE**, CS and Neuroscience, JHU.
Currently a data scientist at APL.

02/15 – 05/16 **Ivan Kuznetsov, BSE**, BME, JHU.
Currently an MD/PhD Candidate at the UPenn, winner of Soros Fellowship.

Summer Interns

Summer '19 **Kareef Ullah**, *Summer Intern*, BME, JHU.

Summer '19 **Shunan Wu**, *Summer Intern*, BME, JHU.

Summer '19 **Shiyu Sun**, *Summer Intern*, BME, JHU.

Summer '19 **Sander Shulhoff**, *Summer Intern*, BME, JHU.

Summer '19 **Kiki Zhang**, *Summer Intern*, BME, JHU.

Summer '18 **Papa Kobina Van Dyck**, *Summer Intern*, BME, JHU.

Examining Committees

2019 **Browne, James**, *Computer Science*, JHU Ph.D. Student, Graduated 2019.

2019 **Mhembere, Disa**, *Computer Science*, JHU Ph.D. Student, Graduated 2019.

2018 **Kutten, Kwame**, *JHU Ph.D. Student*, Graduated 2018.

2018 **Wang, Shangsi**, *Applied Mathematics and Statistics*, JHU Ph.D. Student, Graduated 2018.

2018 **Tang, Runze**, *Applied Mathematics and Statistics*, JHU Ph.D. Student, Graduated 2018.

2018 **Lee, Youjin**, *Biostatistics*, JHU Ph.D. Student, Graduated 2018.

2017 **Zheng, D**, *Computer Science*, JHU Ph.D. Student, Graduated 2017.

2017 **Binkiewicz, Norbert**, *Statistics*, University of Wisconsin Ph.D. Student, Graduated 2017.

2016 **Gray-Roncal, Will**, *Computer Science*, JHU Ph.D. Student, Graduated 2016.

Awards and Recognition

Individual

2002 **Dean's List**, *Washington University*.

Shared

2019 **Kavli NDI Distinguished Postdoctoral Fellow**, *Celine Drieu, PhD*.

2019 **Kavli NDI Distinguished Postdoctoral Fellow**, *Austin Graves, PhD*.

2017 **Kavli NDI Distinguished Postdoctoral Fellow**, *Audrey Branch, PhD*.

2017 **Best Presentation Award HPDC**, *Mhembere et al. (2017)*.

2017 **Nonparametric Statistics of the American Statistical Association Student Paper Award**, *Lee et al. (2017)*.

2014 **F1000 Prime Recommended**, *Vogelstein et al. (2014)*.

2013 **Spotlight**, *Neural Information Processing Systems (NIPS)*.

2011 **Trainee Abstract Award**, *Organization for Human Brain Mapping*.

2008 **Spotlight**, *Computational and Systems Neuroscience (CoSyNe)*.

Service

University Service

2019 **Member**, *Search Committee*, BME, Neuroengineering, 2019.

2019 **Member**, *Search Committee*, BME, Data Science, 2019.

2018	**Member**, *Search Committee*, BME, Neuroengineering, 2018.
Winter '17	**Faculty Superviser**, *MedHacks*, http://medhacks.org/.
Winter '16	**Faculty Superviser**, *MedHacks*, http://medhacks.org/.
Spring '16	**Organizer**, *Global Brain Workshop*, http://brainx.io, First ever international Brain Initiative workshop, bringing together leaders from around the world, covered by Nature and Science (~ 75 participants).
Winter '15	**Faculty Superviser**, *MedHacks*, http://medhacks.org/.

