

Courtney R Armour PhD

COMPUTATIONAL BIOLOGIST

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Summary.

I'm a computational biologist specializing in statistical analysis of complex datasets. I am experienced with multiple programming languages as well as a variety of data exploration and visualization techniques. I'm passionate about open science and reproducible research. I'm seeking the opportunity to work on cutting edge research at the frontier of human health.

Skills and Expertise _____

Research	Computational	Laboratory
Machine learning – Data visualization Gut microbiome – Metagenomics Amplicon sequencing – Statistical analysis – Network analysis Reproducible research	R – R Markdown – Quarto Git/GitHub – Snakemake Conda/mamba – Bash Slurm – Perl – Python	DNA extraction – Library Preparation – Cell Culture – Cloning – Gel electrophoresis – Western blot – PCR

m Education

PhD Molecular and Cellular Biology

DEPTARTMENT OF MICROBIOLOGY

• Advisor: Thomas Sharpton

BS Biological Sciences

COLLEGE OF LIBERAL ARTS AND SCIENCES

• Minor: Mathematics

Oregon State University

2014-2020

Arizona State Univeristy

2008-2012

<u>IIII</u> Experience

Postdoctoral Research Fellow (remote)

SCHLOSS LAB

Univeristy of Michigan

2020-present

- · Optimizing machine learning models to predict colorectal cancer based on gut microbiome taxonomic abundace
- Identifying early markers of colorectal cancer development in a large clinical cohort of individuals with genetic predisposition to cancer
- Developing curriculum and teaching computational skills to coding beginners in virtual workshops

Graduate Research Assistant

Oregon State University

SHARPTON LAB

2014-2020

- Metagenomic analysis of gut microbiom samples to quantify associations with health
- Assembly of metagenomic data from gut microbiome samples to build integrated gene catalogues of microbiome genomic diversity
- Collaborate with a diverse group to develop methodology for analyzing analyzing microbiome data

Research/Laboratory Assistant

University of Arizona

2010-2013

DR. PAUL BOEHMER AND DR. JUI-CHENG HSIEH

- Creation of HSV-1 viral mutants to quantify impact on replication
- · Mentor new members of the lab
- Maintain tissue culture and order supplies

Publications_

- 1. **Courtney R. Armour**, Kelly L. Sovacool, William L. Close, Beg<U+00FC>m D. Top<U+00E7>uo<U+011F>lu, Jenna Wiens, Patrick D. Schloss. (2022). Streamlined implementation of a machine learning model to classify screen relevant neoplasia using reference-based OTU clustering. *bioRxiv preprint*. doi: 10.1101/2022.09.01.506299
- 2. **Courtney R. Armour**, Beg<U+00FC>m D. Top<U+00E7>uo<U+011F>lu, Andrea Garretto, Patrick D. Schloss. (2022). A goldilocks principle for the gut microbiome: taxonomic resolution matters for microbiome-based classification of colorectal cancer. mBio. doi: 10.1128/mbio.03161-21
- 3. Zena Lapp, Kelly L. Sovacool, Nick Lesniak, Dana King, Catherine Barnier, Matthew Flickinger, Jule Kr<U+00FC>ger, Courtney R. Armour, Maya M. Lapp, Jason Tallant, Rucheng Diao, Morgan Oneka, Sarah Tomkovich, Jacqueline Moltzau Anderson, Sarah K. Lucas, Patrick D. Schloss (2022). Developing and deploying an integrated workshop curriculum teaching computational skills for reproducible research Journal of Open Source Education. doi: 10.21105/jose.00144
- 4. Christopher A. Gaulke, Laura M. Beaver, **Courtney R. Armour**, Ian R. Humphreys, Carrie L. Barton, Robyn L. Tanguay, Emily Ho, Thomas J. Sharpton (2020). An integrated gene catalog of the zebrafish gut microbiome reveals significant homology with mammalian microbiomes *bioRxiv preprint*. doi: 10.1101/2020.06.15.153924
- 5. Christopher A. Gaulke*, **Courtney R. Armour***, Ian R. Humphreys, Laura M. Beaver, Carrie L. Barton, Lucia Carbone, Emily Ho, Robyn L. Tanguay, Yuan Jiang, Thomas Sharpton" (2020). Interspecies comparative metagenomics reveals correlated gut microbiome functional capacities among vertebrates *bioRxiv preprint*. doi: 10.1101/2020.06.15.153320
- 6. Rufa L Mendez, Cristobal Miranda, **Courtney R Armour**, Thomas J Sharpton, Jan Frederik Stevens, Jung Yeon Kwon (2020). Supplementation with sea vegetables textsuperscript*Palmaria mollistextsuperscript* and textsuperscript*Undaria pinnatifidatextsuperscript* exerts metabolic benefits in diet-induced obesity in mice Current Developments in Nutrition. doi: 10.1093/cdn/nzaa072
- 7. Duo Jiang, **Courtney R. Armour**, Chenxiao Hu, Meng Mei, Chuan Tian, Thomas J. Sharpton, Yuan Jiang (2019). Microbiome Multi-Omics Network Analysis: Statistical Considerations, Limitations, and Opportunities Frontiers in Genetics. doi: 10.3389/fgene.2019.00995
- 8. **Courtney R. Armour**, Stephen Nayfach, Katherine S. Pollard, Thomas J. Sharpton (2019). A metagenomic meta-analysis reveals functional signatures of health and disease in the human gut microbiome mSystems. doi: 10.1128/mSystems.00332-18
- 9. Thomas Sharpton, Svetlana Lyalina, Julie Luong, Joey Pham, Emily M. Deal, **Courtney R. Armour**, Christopher Gaulke, Shomyseh Sanjabi, Katherine S. Pollard (2017). Development of Inflammatory Bowel Disease Is pathed to a Longitudinal Restructuring of the Gut Metagenome in Mice mSystems. doi: 10.1128/mSystems.00036-17
- 10. Jui-Cheng Hsieh, Ryan Kuta, **Courtney R. Armour**, Paul E. Boehmer (2014). Identification of two novel functional p53 responsive elements in the herpes simplex virus-1 genome Virology. doi: 10.1016/j.virol.2014.04.019

