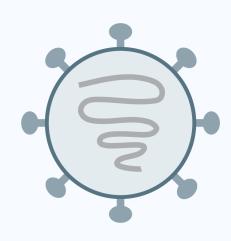
SARAH ARCOS

I am passionate about using computational methods to explore highthroughput biological datasets in a reproducible and rigorous way. I am a postdoctoral fellow in Dr. Adam Lauring's lab at the University of Michigan, where I use molecular biology, computational, and statistical toolboxes to study RNA virus evolution.



EDUCATION

2020 2015

PhD., Biochemistry

Vanderbilt University

Nashville, TN

 \cdot Studying RNA-protein interactions and RNA virus replication

2015 2011

B.S., Neurobiology, Cum laude

Georgetown University

• Washington, DC

· Studied developmental biology of the central nervous system



RESEARCH EXPERIENCE

Current 2021

Postdoctoral Fellow

Lauring Laboratory

University of Michigan

- · RNA virus evolution
- · Influenza A polymerase speed and fidelity
- · Epistatic interactions within the Influenza A polymerase complex

2020 2015

Graduate Researcher

Ascano Laboratory

- **♥** Vanderbilt University
- · RNA virus replication, innate immune evasion
- · Protein-RNA interactions
- · Post-transcriptional gene regulation in the innate immune system

2015 2012 HHMI Undergraduate Research Assistant

Silva Laboratory

- Georgetown University
- · Investigated the role and regulation of SOX14 during central nervous system development
- · Used the model organism Xenopus laevis

CONTACT

- sarcos@med.umich.edu
- **y** sarah_arcos
- github.com/saraharcos
- Saraharcos.github.io

LANGUAGE SKILLS

R		
C++		
Python		
Bash		

Made with the R package pagedown.

The source code is available on github.com/nstrayer/cv.

Last updated on 2023-03-02.



INDUSTRY EXPERIENCE

2018 2018

Infectious Disease/Microbiology Intern

Advisors: Dr. Bret Sellman, Dr. Taylor Cohen

MedImmune/Astra Zeneca

- · Antibody design and development
- · mtDNA haplogroup-dependent neutrophil responses to Staphylococcus aureus alpha-toxin
- · Neutrophil extracellular trap release



♣ TEACHING EXPERIENCE

Current 2019

R Fridays Founder

Vanderbilt Biochemistry Department

♥ Vanderbilt University

- · Leader of a peer group dedicated to discussion and problem solving related to biological data analysis in R
- · The R fridays blog is updated with interesting discussion topics from previous meetings

SERVICE POSITIONS

2018 2017 Vice President, Biochemistry Student's Association

♥ Vanderbilt University

· Coordinated the 2018 Biochemistry Department Annual Symposium



■ SELECTED PRESS (ABOUT)

2020

New method captures early viral-host protein interactions¹

VUMC Reporter

· Story of the VIR-CLASP method developed with Dr. Byungil Kim and Dr. Manny Ascano



■ SELECTED PRESS (BY)

2019

Dr. David Mitchell²

RNA Society

· Authored a spotlight on fellow RNA Society member Dr. David Mitchell

I believe that scientific reproducibility depends upon improved communication between data producers and data analyzers, and I am passionate about increasing data analysis accessibility for wet lab biologists.

PUBLICATIONS ELAVL1 primarily couples mRNA stability with the 3' UTRs of interferon 2021 stimulated genes3 Cell Reports

· Katie Rothamel, Sarah Arcos, Byungil Kim, Clara Reasoner, Neelanjan Mukherjee, and Manuel Ascano

Discovery of Widespread Host Protein Interactions with the Pre-2020 replicated Genome of CHIKV using VIR-CLASP

Molecular Cell

- · Byungil Kim*, Sarah Arcos*, Katherine Rothamel, Jeffrey Jian, Kristie L Rose, W Hayes McDonald, Yuqi Bian, Seth Reasoner, Nicholas J Barrows, Shelton Bradrick, Mariano A Garcia-Blanco, and Manuel Ascano. *Co-first authors
- Viral cross-linking and solid-phase purification enables discovery of 2020 ribonucleoprotein complexes on incoming RNA virus genomes4 Nature Protocols
 - · Byungil Kim*, Sarah Arcos*, Katherine Rothamel, and Manuel Ascano. *Co-first authors

SELECTED TALKS

Mutual information networks reveal evolutionary relationships within 2022 the Influenza A polymerase

American Society for Virology Annual Meeting

Madison, WI

Discovery of widespread host protein interactions with pre-replicated 2019 RNA virus genomes using VIR-CLASP

> EMBL Protein Synthesis and Translational Control Workshop • Heidelberg, Germany

N6-Methyladenosine-dependent regulation of the pre-replicated 2019 Chikungunya viral genome

Vanderbilt Biochemistry Department Annual Symposium

Nashville, TN

N6-Methyladenosine-dependent regulation of RNA during Chikungunya 2018 virus infection

RNA Society Annual Meeting

Parkeley, CA

The role and regulation of SOX14 in the development of the central 2014 nervous system in Xenopus laevis

> Georgetown- Howard Hughes Medical Institute Summer Research Symposium

> > • Washington, DC

T GRANTS AND AWARDS Molecular Mechanisms of Microbial Pathogenesis Post-doctoral 2022 **Training Grant** 2021 University of Michigan · T32AI007528, PI: Dr. Vern Carruthers Vanderbilt Russell G. Hamilton Graduate Leadership Institute Travel 2020 Grant Vanderbilt University · Used to attend rstudio::conf 2020 in San Francisco, CA 2019 **EMBL Advanced Training Center Corporate Partnership Travel Fellowship EMBL** Heidelberg \cdot Used to attend the 2019 EMBL Protein Synthesis and Translational Control Workshop National Institute of Allergy and Infectious Diseases Pre-doctoral 2018 **Training Grant** 2016 Vanderbilt University · T32AI11254, Pl: Dr. Eric Skaar **RNA Society Travel Grant** 2018 **RNA Society** · Used to attend the 2018 RNA Society Annual Meeting in Berkeley, CA Frank Chytil Travel Award 2018 Vanderbilt University \cdot Used to attend the 2018 RNA Society Annual Meeting in Berkeley, CA

Best Poster Award, Vanderbilt Biochemistry Department Annual

Chikungunya virus infection and innate immune activation

· Poster title: N6-Methyladenosine-dependent regulation of RNA during

PROFESSIONAL MEMBERSHIPS

American Society for Virology

RNA Society

Symposium

Vanderbilt University

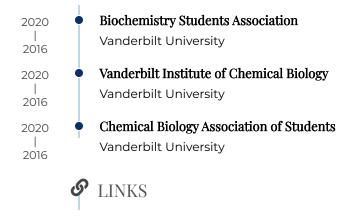
2018

Current

2021

2021

2018



- 1: https://news.vumc.org/2020/05/07/new method captures early viral host protein interactions/
- 2: https://www.rnasociety.org/dr david mitchell
- 3: https://www.biorxiv.org/content/10.1101/2020.08.24.263418v1
- 4: https://www.biorxiv.org/content/10.1101/2020.04.08.032441v1