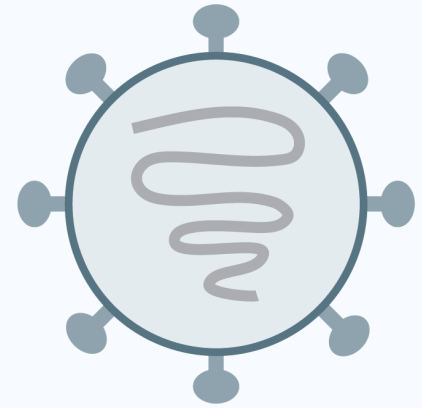




SARAH ARCOS




I am passionate about using computational methods to explore high-throughput biological datasets in a reproducible and rigorous way. I am a post-doctoral fellow in Dr. Adam Luring'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
 - 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

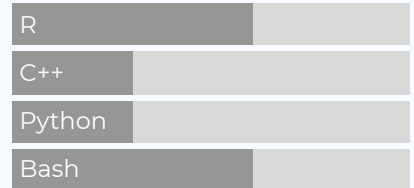
RESEARCH EXPERIENCE

- Current | 2021** **Postdoctoral Fellow**
Luring 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
 [sarah_arcos](https://twitter.com/sarah_arcos)
 github.com/saraharcos
 saraharcos.github.io

LANGUAGE SKILLS



Made with the R package
[pagedown](#).

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

Last updated on 2021-05-05.



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 interactions](#)

VUMC Reporter

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



SELECTED PRESS (BY)

2019



[Dr. David Mitchell](#)

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

- 2020 • **Discovery of Widespread Host Protein Interactions with the Pre-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
- 2020 • **Viral cross-linking and solid-phase purification enables discovery of ribonucleoprotein complexes on incoming RNA virus genomes**
Nature Protocols
• Byungil Kim*, Sarah Arcos*, Katherine Rothamel, and Manuel Ascano. *Co-first authors
- 2020 • **ELAVL1 primarily couples mRNA stability with the 3' UTRs of interferon stimulated genes**
Cell Reports (in press)
• Katie Rothamel, Sarah Arcos, Byungil Kim, Clara Reasoner, Neelanjan Mukherjee, and Manuel Ascano



SELECTED TALKS

- 2019 • **Discovery of widespread host protein interactions with pre-replicated RNA virus genomes using VIR-CLASP**
EMBL Protein Synthesis and Translational Control Workshop
📍 Heidelberg, Germany
- 2019 • **N6-Methyladenosine-dependent regulation of the pre-replicated Chikungunya viral genome**
Vanderbilt Biochemistry Department Annual Symposium
📍 Nashville, TN
- 2018 • **N6-Methyladenosine-dependent regulation of RNA during Chikungunya virus infection**
RNA Society Annual Meeting
📍 Berkeley, CA
- 2014 • **The role and regulation of SOX14 in the development of the central nervous system in *Xenopus laevis***
Georgetown- Howard Hughes Medical Institute Summer Research Symposium
📍 Washington, DC



GRANTS AND AWARDS

- 2020 • **Vanderbilt Russell G. Hamilton Graduate Leadership Institute Travel Grant**
Vanderbilt University
• Used to attend rstudio::conf 2020 in San Francisco, CA

- 2019
|
2016

●

EMBL Advanced Training Center Corporate Partnership Travel Fellowship
 EMBL Heidelberg

- Used to attend the 2019 EMBL Protein Synthesis and Translational Control Workshop
- 2018
|
2016

●

National Institute of Allergy and Infectious Diseases Pre-doctoral Training Grant
 Vanderbilt University

- T32AI11254, PI: Dr. Eric Skaar
- 2018

●

RNA Society Travel Grant
 RNA Society

- Used to attend the 2018 RNA Society Annual Meeting in Berkeley, CA
- 2018

●

Frank Chytil Travel Award
 Vanderbilt University

- Used to attend the 2018 RNA Society Annual Meeting in Berkeley, CA
- 2018

●

Best Poster Award, Vanderbilt Biochemistry Department Annual Symposium
 Vanderbilt University

- Poster title: N6-Methyladenosine-dependent regulation of RNA during Chikungunya virus infection and innate immune activation



PROFESSIONAL MEMBERSHIPS

- Current
|
2018

●

RNA Society
- Current
|
2016

●

Biochemistry Students Association
 Vanderbilt University
- Current
|
2016

●

Vanderbilt Institute of Chemical Biology
 Vanderbilt University
- Current
|
2016

●

Chemical Biology Association of Students
 Vanderbilt University