

Sean Maden

Bioinformatician; Quantitative Oncologist; Computational Biology Ph.D. Student

Contact/Connect

Phone:	<(425)-205-0659>
	<maden@ohsu.edu>; <maden.sean@gmail.com></maden.sean@gmail.com></maden@ohsu.edu>
GitHub:	<https: github.com="" metamaden=""></https:>

Education

Oregon Health & Science University, Department of Biomedical Engineering, Portland. OR August 2018 - May 2021 (expected)

Doctorate of Philosophy (Ph.D.), Computational Biology, advised by Prof. Abhi Nellore, in affiliation with PDXGX (Nellore & Thompson Labs); Dissertation on pan-tissue and disease epigenetics; Emphasis on design of server, auto-compilation, and analysis software for tens of thousands of methylation array samples; Investigating inter-experiment biases, novel normalization techniques, and holistic assay quality assessment across samples; Workflows for multi-assay integration; Application of machine learning for cancer biomarker discovery.

Reed College,

September 2007 - May 2011

Portland, OR

Bachelor of Arts (B.A.), Biology, emphases: population genetics, evolutionary ecology Rigorous liberal arts curriculum in the humanities and sciences; Population biology and statistical genetics research; Thesis characterizing genetic diversity declines from inbreeding in captive laboratory cichlid fishes (*A. burtoni*) with microsatellite molecular markers; Advisor Prof. Suzy Renn.

Selected Research Work and Experience

Research Data Analyst Assistant Fred Hutch, Seattle, WA

October 2015 – June 2018

Principal Investigators: Prof. William Grady, Dr. Ming Yu

Study of gastroesophageal cancer epigenetics; Application of preprocessing and analysis pipelines for methylation arrays and next-generation technologies; Research collaborations across labs, departments, and campuses; Coauthor and editor of original research manuscripts; Significant intellectual contributions to grant applications; Presented lectures, posters at conferences and lab meetings; Learned and applied R, Python, and related programming languages and software for data science and biostatistical analysis; Implemented organized lab repositories for big data.

Bioinformatics Research Intern

April - September 2015

Fred Hutch, Seattle, WA

Principal Investigator: Prof. William Grady; Postdoc project lead: Dr. Ming Yu

Carried out original analyses of cancer epigenetic data; Self-taught analytical programming; Presentation of findings and research discussion at lab meetings.

Laboratory Assistant

January – October 2013

Principle Investigator/Advisor: Prof. Gail Jarvik University of Washington Medical Genetics

Analyzed clinical population and epidemiologic data; Quantified gene-environment interactions with modeling and regression techniques; Programmed preprocessing workflows for next-generation genomics; Coauthored and edited published manuscript of findings (see Publications).

Undergraduate Thesis Researcher

September 2010 - May 2011

Advisor: Prof. Suzy Renn

Reed College, Biology Department

Quantified effects of laboratory breeding on genetic diversity in cichlid fishes (*A. burtoni*) using microsatellites; Authored thesis studying genetic diversity of lab cichlid fishes with microsatellites, and defended thesis before professorial board; Attained department grant funding (see Research Awards).

Scientific Memberships, Fellowships, and Funding

Grad Student Travel Stipend

May 2019

MOMACS Modeling the World's Systems 2019

Washington, DC

Conference travel stipend award, hosted by University of Pittsburgh Dept. of Computer Science.

Affiliate Member, Fred Hutch, Seattle, WA

June 2018 - present

Collaborators: Grady Lab; Luebeck Lab

Affiliate Member of Fred Hutch, with ongoing affiliation and collaboration with epigenetics labs in Clinical Research and Computational Biology divisions.

AAAS Member June 2018 - present

Affiliation: OHSU Graduate School

American Association for the Advancement of Science (AAAS), student member nominated by OHSU leadership.