Department Service

Winter '19	**Organizer**, *Decision Forest Hackathon*.
Summer '19	**Organizer**, *NeuroData Workshop*, https://neurodata.devpost.com, Hackashop to train brain scientists in machine learning for big data (~ 50) participants from around the country..
March '19	**Organizer**, *Neuro Reproducibility Hackashop*, https://brainx3.io/, Hackashop to train brain scientists in best practices in reproducible science, co-organized with two startups: Vathes, LLC and Gigantum (~ 50 participants).
Spring '18	**Organizer**, *NeuroData Hackathon*.
Fall '17	**Organizer**, *NeuroData Mini-Hackathon*.
Summer '17	**Organizer**, *NeuroStorm*, https://brainx2.io, Workshop bring together thought leaders from academia, national labs, industry, and non-profits around the world to take next steps towards accelerating brain science discovery in the cloud (~ 50 participants and 5 observers from funding institutions).
Winter '15	**Organizer**, *Hack@NeuroData*, http://hack.neurodata.io/.
08/15 – now	**Steering Committee**, *Kavli Neuroscience Discovery Institute (KNDI)*.
08/15 – 07/18	**Co-Developer**, *Computational Medicine Minor*.
05/15 – 07/17	**Co-Founder and Faculty Advisor**, *MedHacks*.
08/14 – 08/18	**Director of Undergraduate Studies**, *Institute for Computational Medicine*.

Journal Service

Editorial Board

Guest Associate Editor, *PLoS Computational Biology*.

Editor, *Neurons, Behavior, Data analysis, and Theory*.

Associate Editor, *Journal of the American Statistical Association*.

Conference and Journal Reviewer

Annals of Applied Statistics (AOAS).

Bioinformatics.

International Conference on Learning Representations (ICLR).

Network Science.

Current Opinion in Neurobiology.

Biophysical Journal.

IEEE International Conference on eScience.

IEEE International Conference on Acoustics, Speech, and Signal Processing (ICASSP).

IEEE Global Conference on Signal and Information Processing (GlobalSIP).

IEEE Signal Processing Letters.

IEEE Transactions on Signal Processing.

Frontiers in Brain Imaging Methods.

Journal of Machine Learning Research (JMLR).

Journal of Neurophysiology.

Journal of the Royal Statistical Society B (JRSSB).

> **Nature Communications**.
> **Nature Methods**.
> **Nature Reviews Neuroscience**.
> **Neural Computation**.
> **Neural Information Processing Systems (Neurips)**.
> **NeuroImage**.
> **Neuroinformatics**.
> **PLoS One**.
> **PLoS Computational Biology**.

Professional Service

Fall '16   **Co-Organizer**, *Brains and Bits: Neuroscience Meets Machine Learning, NIPS Workshop*, http://www.stat.ucla.edu/~akfletcher/brainsbits_overview.html.

Fall '15   **Co-Organizer**, *BigNeuro2015: Making Sense of Big Neural Data, NIPS Workshop*, http://neurodata.io/bigneuro2015.

Fall '12   **Co-Organizer**, *Scaling up EM Connectomics Conference*, The world's first connectomics workshop, now run annually alternating between Janelia Research and Max Plank locations ($\sim$ 80 participants).

Translation / Technology Transfer Activities

Open Datasets

1. Allen Atlas These anatomical reference atlases illustrate the adult mouse brain in coronal and sagittal planes of section. They are the spatial framework for datasets such as in situ hybridization, cell projection maps, and in vitro cell characterization. More information at atlas.brain-map.org

2. Amunts et al. (2015) Enabling an unprecedented look into the human brain, BigBrain spans micro- and macro-scopic scales. While previously available reference brains have been restricted to a single scale, such as whole-brain magnetic resonance imaging in humans or electron microscopy of small sections from small animals, BigBrain is an ultrahigh-resolution three-dimensional model of a full human brain at 20 micrometer resolution, coming closer to touching both camps than any previous dataset

3. Bhatla et al. (2015) Using high-pressure freezing, serial section transmission electron microscopy (ssTEM) imaging, digital alignment and manual tracing, Nikhil Bhatla and Rita Droste in Bob Horvitz's Lab reconstructed the anterior half of the C. elegans feeding organ, the pharynx. Volumes are available for three adult hermaphrodite worms and include volumetric tracing of all neurons and selected cell types, as well as synapses identified from the I2 neurons. Sections were approximately 50 nm thick with an image resolution of 2 nm per pixel. The largest volume comprises 1199 slices. These data were published in a paper entitled "Distinct neural circuits control rhythm inhibition and spitting by the myogenic pharynx of C. elegans" (Current Biology, 2015)

