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 mechanism of barnacle cement production and deposition with the ultimate goal of developing a hull coating which could 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

Daniel A. Snellings*, Courtney C. Hong*, Aileen A. Ren*, Miguel A. Lopez-Ramirez*, Romuald Girard*, Abhinav Srinath*, Douglas A. Marchuk, Mark H. Ginsberg, Issam A. Awad, and Mark L. Kahn. Cerebral cavernous malformation: from mechanism to therapy. *Circ Res*, in press, 2021

A. A. Ren*, **D. A. Snellings***, Y. S. Su, C. C. Hong, M. Castro, A. T. Tang, M. R. Detter, N. Hobson, R. Girard, S. Romanos, R. Lightle, T. Moore, R. Shenkar, C. Benavides, M. M. Beaman, H. Mueller-Fielitz, M. Chen, P. Mericko, J. Yang, D. C. Sung, M. T. Lawton, M. Ruppert, M. Schwaninger, J. Korbelin, M. Potente, I. A. Awad, D. A. Marchuk, and M. L. Kahn. PIK3CA and CCM mutations fuel cavernomas through a cancer-like mechanism. *Nature*, 2021

2019

D. A. Snellings, C. J. Gallione, D. S. Clark, N. T. Vozoris, M. E. Faughnan, and D. A. Marchuk. Somatic Mutations in Vascular Malformations of Hereditary Hemorrhagic Telangiectasia Result in Biallelic Loss of ENG or ACVRL1. *Am J Hum Genet*, 105(5):894–906, 2019

J. Koskimaki, D. Zhang, Y. Li, L. Saadat, T. Moore, R. Lightle, S. P. Polster, J. Carrion-Penagos, S. B. Lyne, H. A. Zeineddine, C. Shi, R. Shenkar, S. Romanos, K. Avner, A. Srinath, L. Shen, M. R. Detter, **D. Snellings**, Y. Cao, M. A. Lopez-Ramirez, G. Fonseca, A. T. Tang, P. Faber, J. Andrade, M. Ginsberg, M. L. Kahn, D. A. Marchuk, R. Girard, and I. A. Awad. Transcriptome clarifies mechanisms of lesion genesis versus progression in models of Ccm3 cerebral cavernous malformations. *Acta Neuropathol Commun*, 7(1):132, 2019

2018

M. R. Detter, **D. A. Snellings**, and D. A. Marchuk. Cerebral Cavernous Malformations Develop Through Clonal Expansion of Mutant Endothelial Cells. *Circ Res*, 123(10):1143–1151, 2018

C. Wang, J. N. Schultzhaus, C. R. Taitt, D. H. Leary, L. C. Shriver-Lake, **D. Snellings**, S. Sturiale, S. H. North, B. Orihuela, D. Rittschof, K. J. Wahl, and C. 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 genome graphs.

weaver (github.com/ddsnellings/weaver) Role: Creator & Developer
An open source toolkit for analyzing sequencing data generated by the Tapestri platform.

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"

Angioma Alliance 2020 Annual Scientific Meeting November 2020
Talk: "Biallelic Somatic Mutation of *KRIT1*, *CCM2*, and *PDCD10* in Sporadic CCMs"

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

REFEREES

Douglas Marchuk, PhD
James B. Duke Professor
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Email: douglas.marchuk@duke.edu Phone: (919) 684-1945

Craig Lowe, PhD
Assistant Professor
Department of Molecular Genetics and Microbiology, Duke University
Email: craig.lowe@duke.edu Phone: (919) 613-1754

Mark Kahn, MD
Edward S. Cooper, M.D./Norman Roosevelt and Elizabeth Meriwether McLure Professor
Department of Medicine, University of Pennsylvania
Email: markkahn@pennmedicine.upenn.edu Phone: (215) 898-9007