2016 SAS-BWF Fellow, Fred Hutch, Seattle, WA

May 2016 - 2017

Advisors: Prof. Bill Grady, Dr. Ming Yu

Fellowship hosted and funded by SAS Institute and Burroughs Wellcome Fund. Contributed to ongoing research of epigenetics of colorectal cancer and biomarker discovery for clinical screening. Worked closely with JMP developers and recommended new software features based on real research needs.

AACR Associate Member, Fred Hutch, Seattle, WA

Jan 2016 - 2017

Recognized and supported by Association for Cancer Research (AACR) as promising early-career cancer investigator. AACR is one of the largest cancer research organizations in the US, and it hosts numerous annual meetings and workshops to help cancer researchers network and learn from colleagues.

Undergraduate Research Grant, Reed College, Portland, OR

Winter 2010

Advisor: Prof. Suzy Renn

Granted funding to extend research for undergraduate senior thesis studying genetic diversity of *A. burtoni* fishes (see Publications), awarded by Reed College Biology Department based on submission of original research overview.

Advisor: Prof. Robert Kaplan

Designed and conducted habitat field survey experiment of native at-risk frog *Rana aurora*, constructed breeding habitat and compiled a literature review of amphibian ecology. Authored and coauthored extensive reports presented as contributions to canon of Fisher Fellows past and present.

Peer-reviewed Publications, Preprints, and Works in Progress

- 1. Julianne K. David, **Sean K. Maden**, Benjamin R. Weeder, Reid F. Thompson, Abhi Nellore. "Cancer-specific" exon-exon junctions appear in embryological and other normal cells. 2019 (under review)
- 2. Ting Wang, **Sean K. Maden**, Georg Luebeck, Chris Li, Polly Newcomb, Cornelia Ulrich, Kelly Carter, Michael Luo, Ming Yu, William M. Grady. *Dysfunctional epigenetic aging of the normal colon in association with colorectal adenoma and cancer risk.* 2019 (under review)
- 3. Jenny Smith*, **Sean K. Maden***, David Lee*, Ronald Buie, Vikas Peddu, Ryan Shean, Ben Busby. *Consensus Machine Learning for Gene Target Selection in Pediatric AML Risk.* 2019 BioRxiv, 632166. *contributed equally (preprint)
- 4. Yuna Guo, Kelly Carter, Ming Yu, **Sean K. Maden**, Darwin Edmonds, Polly Newcomb P, Christopher Li, Neli Ulrich, William M. Grady. Senescence associated secreted factors are candidate drivers of the age related risk of colorectal cancer. 2018 (under review).
- 5. Georg E. Luebeck, William D. Hazelton, Kit Curtius, Sean K. Maden, Ming Yu, Kelly T. Carter, Wynn Burke, Paul D. Lampe, Christopher I. Li, Cornelia M. Ulrich, Polly A. Newcomb, Maria Westerhoff¹¹, Andrew M. Kaz, Yanxin Luo, John M. Inadomi, William M. Grady. *Implications of epigenetic drift in colorectal neoplasia*. 2018 Cancer Research (accepted for publication).
- 6. Ming Yu*, Sean K. Maden*, Matthew Stachler*, Andrew M. Kaz, Tai J. Heinzerling, Rachele M O'Leary, Xinsen Xu, Adam Bass, Amitabh Chak, Joseph E. Willis, Sanford D. Markowitz, William M. Grady. Subtypes of Barrett's Esophagus and Esophageal Adenocarcinoma Based on Genome-wide Methylation Analysis. 2017, Gut. *contributed equally.
- 7. Ludovic Barault, Alessio Amatu, Giulia Siravegna, Agostino Ponzetti, Sebastian Moran, Andrea Cassingena, Benedetta Mussolin, Chiara Falcomatà, Alexandra Binder, Carmen Cristiano, Daniele Oddo, Carlotta Cancelliere, Sara Bustreo, Katia Bencardino, **Sean Maden**, Alice Vanzati, Patrizia Zavattari, Mauro Truini, William M. Grady, Patrizia Racca, Karin B. Michels, Salvatore Siena, Manel Esteller, Alberto Bardelli, Andrea Sartore-Bianchi, Federica Di Nicolantonio. *Discovery of methylated circulating DNA biomarkers for comprehensive non-invasive monitoring of treatment response in metastatic colorectal cancer*. 2017 Gut; PMCID: PMC5897187.
- 8. Georg E. Luebeck, Kit Curtius, William D Hazelton, **Sean Maden**, Ming Yu, Prashanthi N Thota, Deepa T Patil, Amitabh Chak, Joseph E Willis, William M Grady. *Identification of a key role of widespread epigenetic drift in Barrett's esophagus and esophageal adenocarcinoma*. 2017 Clinical Epigenetics; PMCID: PMC5644061.
- 9. (Acknowledgement) Kit Curtius, Chao-Jen Wong, William D. Hazelton, Andrew M. Kaz, Amitabh Chak, Joseph E. Willis, William M. Grady, Georg E. Luebeck. A Molecular Clock Infers Heterogeneous Tissue Age Among Patients with Barrett's Esophagus. May 11, 2016 PLoS Comput Bio.; PMID: 27168458