4. Bloss et al. (2016) Neuronal circuit function is governed by precise patterns of connectivity between specialized groups of neurons. The diversity of GABAergic interneurons is a hallmark of cortical circuits, yet little is known about their targeting to individual postsynaptic dendrites. We examined synaptic connectivity between molecularly defined inhibitory interneurons and CA1 pyramidal cell dendrites using correlative light-electron microscopy and large-volume array tomography. We show that interneurons can be highly selective in their connectivity to specific dendritic branch types and, furthermore, exhibit precisely targeted connectivity to the origin or end of individual branches. Computational simulations indicate that the observed subcellular targeting enables control over the nonlinear integration of synaptic input or the initiation and backpropagation of action potentials in a branchselective manner. Our results demonstrate that connectivity between interneurons and pyramidal cell dendrites is more precise and spatially segregated than previously appreciated, which may be a critical determinant of how inhibition shapes dendritic computation

5. Bloss et al. (2018) —

6. Bock et al. (2011) Layer 2/3 - Davi Bock, Ph.D. and Wei-Chung Allen Lee, Ph.D., in the laboratory of Clay Reid, M.D., Ph.D. acquired a beautiful volume of mouse primary visual cortical data, spanning layers 1, 2/3, and upper layer 4. In addition to the electron microscope (EM) data, they used two-photon microscopy to

determine the functional properties of about 14 of the cells in the same volume. Images were collected at approximately 4x4x45 cubic nanometers with a total volume of approximately 450x350x50 cubic microns

7. Branch (2018) Adult generated neurons in aging M. musculus (iDisco)

8. Bumbarger et al. (2013) These serial thin section data were generated by Dan Bumbarger in Ralf Sommer's lab in order to compare the pharyngeal connectomes of the pharyngeal nervous system between Caenorhabditis elegans and Pristionchus pacificus. (Cell 2013, 152:109–119). In P. pacificus they found clearly homologous neurons for all of the 20 pharyngeal neurons in C. elegans, but were surprised to uncover a massive rewiring of synaptic connectivity between the two species. These changes seem to correlate with known behavioral difference, most interestingly with the novel predatory feeding behaviors found in Diplogastrid nematodes such as P. pacificus

9. Collman et al. (2015) Synapses of the mammalian CNS are diverse in size, structure, molecular composition, and function. Synapses in their myriad variations are fundamental to neural circuit development, homeostasis, plasticity, and memory storage. Unfortunately, quantitative analysis and mapping of the brain's heterogeneous synapse populations has been limited by the lack of adequate single-synapse measurement methods. Electron microscopy (EM) is the definitive means to recognize and measure individual synaptic contacts, but EM has only limited abilities to measure the molecular composition of synapses. This report describes conjugate array tomography (AT), a volumetric imaging method that integrates immunofluorescence and EM imaging modalities in voxel-conjugate fashion. We illustrate the use of conjugate AT to advance the proteometric measurement of EM-validated single-synapse analysis in a study of mouse cortex

10. Deisseroth et al. (2015) Twelve CLARITY mouse brains (5 wild type controls and 7 behaviorally challenged) were prepared by Li Ye, and imaged using CLARITY-Optimized Light-sheet Microscopy (COLM) (whole brain COLM imaging and data stitching performed by R. Tomer, in preparation)

