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245 S Chester St. Baltimore, MD 21231

Personal Summary

I am a published molecular virologist with experience investigating respiratory pathogen kinetics and evolution to investigate the arms-race between viruses and the nasal mucosal barrier immune system. At current, I am a 3rd year PhD Candidate at the Johns Hopkins Bloomberg School of Public Health within the Department of Molecular Microbiology and Immunology under the supervision of <u>Dr. Andy Pekosz</u>. Leveraging my graduate studies, I will continue to contribute computational tools and robust evidence to the field of virology in the field of government and/or industry.

Education

Johns Hopkins Bloomberg School of Public Health PhD in Molecular Microbiology and Immunology (In progress)	Sept. 2021 – Current
University of Tennessee, Knoxville Bachelor of Science in Microbiology	August 2015 - May 2019
Awards and Fellowships	
Johns Hopkins T32 Grant Recipient	July 2023 – July 2025
Johns Hopkins University – Department of Molecular Microbiology and Immunology	
Oak Ridge Institute for Science and Education (ORISE) Fellowship Walter Reed Army Institute of Research, Silver Spring, MD	Sept. 2019 – Aug. 2021

Certifications

Coursera – Genomic Data Science Specialization through Johns Hopkins Coursera – R Specialization

Computational Expertise

- Development of software such as <u>pyFLUte</u> for streamlining at-scale influenza genomic surveillance.
- Molecular clock modeling of viral, parasitic and bacterial genomic evolution.
- Strong background in R, Python, Perl, Java, Java Script and Linux-based tools.
- Implementation of bioinformatic pipelines within UNIX environments.
- Competent with MiSeq, HiSeq, and ONT library preparation/QC, data generation and variant analysis pipelines
- Proficient streamlined data cleaning using R and Python tools.
- Proficient in advanced Bulk and scRNAseq analysis using R and Python-based tools.
- Interactive geospatial modeling of infectious disease data
- Efficient with large -omics data visualization

Wet lab Expertise

- Patient-derived Air-Liquid Interface (ALI) Epithelial Cell Culture Expansion and Differentiation
- Transgenic stem cell product development through lentiviral transduction systems for ALI culture systems
- Reverse Genetics of Orthomyxoviruses (Influenza B)
- Flow Cytometry FACS Caliber, LSR II, MacsQuant and BD Symphony Systems
- Luminex Multiplex Serology Analysis (Magpix Platforms)
- Cell/tissue culture for: microbial, mammalian, and parasitic cell lines using aseptic technique
- Protein Extraction, isolation, Western Blotting.
- Serology: ELISA and agglutination testing in various parasite models
- PCR, PCR-RFLP, Multiplex PCR, qPCR, qRT-PCR, and HRM for pathogen detection, quantification, and genotyping
- Multi-Dye Flow Cytometry on MACS Quant systems
- Cloning: Gibson Assembly, Restriction Digest, TOPO, Golden Gate
- CRISPR Library Screening in MMEJ repair systems
- Antibiotic and Fungicide Sensitivity Testing
- BL-1 and BL-2 Certifications relevant to cell culture and gene editing of blood borne pathogens.
- Environmental ATP Testing/quantitative analysis

Grants and Awards

- Fall 2023: The A. Ralph and Sylvia E. Barr Fellowship in Vector Biology
- Fall 2022: The A. Ralph and Sylvia E. Barr Fellowship in Vector Biology
- Fall 2021: The A. Ralph and Sylvia E. Barr Fellowship in Vector Biology
- Summer 2018: University of Tennessee Research IMPACT Independent Research Funding
- 2015 2019: Tennessee HOPE Scholarship recipient
- Summer 2015: Eagle Scout Recipient

Implemented CRISPR/Cas9 editing protocols of P. falciparum SNP targets investigating PfCRT resistance to Chloroquine in CHMI workflows.