- 10. Daniel Seung Kim, Sean K. Maden, Amber A Burt, Jane E Ranchalis, Clement E Furlong and Gail P Jarvik. Dietary fatty acid intake is associated with paraoxonase 1 activity in a cohortbased analysis of 1,548 subjects. 2013 Lipids in Health and Disease; PMCID: PMC3878825.
- 11. **Sean Maden**, Advisor: Suzy P Renn. *Observed Declines in Genetic Diversity Across Successive Generations of a Captive Astatotilapia burtoni Lineage, Using Microsatellite Molecular Markers*. Senior Thesis, 2011 Reed College.

Scientific Lay Press Works

1. (**Acknowledgement**) Anne-Sophie Kuhlman. "Esophageal adenocarcinoma: when DNA methylation informs the treatment". Fred Hutch Science Spotlight July 16, 2018.

Selected Bioinformatics Skills and Interests

Scientific software development

Bash and shell scripting;

Python and R statistical software libraries;

Tibbles, tidyverse, dply, Rstudior;

R modules, ggplot2, Gviz;

R-shiny apps, reactive programming;

GitHub, version control;

Nix and Windows operating systems;

Embarrassingly parallel processing;

TCGA, GEO, public genomics repos;

MySQL, MongoDB, database applications;

Statistical modeling and regression;

Machine learning:

ROC/AUC and predictive analyses;

Biomarker discovery;

Next-generation sequencing; Methylation array data mining;

Epigenetic regulation of expression;

Enhancers, insulators, super-enhancers;

Metadata mining and cleaning

Selected Science Presentations

- 1. (poster/abstract) Sean K. Maden, Kasper D. Hansen, Abhinov Nellore. *Comprehensive Reanalysis of DNA Methylation Array Data*. 2019 MOMACS Modeling the World's Systems Conference.
- 2. (poster) Mitra Barahimi, **Sean Maden**, Kelly Carter, Ming Yu, Kathy Vickers, Chris Li, William Grady. Sa1617-Discovery and Validation of Potential Field Cancerization Molecular Markers that Associate with Metachronous Polyp Formation. Gastroenterology 154 (6), S-331
- 3. (poster) Erik J. Snider, E. Georg Luebeck, Ming Yu, Kelly T. Carter, **Sean K. Maden**, Amitabh Chak, Joseph E. Willis, John M. Inadomi, Michael D. Saunders, Adam W. Templeton, William M. Grady. *Medication effects on methylation-related biological aging in Barrett's esophagus*. 2018. Gastroenterology 156 (6), S-509
- 4. (lecture) Sean Maden. "Recount Methylation: A generalizable platform for re-analysis of array-based epigenetic data". Biophysics Journal Club meeting, October 2018.
- 5. (poster) Mitra Barahimi, **Sean Maden**, Ming Yu, Kelly Carter, William Grady. *Discovery and Validation of Potential Field Cancerization Molecular Markers That Associate With Metachronous Polyp Formation*. Presented at 2018 Digestive Disease Week.
- 6. (lecture) **Sean Maden**. *Identification of Novel Molecular Characteristics of Methylation Subtypes in Esophageal Adenocarcinoma by Integrated Analysis*. Cancer Epigenetics Affinity Group meeting. Fred Hutch campus. Nov. 7, 2017.