11. Dyer et al. (2016) Methods for resolving the 3D microstructure of the brain typically start by thinly slicing and staining the brain, and then imaging each individual section with visible light photons or electrons. In contrast, X-rays can be used to image thick samples, providing a rapid approach for producing large 3D brain maps without sectioning. Here we demonstrate the use of synchrotron X-ray microtomography (microCT) for producing mesoscale (1 cubic micron resolution) brain maps from millimeter-scale volumes of mouse brain. We introduce a pipeline for mircoCT-based brain mapping that combines methods for sample preparation, imaging, automated segmentation of image volumes into cells and blood vessels, and statistical analysis of the resulting brain structures. Our results demonstrate that X-ray tomography promises rapid quantification of large brain volumes, complementing other brain mapping and connectomics efforts

12. Harris et al. (2015) From the laboratory of Kristen M Harris, PhD, three volumes of hippocampal CA1 neuropil in adult rat were imaged at an XY resolution of 2 nm on serial sections of 50-60 nm thickness. All axons, dendrites, glia, and synapses were reconstructed in a cube surrounding a large dendritic spine, a cylinder surrounding an oblique dendritic segment, and a parallelepiped surrounding an apical dendritic segment

13. Hildebrand et al. (2017) Hildebrand and colleagues acquired a multi-resolution serial-section electron microscopy data set containing the anterior quarter of a 5.5 days post fertilization larval zebrafish, including its complete brain. A draft projectome consisting of central and peripheral myelinated neurons was then reconstructed. Electron micrographs and reconstructions are available for view in CATMAID. A manuscript describing the data and methods used to generate it has been published in Nature

14. Kasthuri et al. (2015) We describe automated technologies to probe the structure of neural tissue at nanometer resolution and use them to generate a saturated reconstruction of a sub-volume of mouse neocortex in which all cellular objects (axons, dendrites, and glia) and many sub-cellular components (synapses, synaptic vesicles, spines, spine apparati, postsynaptic densities, and mitochondria) are rendered and itemized in a database. We explore these data to study physical properties of brain tissue. For example, by tracing the trajectories of all excitatory axons and noting their juxtapositions, both synaptic and non-synaptic, with every dendritic spine we refute the idea that physical proximity is sufficient to predict synaptic connectivity (the so-called Peters' rule). This online minable database provides general access to the intrinsic complexity of the neocortex and enables further data-driven inquiries

15. Lee et al. (2016) Electron Microscopy data used in a study of an excitatory network in Mouse V1

16. Micheva et al. (2015) Multi-channel array tomography data which is barrel cortex from an adult mouse (C57BL/6J)

17. Ohyama et al. (2015) Understanding brain function and development would be facilitated enormously by

being able to perform all experiments on the basis of known circuitry. Over 20 laboratories world wide have contributed towards the reconstruction of neurons in the central nervous system of Drosophila larva, led by the Cardona lab at HHMI Janelia. Here, we see a side view of the approximately 7,000 neurons reconstructed so far, either in full or partially, of the approximately 12,000 neurons of this animal. The 0111-8 data set was originally sectioned and imaged by Richard D. Fetter and his two tech assistants, and funded by the HHMI Janelia Fly EM Project Team. There are now many more papers now using the 0111-8 data (see publications below)

18. Takemura et al. (2013) The right part of the brain of a wild-type Oregon R female fly was serially sectioned into 40-nm slices. A total of 1,769 sections, traversing the medulla and downstream neuropils, were imaged at a magnification of 35,000X

19. Templier et al. (2019) The non-destructive collection of ultrathin sections onto silicon wafers for post-embedding staining and volumetric correlative light and electron microscopy traditionally requires exquisite manual skills and is tedious and unreliable. In MagC introduced here, sample blocks are augmented with a magnetic resin enabling remote actuation and collection of hundreds of sections on wafer. MagC allowed the correlative visualization of neuroanatomical tracers within their ultrastructural volumetric electron microscopy context