Supported active clinical trial as diagnostic hub for malarial infections. Maintained routine cell-line culture of parasites for various experiments

Dr. Andy Pekosz Lab - Johns Hopkins Bloomberg School of Medicine

Walter Reed Army Institute of Research - Silver Spring, Maryland

Genetics and Parasite Biology Malarial Biologics Branch

Thesis Project: "The Evolutionary Dynamics of Influenza B at the Nasal Barrier"

Worked routinely with clinicians, scientists, software engineers across multiple departments within a government-restricted environment.

University of Tennessee: Hansen Laboratory, Knoxville, TN

Department of Etymology and Plant Pathology

Undergraduate Laboratory Manager

Research Experience

PhD Candidate

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ORISE Fellow

Malaria Infections (CHMI)"

from endemic regions.

Project: "Investigations into azoxystrobin fungicide sensitivity dynamics in the frogeve leaf spot Cercospora nicotianae"

- Fungal Cell Culture Isolated unknown plant pathogens on selective or nutrient rich media in a BSL-1 setting
- Molecular Biology Performed various methods of PCR to identify unknown plant pathogens in fungal and bacterial species
- Fungicide Resistance Investigated the level of azoxystrobin fungicide sensitivity of Cercospora nicotianae through spore germination microscopy
- Assisted in training for 1 graduate and 2 undergraduate students in basic laboratory techniques: pathogen isolation, sterile technique, PCR, Media Preparation, basic bioinformatics, and agricultural extension protocols.

University of Tennessee: Su laboratory, Knoxville TN

Department of Microbiology - Population Genetics, BSL-1

Laboratory Technician and Research Assistant

- Developed a computational GIS-based spatial tracking documented application for of Τ. genotypes gondii (https://myutk.maps.arcgis.com/apps/webappyiewer/index.html?id=032057aa0bc84cbcaa2c8e18c6fd58c8)
- Delivered a step-by-step protocol for data entry and computational utilization of WebGIS platform.
- Parasite Cell Culture Isolated DNA from muscle, brain, liver and spleens of T. gondii infected mammals to be screened for antibody titers.
- Prepared Mammalian muscle and brain tissue for antigen testing and microscopy.
- Parasite Propagation Cultured and Managed T. gondii cell lines in select human fibroblast and liver cells.
- Serotyping Performed serological testing on mammalians tissue samples via MAT immunological testing.
- **Molecular Biology** Utilized PCR-RFLP to target and characterized over 100+ unique *T. gondii* genotypes.
- Computational Biology Compiled sequences and curated phylogenetic trees for genomic characterization of T. gondii.

UT Institute of Agriculture: Stewart Laboratory, Knoxville, TN

Department of Plant Sciences - Synthetic biology, BSL-1

Undergraduate Research Assistant Advisor-Reggie Millwood

- Develop A database for molecular clones by Stewart Lab scientists utilizing Microsoft Excel and Access •
- Performed PCR-based screening on bacterial clone isolates and software debugging for database development.
- Shadowed Postdoctoral and Graduate staff working on crop genetics, chloroplast engineering, and CRISPR/Cas platforms. .
- Trained and Assisted -several undergraduate students in molecular techniques and general laboratory maintenance.

UT Institute of Agriculture: Stewart Laboratory, Knoxville, TN

Department of Plant Sciences - Synthetic biology, BSL-1

Undergraduate Research Assistant

- Bacterial Cell Culture Responsible for plate and liquid cultures of vector-carrying isolates in LB and YEP media.
- Explored novel methods of protein-induced transcriptional regulation in Nicotiana tabacum through targeted genomic techniques utilizing Agrobacterium tumefaciens.
- Fluorescent/ UV Microscopy- Utilization of fluorescent protein spectroscopy technology for gene dosage, detection, and translational quantification.
- Bacterial Vector Constriction Utilized TOPO and Gibson assembly techniques to construct vectors appropriate for JM109 and EHA 105 strains of E. coli and A. tumefaciens respectively. Primarily worked with GFP/ RFP combination reporters for translation efficiency screening.
- protein inducible promoter in plant systems.
- Statistical Analysis Utilization of R and SAS platforms for analysis of experimental data.
- . Plant Horticulture — In tobacco and soybean model systems.

