

Paul W. Hook

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300 E. 26th Street, Baltimore, MD 21218, USA

EDUCATION

Johns Hopkins School of Medicine, Baltimore, MD
Ph.D., Human Genetics

2014 – 2020

The Pennsylvania State University, University Park, PA
B.S., Biochemistry and Molecular Biology
Area of concentration: Molecular and Cell Biology

May 2012

RESEARCH EXPERIENCE

Postdoctoral Fellow

2020 – Present

Whiting School of Engineering
Johns Hopkins University, Baltimore, MD
PI: Winston Timp

- Developing genomics approaches utilizing long-read sequencing to explore subjects including DNA methylation, protein-DNA binding, and information storage in RNA

Graduate Student

2014 – 2020

Johns Hopkins School of Medicine, Baltimore, MD
Advisor: Andrew McCallion

- Designed, performed, and analyzed RNA-seq and single-cell RNA-seq experiments on mouse dopaminergic neurons and established a scoring paradigm for prioritizing genes from Parkinson disease GWAS loci
- Analyzed chromatin data in order to identify cell populations relevant to schizophrenia disease risk and to prioritize disease-relevant variation in disease-associated loci

Research Technologist

2012 – 2014

Johns Hopkins School of Medicine, Baltimore, MD
Advisor: Andrew McCallion

- Elucidated the functional consequences of disrupting genes on somitogenesis and heart development in zebrafish and implemented the use of Cas9 nuclease genome editing in zebrafish and human cell culture

Science Undergraduate Laboratory Internship

Summer 2011

Department of Energy, Pacific Northwest National Laboratory, Sequim, WA
Advisor: Michael Huesemann

- Explored how temperature affects algal growth and algal lipid composition for the National Alliance for Advanced Biofuels and Bio-products team

Chemical Research Intern

2010 – 2011

The Pennsylvania State University, University Park, PA
Advisor: Joseph Keiser

- Adapted and developed experiments focused on exploring the biochemical components of peanuts for an undergraduate laboratory class
- Assisted in designing and building demonstrations for undergraduate chemistry lectures

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TEACHING EXPERIENCE

Teaching Assistant

Fall 2016

Evolution of Ideas in Human Genetics

Johns Hopkins University School of Medicine, Baltimore, MD

Presenter

November 14, 2015

The Genome Geeks Are In

Smithsonian National Museum of Natural History, Washington, DC

Peer Learning Assistant

Fall 2011

BMB 430: Developmental Biology

The Pennsylvania State University, University Park, PA

LEADERSHIP AND MENTORING

Shared Interest Group (SIG) Leader

2022 – Present

Bioinformatics and Computational Methods SIG

The American Society of Human Genetics

Membership Engagement Committee

2020 – Present

The American Society of Human Genetics

ASHG Annual Meeting Advisory Group

2022

The American Society of Human Genetics

Program Committee

2022

NIH Advanced Genomic Technology Development Annual Meeting

Student Faculty Representative

2016 – 2020

Human Genetics Pre-Doctoral Training Program

Johns Hopkins University School of Medicine, Baltimore, MD

Peer Mentoring Leader

2017 – 2020

Institute of Genetic Medicine Peer Mentoring Families

Johns Hopkins University School of Medicine, Baltimore, MD

PROFESSIONAL ASSOCIATIONS

The American Society of Human Genetics

2015 - Present

PREPRINTS AND PEER-REVIEWED PUBLICATIONS

Rhie, A., Nurk, S., Cechova, M., Hoyt, S.J., Taylor, D.J., Altemose, N., **Hook, P.W.**, Koren, S., Rautiainen, M., Alexandrov, I.A., ..., Eichler, E.E., O'Neill, R., Schatz, M.C., Miga, K.H., Makova, K.D., & Phillippy, A.M. (2023). The complete sequence of a human Y chromosome. *Nature*, 1–11. <https://doi.org/10.1038/s41586-023-06457-y>.

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Hook, P.W., and Timp, W. (2023). Beyond assembly: the increasing flexibility of single-molecule sequencing technology. *Nat. Rev. Genet.* 24, 627–641.

Boyd, R.J., McClymont, S.A., Barrientos, N.B., **Hook, P.W.**, Law, W.D., Rose, R.J., Waite, E.L., Rathinavelu, J., Avramopoulos, D., and McCallion, A.S. (2023). Evaluating the mouse neural precursor line, SN4741, as a suitable proxy for midbrain dopaminergic neurons. *BMC Genomics* 24, 306.

Razaghi, R., **Hook, P.W.**, Ou, S., Schatz, M. C., Hansen, K. D., Jain, M., & Timp, W. (2022). Modbamtools: Analysis of single-molecule epigenetic data for long-range profiling, heterogeneity, and clustering. *bioRxiv*, 2022.07.07.499188.

Gershman, A., Sauria, M.E.G., Guitart, X., Vollger, M.R., **Hook, P.W.**, Hoyt, S.J., Jain, M., Shumate, A., Razaghi, R., Koren, S., Altemose, N., Caldas, G.V., Logsdon, G.A., Rhie, A., Eichler, E.E., Schatz, M.C., O'Neill, R.J., Phillippy, A.M., Miga, K.H., & Timp, W. (2022). Epigenetic patterns in a complete human genome. *Science*, 376 (6588), eabj5089.