20. Tobin et al. (2017) Wiring variations that enable and constrain neural computation in a sensory microcircuit

21. Wanner et al. (2016 Large-scale reconstructions of neuronal populations are critical for structural analyses of neuronal cell types and circuits. Dense reconstructions of neurons from image data require ultrastructural resolution throughout large volumes, which can be achieved by automated volumetric electron microscopy (EM) techniques. We used serial block face scanning EM (SBEM) and conductive sample embedding to acquire an image stack from an olfactory bulb (OB) of a zebrafish larva at a voxel resolution of $9.25 \times 9.25 \times 25$ nm3 (Wanner et al., 2016). Skeletons of 1,022 neurons, $\sim 98\%$ of all neurons in the OB, were reconstructed by manual tracing and efficient error correction procedures

22. Weiler (2014) The lab of Stephen J Smith has been developing array tomography technology for nearly a decade (Micheva et al., 2007). This technology is unique in its ability to measure many proteins (20 or more) in biological tissue samples with superresolution precision. In this dataset, we are using array tomography to uncover molecular signatures synaptic diversity (O'Rourke et al., 2012), which is fundamental to neural circuit design and function. Images generously donated by Nick Weiler

23. Randlett et al. (2015) Light microscopy data

Open-source Software: Active

1. GraSPy (Graph Statistics) Utilities and algorithms designed for processing and analysis of graphs with specialized graph statistical algorithms
2. MGC (Non-parametric hypothesis testing) Multiscale Graph Correlation (MGC) is a framework for universally consistent testing high-dimensional and non-Euclidean data.
3. ndcloud (NeuroData Cloud) The deployment of tools which support the Open Connectome Project
4. Sparse Projection Oblique Randomer Forests (Classification and regression) SPORF is an improved random forest algorithm that achieves better accuracy and scaling than previous implementations on a standard suite of >100 benchmark problems.
5. LOL (Supervised dimensionality reduction) Linear Optimal Low-rank (LOL) projection for improved classification performance in high-dimensional classification tasks
6. m2g (MR graph analysis) m2g uses diffusion MRI data from individuals to estimate connectomes reliably and scalably.
7. reg (Image registration) Performs non-linear affine and deformable image registration.
8. Uncertainty-Forest A Python package containing estimation procedures for posterior distributions, conditional entropy, and mutual information between random variables X and Y
9. Open-Data-Registry This bucket contains multiple neuroimaging datasets (as Neuroglancer Precomputed Volumes) across multiple modalities and scales, ranging from nanoscale (electron microscopy), to microscale (cleared lightsheet microscopy and array tomography), and mesoscale (structural and functional magnetic resonance imaging). Additionally, many of the datasets include segmentations and meshes.
10. OCP The Open Connectome Project
11. neuroparc This repository contains a number of useful parcellations, templates, masks, and transforms to (and from) MNI152NLin6 space. The files are named according to the BIDs specification.
12. ndex Python 3 command-line program to exchange (download/upload) image data with NeuroData's

cloud deployment of APL's BOSS spatial database

13. ndwebtools ndwebtools (ndweb) is a Django application to provide a user-friendly interface for interacting with NeuroData resources and data.
14. Non-Parametric-Clustering

Open-source Software: Contributed

1. scipy
2. render
3. neuroglancer
4. boss
5. cloud-volume
6. igraph
7. C-PAC

Open-source Software: Archived

1. FlashGraph (Scalable Analytics)
2. FlashX (Scalable machine learning)
3. knor (Clustering)
4. MEDA (Matrix Exploratory Data Analysis)
5. oopsi (Calcium Spike Sorting)
6. SynapseAnalysis (Synapse Detection)
7. VESICLE (EM Synapse Detection)
8. ndviz
9. ndstore
10. CAJAL
11. DMG
12. vesicle

Consultancy

2017   **Consultant**, *Greenspring Associates*.

2016   **Consultant**, *Scanadu*.

Advisory Board Appointments

10/18 – now   **Advisory Board**, *Mind-X*.

01/17 – now   **Advisory Board**, *PivotalPath*.

Startups

01/17 – now   **Co-Founder**, *gigantum*.

01/16 – now   **Co-Founder**, *d8alab*.

01/11 – now   **Co-Founder & Co-Director**, *NeuroData* (formerly Open Connectome Project).