Presentations

Talks

Comparative metagenomic investigations link the functional capacity of the gut microbiome to vertebrate physiology

DISSERTATION DEFENSE

The functional diversity of the gut microbiome in association with host physiology MICROBIOLOGY AND IMMUNOLOGY SEMINAR

The power of poop: diagnostic potential of the gut microbiome in human disease Center for Genome Research and Biocomputing Spring Conference

A metagenomic meta-analysis reveals functional signatures of health and disease in the human gut microbiome

MICROBIOLOGY SEMINAR SERIES

A metagenomic meta-analysis reveals functional signatures of health and disease in the human gut microbiome

INTERNATIONAL SYMBIOSIS SOCIETY.

Interaction between herpes simplex virus-1 DNA polymerase and uracil-DNA glycosylase. awarded best undergraduate presentation

BASIC MEDICAL SCIENCES RESEARCH RETREAT

Oregon State University

MAR 2020

University of Michigan

FEB 2020

Oregon State University

APR 2019

Oregon State University

NOV 2018

11012010

Oregon State University

JUL 2018

University of Arizona

JUN 2013

Interaction between herpes simplex virus-1 DNA polymerase and uracil- DNA glycosylase

BASIC MEDICAL SCIENCES INTRA-DEPARTMENTAL SEMINAR SERIES

University of Arizona
MAR 2013

Posters

Modeling the functional variation of the gut microbiome across vertebrates

ASM MICROR

San Francisco CA JUN 2019

A metagenomic meta-analysis reveals functional signatures of health and disease in the

human gut microbiome

Lake Arrowhead CA

LAKE ARROWHEAD MICROBIAL GENOMICS CONFERENCE

SEP 2018

A Metagenomic Meta-Analysis Reveals Functional Signatures of Health and Disease in the

Human Gut Microbiome. awarded best graduate student poster

Oregon State University

FALL CENTER FOR GENOME RESEARCH AND BIOCOMPUTING CONFERENCES

OCT 2018

Large-scale analysis of human gut metagenomes reveals functional indicators of disease in the gut microbiome

Pasco WA AUG 2017

Integrating clinical data reveals functional indicators of disease in the gut microbiome

EMSL/PNNL MULTI-OMICS FOR MICROBIOMES CONFERENCE

Oregon State University

OSU MICROBIOME INITIATIVE FORUM

MAY 2017

Integrating clinical metagenomic data reveals microbiome signatures of dysbiosis

University of Oregon

META CENTER SYMPOSIUM

University of Oregon

An automated workflow for the quality control and functional analysis of host-associated metagenomes

META CENTER SYMPOSIUM

AUG 2015

AUG 2016

Trellowships

Larry W. Martin and Joyce B. O'Neill Fellowship

\$25,000

Oregon State University

2018 - 2019

Nicholas R. Tartar Graduate Student Fellowship

\$6,671

Oregon State University

Winter 2018

Pacific Northwest National Lab Travel Grant

EMSL/PNNL Multi-omics for Microbiomes Conference

COST OF CONFERENCE REGISTRATION

AUG 2017

ASM SIGHPC and Intel Data and Computational Science Fellowship

Oregon State University

\$15,000 PER YEAR - \$60,000 TOTAL

2016 - 2020

President's Scholarship

Arizona State University

\$9,000 PER YEAR - \$36,000 TOTAL

2008 - 2012

Specialized Training _____

Deep Learning for Life Scientists

Oregon State University

2019

Specialized workshop courses

CENTER FOR GENOME RESEARCH AND BIOCOMPUTING

CENTER FOR GENOME RESEARCH AND BIOCOMPUTING

Oregon State University

2014 - 2019

• RNAseq I and II

· Computing in Life Sciences - Perl

• Python I and II

• Command-line Data Analysis

• Introduction to Unix and Linux

Certificate in Applied Biostatistics

University of Washington

DEPARTMENT OF BIOSTATISTICS

2014



Teaching

Teaching Assistant (ST599) *Introduction to Quantitative Genomics*Oregon State University

DEPARTMENT OF STATISTICS Fall 2016

Teaching Assistant (BI211) *Principles of Biology for Life Science Majors*Oregon State University

DEPARTMENT OF STATISTICS

Fall 2016

Mentorship

Austin Hammer (PhD Student)

Oregon State University

DEPARTMENT OF MICROBIOLOGY 2019 - 2020

Ian Humphreys (Accelerated Masters Student)Oregon State University

DEPARTMENT OF MICROBIOLOGY 2017 - 2019

Ryan Kuta (Undergraduate Student)University of Arizona

DEPARTMENT OF BASIC MEDICAL SCIENCES 2012 - 2013