Team lead on Influenza B surveillance, genome and phenotypic characterization Development of novel transgenic primary cell cultures derived for respiratory virus characterization

- Rescue of isogenic influenza B viruses for targeted genomic determinants of viral and host phenotypes.

Facilitated implementation of the department's first high-performance computing platform for GWAS analysis.

Development of bioinformatic and database pipelines for Johns Hopkins Hospital collaborative seasonal influenza A and B surveillance.

Project: "Developing a Suite of Geographically and Genetically Diverse Human Malaria Parasite Strains to Enable Heterologous Controlled Human

Utilized next generation sequencing for whole genome and amplicon-based genotyping and drug resistance profiles of Plasmodium field isolates

Designed and optimized SOPs for antigenic diversity of Plasmodium falciparum field isolates for detection vaccine and surveillance studies.

September 2019 - August 2021

May 2018 - July 2019

Jan. 2019 - May 2019

May 2017- Aug 2017

May 2016 - Aug. 2018

August 2021-Present

Volunteer and Leadership Experience

UT Crew Club – Knoxville, TN Secretary	Aug. 2015-May 2018
 Utilized Excel skills to track volunteer hours, organization Designed core training programs for both the novice and varsity teams 	
East Tennessee Children's Hospital – Knoxville, TN Emergency Department Volunteer • Accumulated over 100 hours of volunteer work assisting clinicians through	Aug. 2017 – May2019
Ask a Scientist STEAM Community engagement	Jan 2018 – Jan 2019
Microbiology Undergraduate Club journal and education outreach club	Aug 2016 – May 2019 Microbiology

Volunteer Experience

Logged over 100+ total service hours in and around the Knoxville community over the course of four years

 Achieved recognition from the University of Tennessee Center for Leadership and Service for participation in more than 100 hours of volunteer service

Leadership Experience

2018	Writer - Ask A Scientist, The University of Tennessee - Knoxville 2018 - current
2018	Photography Media Manager - Imagine UT, The University of Tennessee – Knoxville, 2018
2018	Secretary - Microbiology Undergraduate. Club - The University of Tennessee – Knoxville, 2018
2018	Ambassador – The Division of Biology, The University of Tennessee
2016	TEDxUTK – Logistics Team Leader – University of Tennessee, 2016-2018
2016	Secretary - Tennessee Crew Club - Knoxville, 2016-2017 term.

Publications *co-first authored work

*Akin E, *Wilson JL, Zhou R, Jedlicka A, Dziedzic A, Liu H, Fenstermacher KZ, Rothman R, Pekosz A. The Influenza B Virus Victoria and Yamagata Lineages Display Distinct Cell Tropism and Infection Induced Host Gene Expression in Human Nasal Epithelial Cell Cultures. bioRxiv [Preprint]. 2023 Aug 4:2023.08.04.551980. doi: 10.1101/2023.08.04.551980. PMID: 37577630; PMCID: PMC10418153.

Creisher PS, Perry JL, Zhong W, Lei J, Mulka KR, Ryan H, Zhou R, **Akin EH**, Liu A, Mitzner W, Burd I, Pekosz A, Klein SL. Adverse outcomes in SARS-CoV-2 infected pregnant mice are gestational age-dependent and resolve with antiviral treatment. bioRxiv [Preprint]. 2023 Mar 23:2023.03.23.533961. doi: 10.1101/2023.03.23.533961. PMID: 36993658; PMCID: PMC10055386.

Robben PM, **Akin EH**, Dunbar CR, Pichugin A, Regules JA. Late-presenting Plasmodium falciparum Malaria in a Non-Endemic Setting During COVID-19 Travel Restrictions. Mil Med. 2023 May 16;188(5-6):e1335-e1337. doi: 10.1093/milmed/usab393. PMID: 34557926; PMCID: PMC8500131.