- 7. (poster) Sean Maden* and Georg Luebeck*, Kit Curtius, William Hazelton, Ming Yu, Prashanti Thota, Deepa Patil, Amitabh Chak, Joseph Willis, William Grady *co-first authors. Scope and Significance of Epigenetic Drift in Barrett's Esophagus. Cancer Systems Biology Consortium (CSBC) Principal Investigator Meeting, Broad Institute MIT, Cambridge, MA. Oct 3, 2017.
- 8. (lecture) **Sean Maden**. Spatial and temporal epigenetic pattern gradients differentiate normal and progressed tissues in esophagus. Cancer Intervention and Surveillance Modeling Network (CISNET) 2017 meeting, Esophageal Cancer projects, June 6, 2017.
- 9. (poster) **Sean Maden***, Ming Yu*, Matthew Stachler*, Andrew M. Kaz, Tai J. Heinzerling, Rachele M O'Leary, Xinsen Xu, Adam Bass, Amitabh Chak, Joseph E. Willis, Sanford D. Markowitz, William M. Grady (*co-first authors). *Genome-wide methylation analysis reveals methylator subtypes of Barrett's esophagus and esophageal adenocarcinoma*. 2016 AACR Annual Meeting, Abstract: #3192, Session: DNA Methylation 1.

Graduate Level Science Courses Completed

Oregon Health and Science University, Ph.D. Computational Biology, Biomedical Engineering Summer 2018 – Spring 2021 (expected)

BMI 550/650: Bioinformatics and Computational Biology I: Algorithms

CS 545: Machine Learning

CONJ 650: Practice and Ethics of Science **BME 601:** Prequalifying PhD Research

University of Washington, M.Sc. Genetic Epidemiology, Public Health Genetics Fall 2012 - Fall 2013 (no degree obtained)

PHG 536: Bioinformatics and

Sequence Analysis

BIOSTAT 517/518: Applied

Biostatistics I and II

EPI 512/513: Epidemiological

Methods I and II

PHG 512: Law and Ethics in Public

Health Genetics

PHG 513: Pharmacogenetics and

Toxicogenomics

BIOST 580: Biostatistics Seminar **EPI 583:** Epidemiology Seminar

GENOME 525: Topics in Human Genetics

PHG 511: Genetic Epidemiology

PHG 521: Culture and Societal Genomics

PHG 523: Genetics and the Law

Undergraduate Level Science Courses Completed

Shoreline Community College, Biotechnology Lab Specialist Certificate Fall 2014 - Winter 2015 (no degree obtained)

BIOL 270: Molecular Biology BIOL 265/266: Solution and Media

BIOL 286: Molecular Diagnostics BIOL 275: Recombinant DNA

Reed College, B.A. Biology, Fall 2007 - Spring 2011

MATH 111: Calculus MATH 112: Intro to Analysis PHYS 100: General Physics I HIST 315: Medieval/Renaissance

Science and Religion

BIOL 332: Vascular Plant Diversity **BIOL 101/102:** Introductory Biology

CHEM 101/102: Introductory

Chemistry

BIOL 361: Genetics

BIOL 431: Seminar: Ecology and

Evolution of Plant-Human

Interactions

BIOL 358: Microbiology

BIOL 366: Population Ecology and

Evolution

CHEM 201/202: Organic Chemistry I

and II

PHIL 201: Logic

BIOL 351: Developmental Biology

(lecture only)

BIOL 431: Seminar: Chromosome

Structure

PHIL 316: Philosophy of Science

BIOL 431: Seminar: Gene Duplication/Overexpression

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