Soto-Beasley, A.I., Walton, R.L., Valentino, R.R., **Hook, P.W.**, Labbé, C., Heckman, M.G., Johnson, P.W., Goff, L.A., Uitti, R.J., McLean, P.J., Springer, W., McCallion, A.S., Wszolek, Z.K., & Ross, O.A. (2020). Screening non-MAPT genes of the Chr17q21 H1 haplotype in Parkinson's disease. *Parkinsonism & Related Disorders*, 78, 138–144.

Hook, P.W., & McCallion, A.S. (2020). Leveraging mouse chromatin data for heritability enrichment informs common disease architecture and reveals cortical layer contributions to schizophrenia. *Genome Research*, 30 (4): 528–39.

McClymont, S.A, **Hook, P.W.**, Soto, A.I., Reed, X., Law, W.D., Kerans, S.J., Waite, E.L., Briceno, N.J., Thole, J.F., Heckman, M.G., Diehl, N.N., Wszolek, Z.K., Moore, C.D., Zhu, H., Akiyama, J.A., Dickel, D.E., Visel, A., Pennacchio, L.A., Ross, O.A., Beer, M.A., McCallion, A.S. (2018). Parkinson Associated SNCA Enhancer Variants Revealed by Open Chromatin in Mouse Dopamine Neurons. *The American Journal of Human Genetics*, 103 (6), 874–892.

Hook, P.W., McClymont, S.A., Cannon, G.H., Law, W.D., Morton, A.J., Goff, L.A., & McCallion, A.S. (2018). Single-Cell RNA-Seq of Mouse Dopaminergic Neurons Informs Candidate Gene Selection for Sporadic Parkinson Disease. *The American Journal of Human Genetics*, 102 (3), 427–446.

Turner, T.N., Hormozdiari, F., Duyzend, M.H., McClymont, S.A., **Hook, P.W.**, Iossifov, I., ... Eichler, E.E. (2016). Genome Sequencing of Autism-Affected Families Reveals Disruption of Putative Noncoding Regulatory DNA. *The American Journal of Human Genetics*, 98 (1), 58–74.

Maragh, S., Miller, R.A., Bessling, S.L., Wang, G., **Hook, P.W.**, & McCallion, A.S. (2014). Rbm24a and Rbm24b are required for normal somitogenesis. *PLoS ONE*, 9 (8).

Van Wagenen, J., Miller, T.W., Hobbs, S., **Hook, P.**, Crowe, B., and Huesemann, M. (2012). Effects of light and temperature on fatty acid production in *Nannochloropsis salina*. *Energies* 5, 731–740.

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PLATFORM AND INVITED TALKS

Hook, P.W. “Targeted long-read sequencing for interrogation of cancer genetic loci.” Association of Biomolecular Resources Facilities Annual Meeting, Palm Springs, CA, 2022.

Hook, P.W., McCallion, A.S. “Refining cell populations and fine-mapping variants for schizophrenia and bipolar disorder using mouse open chromatin profiles” The American Society of Human Genetics, Houston, TX, 2019.

Hook, P.W., McClymont S.A., Cannon, G.H., Law, W.D., Morton, A.J., Goff, L.A., McCallion, A.S. “Prioritizing genes for sporadic Parkinson disease using single-cell expression profiling of mouse dopaminergic neurons” 11th Leena Peltonen School of Human Genomics, Les Diablerets, Switzerland, 2018.

Hook, P.W., McClymont, S.A., Goff, L.A., McCallion, A.S. “RNA-seq analysis identifies phenotypic heterogeneity among *ex vivo* purified dopamine neurons and highlights their progressive temporal diversification” The American Society of Human Genetics, Vancouver, BC, Canada, 2016.

POSTER PRESENTATIONS

Hook, P.W., Hosea, J.A., Morina, L.B., Ebenstein, Y., Simpson, J., Timp, W. “Measuring the epigenome with nanopore sequencing.” The Advances in Genomic Technology Development Annual Meeting, La Jolla, California, 2023.

Hook, P.W., Timp, W. “Protein-DNA interactions at the bench: CUT&RUN/Tag with nanopore sequencing.” The Advances in Genome Biology and Technology Annual Meeting, Hollywood, FL, 2023.

Hook, P.W., Krueger, F., Timp, W. “Adapting Enzymatic Methyl-seq (EM-seq) for long-read nanopore sequencing.” Nanopore Community Meeting, New York City, NY, 2022.

Hook, P.W., Krueger, F., Timp, W. “Adapting Enzymatic Methyl-seq (EM-seq) for long read sequencing.” The Advances in Genome Biology and Technology Annual Meeting, Orlando, FL, 2022.

HONORS AND AWARDS

ASHG/Charles J. Epstein Trainee Award for Excellence in Human Genetics Research – Semifinalist 2019
The American Society of Human Genetics - Houston, TX
“Refining cell populations and fine-mapping variants for schizophrenia and bipolar disorder using mouse open chromatin profiles”

C.W. Cotterman Award 2018
The American Society of Human Genetics – San Diego, CA
“Single-Cell RNA-Seq of Mouse Dopaminergic Neurons Informs Candidate Gene Selection for Sporadic Parkinson Disease”

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Leena Peltonen School of Human Genomics Trainee Les Diablerets, Switzerland	Summer 2018
Graduated with Distinction <i>Eberly College of Science</i> The Pennsylvania State University, University Park, PA	May 2012
Dean's List <i>Eberly College of Science</i> The Pennsylvania State University, University Park, PA	2008-2012
Gail A. and Thomas G. Ernst Scholarship	2009 – 2011
Kimberly Clark Bright Futures Scholarship	2008 – 2012