DANIEL A. SNELLINGS

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EDUCATION

Ph.D. Molecular Genetics and Microbiology 2017 - Present
Program in Cell and Molecular Biology
Duke University

B.S. Biochemistry and Molecular Biology 2013 - 2017
Pennsylvania State University

RESEARCH

The Role of Somatic Mutations in Vascular Malformations 2017 - Present
Douglas A Marchuk, Duke University
My work in the Marchuk Lab focuses on the genetic changes that lead to hereditary and sporadic neurovascular malformations. Specifically, I have shown that vascular malformations in Hereditary Hemorrhagic Telangiectasia follow a Knudsonian two-hit mechanism; and that cerebral cavernous malformations accumulate multiple synergistic somatic mutations which contribute to pathogenesis.

Environmental Factors Influencing Bumblebee Pigmentation Academic Year 2014 - 2017
Heather M Hines, Pennsylvania State University
In the Hines Lab I studied the mechanism of pigment biosynthesis and deposition in developing bumblebees. I also investigated the impact of foraging success and nutrient diversity on the pigment intensity of adult bees for potential use in the field as a bioindicator of nutritional fitness.

The Mechanism of Cement Production in Barnacles Summers 2015 - 2016
Christopher M Spillmann, Naval Research Laboratory
At the Naval Research Lab I worked with a group focused on understanding the biological mechanism of barnacle cement production and deposition with the ultimate goal of developing a hull coating which would prevent barnacle biofouling of naval vessels. Towards this end, I studied a previously undescribed tissue and helped characterize its role in barnacle development.

PUBLICATIONS

* Authors contributed equally

2021

Aileen A. Ren*, **Daniel A. Snellings***, Sophie Y. Su, Courtney C. Hong, Marco Castro, Alan T. Tang, Matthew R. Detter, Nicholas Hobson, Romuald Girard, Sharbel Romanos, Rhonda Lightle, Thomas Moore, Robert Shenkar, Christian Benavides, M. Makenzie Beaman, Helge Mueller-Fielitz, Mei Chen, Patricia Mericko, Jisheng Yang, Derek C. Sung, Michael T. Lawton, Michael Ruppert, Markus Schwaninger, Jakob Körbelin, Michael Potente, Issam A. Awad, Douglas A. Marchuk, and Mark L. Kahn. **Cerebral cavernomas arise through cancer-like PIK3CA and CCM mutations. *Nature*, in press, April 2021**

2019

Daniel A Snellings, Carol J Gallione, Dewi S Clark, Nicholas T Vozoris, Marie E Faughnan, and Douglas A Marchuk. **Somatic Mutations in Vascular Malformations of Hereditary Hemorrhagic Telangiectasia Result in Bi-allelic Loss of ENG or ACVRL1.** ***Am J Hum Genet*****, Oct 2019** Janne Koskimäki, Dongdong Zhang, Yan Li, Laleh Saadat, Thomas Moore, Rhonda Lightle, Sean P Polster, Julián Carrión-Penagos, Seán B Lyne, Hussein A Zeineddine, Changbin Shi, Robert Shenkar, Sharbel Romanos, Kenneth Avner, Abhinav Srinath, Le Shen, Matthew R Detter, **Daniel Snellings**, Ying Cao, Miguel A Lopez-Ramirez, Gregory Fonseca, Alan T Tang, Pieter Faber, Jorge Andrade, Mark Ginsberg, Mark L Kahn, Douglas A Marchuk, Romuald Girard, and Issam A Awad. **Transcriptome clarifies mechanisms of lesion genesis versus progression in models of Ccm3 cerebral cavernous malformations.** ***Acta Neuropathol Commun*****, 7(1):132, Aug 2019**

2018
Matthew R Detter, **Daniel A Snellings**, and Douglas A Marchuk. **Cerebral Cavernous Malformations Develop Through Clonal Expansion of Mutant Endothelial Cells.** ***Circ Res*****, 123(10):1143–1151, 10 2018** Chenyue Wang, Janna N Schultzhaus, Chris R Taitt, Dagmar H Leary, Lisa C Shriver-Lake, **Daniel Snellings**, Samantha Sturiale, Stella H North, Beatriz Orihuela, Daniel Rittschof, Kathryn J Wahl, and Christopher M Spillmann. **Characterization of longitudinal canal tissue in the acorn barnacle Amphibalanus amphitrite.** ***PLoS One*****, 13(12):e0208352, 2018**

SOFTWARE

gonomics (github.com/vertgenlab/gonomics) Role: Developer
A collection of genomics software tools written in Go (golang).
My work in gonomics focuses on developing a somatic variant caller that operates on sequencing data aligned to traditional linear references as well as data aligned to graph references as have been implemented in gonomics.

weaver (github.com/ddsnellings/weaver) Role: Creator & Developer
An open source toolkit for analyzing sequencing data generated by the Tapestri platform.
Currently available analysis pipelines for the Tapestri platform are proprietary and are only available through an AWS instance provided by Mission Bio. I have written several functions to analyze data I have generated with this platform. Though still very new, weaver is an outlet for me to formalize these functions for future use, and hopefully be a useful resource for the community.

FUNDING

F31 NIH/NHLBI (1F31HL152738-01) Role: PI April 2020 - March 2023
Investigating the Role of Somatic Mutations in Arteriovenous Malformations

SELECTED PRESENTATIONS

Invited Mission Bio Tapestri Webinar February 2021
Talk: "Multiple Somatic Mutations in a Single Clonal Population Drive CCM Pathogenesis"

American Society of Human Genetics 2020 Annual Meeting October 2020
Poser 1720: "A Novel Mutation in *GNAQ* Identified in Sturge-Weber Syndrome"

American Society of Human Genetics 2019 Annual Meeting October 2019
Flash Talk: "A Genetic Two-Hit Mechanism Drives Vascular Malformation in HHT"

American Society of Human Genetics 2019 Annual Meeting October 2019
Poster 1238/F: "A Genetic Two-Hit Mechanism Drives Vascular Malformation in HHT"

13th HHT International Scientific Conference June 2019
Talk: "HHT Telangiectases Contain Biallelic Mutations in *ENG* or *ACVRL1*"

OUTREACH

Undergraduate Career Development Panel October 2019
Served as a panelist detailing my path to graduate school and discussed career options with 1st year undergraduates.

The Great Insect Fair May 2016
Displayed samples and taught children about the importance of bumblebee coloration and the presence of color mimics in the wild.

MENTORSHIP

Jeff Reitano, Rotation Student 2021
Daichi Shonai, Rotation Student 2021
Makenzie Beaman, Rotation Student 2020
Taylor Anglen, Rotation Student 2020
Nicole Kastelic, Undergraduate Researcher 2019 - 2020
Makala Moore, Rotation Student 2019
Layne Clements, Undergraduate Summer Student 2018

PROFESSIONAL MEMBERSHIPS

American Society of Human Genetics (ASHG) 2019 - Present

American Heart Association (AHA) 2019 - Present

American Association for the Advancement of Science (AAAS) 2019 - Present

HONORS AND AWARDS

Reviewers Choice Abstract ASHG 2019 Annual Meeting October 2019

Best Scientific Oral Presentation 13th HHT International Scientific Conference June 2019

Molecular Genetics and Microbiology Travel Award Duke University April 2019

Eberly College of Science Research Award Pennsylvania State University November 2016

Apes Valentes Research Award Center for Pollinator Research, Penn State May 2015