Andrew Hillandrewhill157@gmail.com | (860) 303-8849 | www.andrewjohnhill.com

Education

University of Washington, Seattle

Ph.D. in Genome Sciences (entering class of 2014)

University of Washington, Seattle

B.S. in Bioengineering with College Honors (2012)

3.79 Departmental GPA ; 3.69 Overall GPA

Research Experience

10X Genomics 9/2016 – 1/2017

**Consultant**

* Software development for single-cell VDJ product.

10X Genomics 6/2016 – 9/2016

**Computational Biology Intern**

* Research and development to enable co-assay for single-cell measurement of both gene expression profiles and VDJ sequences (T-cell and B-cell receptor sequences).
* Software developer for production and customer facing versions of CellRanger 1.2, a Python and Rust based distributed computing pipeline for 10X single-cell RNA-seq and VDJ datasets.
* Co-developed and deployed secondary analysis toolkit for 10X single-cell RNA-seq datasets in R.

Shendure Lab of University of Washington Department of Genome Sciences 9/2014 – Present

**Ph.D. Candidate**

* Computational and molecular biology methods development for:
* Simultaneous measurement of multiple data types from single-cell genomics datasets.
* Developing novel forward genetics paradigms.
* Examining regulation during cell-fate decisions using single-cell approaches.
* Contributed to development of a novel method for determining which cell types are contributing to cell-free DNA in blood plasma samples from cancer patients using changes in inferred nucleosome positioning.
* One of two main developers of an interactive data visualization tool utilizing D3.js and jQuery to explore spatiotemporal measurement of gene expression in developing *C. Elegans* embryos ([EPICViz](http://andrewjohnhill.com/projects/uw-genome-sciences/#epicviz)).

MacArthur Lab of Massachusetts General Hospital/Broad Institute of MIT and Harvard 9/2013 – 9/2015

**Research Assistant**

* Developed python API and scripts to extract/refine data from Leiden Open Variation Databases.
* Performed extensive automated validation of variants mapped from HGVS to VCF format.
* Developed methods for automating detection of misannotated protein coding exons in GENCODE using metrics for evolutionary conservation, constraint in a large reference panel, and tissue expression levels.
* Wrote software to detect and analyze multi-nucleotide polymorphisms derived from ~65K exome sequencing samples that change variant interpretation compared to individual variants.

**Tekscan, Inc.** 9/2012 – 1/2014

**Applications Engineer**

* Conducted engineering research projects for new applications of force and pressure sensors.
* Implemented algorithms to estimate shoe-size from noisy pressure sensor data.
* Signal processing and data analysis for IMU position/angle tracking of human gait.
* Improved algorithms for gait-analysis from Tekscan pressure sensor data.
* Developed automated test fixtures and data-analysis scripts with MATLAB and LabVIEW.
* Provided engineering support and/or training to customers and all internal departments.

**UW Biorobotics Lab (Professors Blake Hannaford and Howard Chizeck)** 1/2010 – 6/2012  
**Undergraduate Research Assistant**

* Thesis: Online Modeling of the *In Vivo* Mechanical Properties of Soft Tissue for Robotic Surgery
* Designed, built, and programmed electromechanical device to quantify in vivo tissue dynamics.
* Developed Unscented Kalman Filter/signal processing using MATLAB/C++.
* Co-developed hardware and microcontroller code for haptic-enabled glove.
* Developed hardware and microcontroller code to detect peg-contact in FLS block-transfer task.

Professor Joan Sanders Lab 8/2009 – 1/2010

**Undergraduate Research Assistant**

* Collected/analyzed data to calibrate tri-axis piezoelectric force sensor for amputee gait analysis.
* Designed and built Plexiglas housing for patient-mounted electronics.