(expected) Akin, E., Kamau, E. Pichugin, A. et al., 2024 "Down Selection of 4 Kenyan Plasmodium falciparum strains for Heterologous non-3D7 CHMI challenge", Walter Reed Army Institute of Research, Malaria Biologics Branch, Silver Spring, MD.

Presentations and Abstracts *Accomplished During PhD Work

- *Akin E, 2023. <u>Seminar Talk</u>. Centers of Excellence for Influenza Research and Response (CEIRR). Title: Distinct Transcriptional Landscapes of Influenza B are Independent of Viral Load in Infected Human Nasal Epithelial Cells. Johns Hopkins Bloomberg School of Public Health. Baltimore, MD.
- *Akin E, 2023. <u>Seminar Talk</u>, American Society of Virology Conference. Title: Distinct Transcriptional Landscapes of Influenza B are Independent of Viral Load in Infected Human Nasal Epithelial Cells. Johns Hopkins Bloomberg School of Public Health. Athens, GA
- *Akin E., 2022. Poster. American Society of Virology Conference. Title: **Evolutionary Dynamics of Influenza B: Characterizing** Lineage Determinants of Host Response at the Nasal Barrier. Johns Hopkins Bloomberg School of Public Health. Madison. WI
- *Akin E., 2022. Flash Talk. Centers of Excellence for Influenza Research and Response (CEIRR) Annual Meeting. Title: **Evolutionary Dynamics of Influenza B: Characterizing Lineage Determinants of Host Response at the Nasal Barrier.** Johns Hopkins Bloomberg School of Public Health. Memphis, TN

- *Akin E., 2022 Poster: Centers of Excellence for Influenza Research and Response (CEIRR) Annual Meeting Title: Evolutionary Dynamics of Influenza B: Characterizing Lineage Determinants of Host Response at the Nasal Barrier. Johns Hopkins Bloomberg School of Public Health. Memphis, TN
- Akin E., 2020. Department Talk. Drug Resistance Mechanisms in P. falciparum: Chloroquine and other 4-aminoquinalones and to how diminish it through CRISPR/Cas9 mediated mutagenesis. Malaria Biologics Branch. Silver Spring, MD
- Akin E., 2020. Department Presentation. Whole genome analysis of down-selected parasites against recorded Pf3K database genotypes. Malaria Biologics Branch. Silver Spring, MD
- Akin E., 2020. Department Presentation. Targeted Analysis of Circumsporozoite Protein in *Plasmodium ovale* subspecies reveal novel haplotypes of central repeat region. Malaria Biologics Branch. Silver Spring, MD
- Akin E., 2020. Department Presentation. Application of machine learning in estimating genetic relatedness in Plasmodium field isolates. Malaria Biologics Branch. Silver Spring, MD
- Akin E., 2020. Department Presentation. Down selection of Kenyan Plasmodium falciparum field strains using microsatellite and targeted sequencing analysis for heterologous CHMI. Malaria Biologics Branch. Silver Spring, MD
- Akin E., 2019. Keystone Presentation. Continued investigations into azoxystrobin fungicide sensitivity dynamics in the frogeye leaf spot pathogen, Undergraduate Research and Creative Achievement. Knoxville, TN
- Akin et. al, 2018. The Q-System: A New Technology to Regulate Transgene Expression in Plants. Abstract.
- Akin, E., 2018. The Q system: A new transcriptional technology to regulate gene expression in plants. The Exhibition of Undergraduate Research and Creative Achievement. Knoxville, TN
- Akin, E., 2018. The Q system: A new transcriptional technology to regulate gene expression in plants. Annual Student Conference of current College of Agriculture and Natural Science work. Knoxville, TN
- Akin, E., 2017. The Q system: A protein- based transcriptional regulation module for gene expression in plants. College of Agriculture and Natural Science Alpha Gamma Rho Honors Society Induction. Knoxville, TN.