Selected Awards and Honors

* National Science Foundation Graduate Research Fellowship (5 year fellowship, 3 years funding)
* Mary Gates Research Scholarship
* USA Gymnastics Men’s Program Scholarship
* Friends of Gymnastics Scholarship

Publications

* Xiaojie Qiu, **Andrew Hill**, Jonathan Packer, Dejun Lin, Yian Ma, Cole Trapnell. Single-cell mRNA quantification and differential analysis with Census. **In Press, Nature Methods.**
* Exome Aggregation Consortium, Monkol Lek, Konrad J Karczewski\*, Eric V Minikel\*, Kaitlin E Samocha\*, Eric Banks, Timothy Fennell, Anne H O'Donnell Luria, James S Ware, **Andrew J Hill**, Beryl B Cummings, Taru Tukiainen, Daniel P Birnbaum, Jack A Kosmicki, Laramie Duncan, Karol Estrada, Fengmei Zhao, James Zou, ***[54 additional authors]*,** Mark J Daly, Daniel G MacArthur. Combined analysis of protein-coding genetic variation in 60,706 humans. **August 2016**. **Nature**536,285–291.
* Matthew W Snyder\*, Martin Kircher\*, **Andrew J Hill**, Riza Daza, and Jay Shendure. Cell-free DNA Comprises an *In Vivo* Nucleosome Footprint that Informs Its Tissues-Of-Origin. **January 2016**. ***Cell***, *164*(1-2), 57–68.
* Xinxian Deng, Wenxiu Ma, Vijay Ramani, **Andrew Hill**, Fan Yang, Ferhat Ay, Joel B. Berletch, Carl Anthony Blau, Jay Shendure, Zhijun Duan, William S. Noble, and Christine M. Disteche. Bipartite structure of the inactive mouse X chromosome. **August 2015**. **Genome Biology**, 16:152.

Invited Talks

* **Andrew Hill**, Beryl Cummings, Konrad Karczewski, Monkol Lek, and Daniel MacArthur. “Phased annotation of protein-coding variants across 60,706 human exomes.” Presented at the 65th Annual Meeting of The American Society of Human Genetics, **October, 2015** in Baltimore, MD.
* **Andrew Hill**, Xiaojie Qiu, and Cole Trapnell. “Pseudotemporal ordering of cells undergoing immune stimulation and perturbations to cell-cell signaling.” Genome Training Grant Symposium invited trainee speaker. **July, 2015**.
* **Andrew Hill**. “Calibration and Synchronized Data Acquisition for High-Speed Applications.” Tekscan North American Distributor Meeting. Boston, MA. **April, 2013**.
* **Andrew Hill**, Sina Kosari, Blake Hannaford, and Howard Chizeck. “Online Modeling of the *In Vivo* Mechanical Properties of Soft Tissue for Robotic Surgery.” University of Washington Mary Gates Undergraduate Research Symposium. Seattle, WA. **May 2012**.

Selected Coursework

Data Visualization (CSE 512) Applied Biostatistics I/II (BIOSTAT 514/517) Linear algebra/Diff. Equations

Embedded Microcomputer Systems (EE 472) Digital Signal Processing Computational Molecular Biology

**Independent Coursework**

* Machine Learning Coursera (Stanford)
* Algorithms Design and Analysis – Part 1 Coursera (Stanford)
* Circuits and Electronics MIT Open Courseware
* Manual Machining and Layout (Mill and Lathe) Artisan’s Asylum

Skills

* **Computing:** Python, R, bash, Java, C#, MATLAB, LabVIEW
* **Web Development:** HTML, CSS, JavaScript, D3.js, React.js, jQuery
* **Operating Systems:** Mac OSX, Linux, Windows
* **Embedded Systems:** ARM and Arduino embedded system programming
* **Machining:** CNC mill, lathe, band-saw, drill-press, various hand tools

Coaching and Teaching Experience

Genome Sciences 361: Fundamentals of Genetics and Genomics Autumn 2016

**Teaching Assistant**

iD Tech Camps Summer 2012

**Summer Camp Instructor: Programming in Java and Adventures in Robotics**

**UW Bioengineering Department Circuitry Workshops** Winter 2012

**Volunteer Instructor**

UW Bioengineering Outreach Program 12/2011 – 6/2012

**Ultrasound Education Module Co-Developer and Instructor**

**United States Gymnastics Training Camps** Summers 2005 – 2010

**Counselor and Coach**

Leadership Experience and Activities

**Dana-Farber Cancer Institute, Brigham and Women’s Hospital** 9/2012 – 4/2013

**Volunteer – Kraft Family Blood Donor Center**

**UW Biomedical Engineering Society** 6/2011 – 6/2012

**Vice President and Webmaster**

**UW Honors Department** 9/2008 – 9/2009

**Peer Mentor**

**Washington Men’s Gymnastics Team** 8/2008 – 10/2009

***Team Member***

Study Abroad

* Creative Travel Writing and Sustainability in Ecuador Summer 2010