

# Michael J. Akinyemi

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## RESEARCH INTERESTS

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Computational Biology, Immune Engineering, Systems Biology/Immunology, Molecular Biophysics, Autoimmune Disorders, Probabilistic Algorithms, Biostatistics

## EDUCATION

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### University of Central Florida

Biomedical Sciences (B.S.)

Computer Science (Minor)

Genomics & Bioinformatics (Minor)

- Ronald E. McNair Scholar
- Relevant Coursework:
  - *CS: Discrete Structures I, Computer Science II, Computer Logic & Organization, Object Oriented Programming, Matrix & Linear Algebra*
  - *Bio: Immunology, General Microbiology, Biochemistry I, Molecular Biology II, Quantitative Biological Methods*

Orlando, FL

Spring 2025

### Tallahassee Community College

Associate of Arts (A.A.)

- Phi Theta Kappa Honors Society

Tallahassee, FL

Summer 2022

## RESEARCH EXPERIENCE

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### University of Central Florida

College of Medicine

**Advisor:** Dr. Hung Nguyen

**Research Assistant**

- Independently developed a new immune cell datamining pipeline to analyze PBMC scRNA-Seq datasets to discover correlations between metabolic activity and the grafts vs leukemia (GvL) effect.
- Investigated correlation between the gut microbiome and immune cell behavior using machine learning models to integrate 16S rRNA-Seq & various single-cell assays.

Orlando, FL

December 2023 — Present

### Massachusetts Institute of Technology

Department of Chemical Engineering

**Advisor:** Dr. Brandon DeKosky

**Research Intern** | MIT Summer Research Program

*Sequential Insertion Project (Summer 2023)*

- Contributed to a project improving the efficiency of cell line generation for antibody library display systems by reducing the length of the donor plasmid.
- Successfully performed troubleshooting of restriction cloning experiments to construct donor plasmid for CRISPR/Cas9-mediated genome integration.
- Developed a suite of automated data visualization tools capable of handling high-throughput sequence data.
- Optimized clustering algorithms within bioinformatics pipelines to analyze BCR/TCR immune repertoire data collected through NGS, resulting in a ~70% faster runtime.
- Developed a novel tool using **Hidden Markov Models (HMMs)** and probabilistic approaches to correct errors in next-generation sequencing (NGS) of antibody libraries.

Cambridge, MA

June 2023 — Present

### Bioinformatics & HPC Algorithm Optimization

- Contributed to a project improving the efficiency of cell line generation for antibody library display systems.
- Successfully troubleshooted restriction cloning experiments for construction of donor plasmid for CRISPR/Cas9 system.
- Optimized clustering algorithms within bioinformatics pipelines to analyze BCR/TCR immune repertoire data collected through NGS.
- Developed automated data visualization tools capable of handling high-throughput sequence data.

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**HONORS, AWARDS & SCHOLARSHIPS**


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<b>McNair Summer Research Institute Scholarship</b> <i>Ronald E. McNair Scholars Program</i>	Orlando, FL August 2023
<b>Conference Presentation Travel Award</b> <i>UCF Office of Undergraduate Research</i>	Orlando, FL November 2023
<b>Florida Bright Futures Academic Scholars</b> <i>Florida Department of Education</i>	Tallahassee, FL December 2020
<b>President's List (3 Times)</b>	
<b>Dean's List (2 Times)</b>	

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**PUBLICATIONS**


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**Akinyemi, M., & DeKosky, B. (1979). Probabilistic Approaches to Correct Antibody Library Sequence Errors**  
Econometrica, 47(2), 263-291. (Published)

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**PRESENTATIONS**


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<b>Leadership Alliance National Symposium</b> <u>Oral</u> <i>Probabilistic Approaches to Correct Antibody Library Sequence Errors</i>	Hartford, CT July 2024
<b>UCF Student Scholar Symposium</b> <u>Poster</u> <i>Overcoming Challenges in CRISPR Vector Restriction Cloning to Optimize Generation of Immune Repertoires.</i>	Orlando, FL March 2024
<b>Annual Biomedical Research Conference For Minoritized Scientists (ABRCMS)</b> <u>Poster</u> <i>Generating a Stable Cell Line: Troubleshooting Restriction Cloning of a CRISPR Vector For Sequential Genome Insertion.</i>	Phoenix, AZ November 2023
<b>MIT Summer Research Program Fall Extension Research Showcase</b> <u>Poster</u> <i>Investigating Diversity of V(D)J Gene Recombinants Within Antibody Repertoires</i>	Virtual November 2023
<b>MIT Summer Research Program Showcase</b> <u>Poster</u> <i>Generating a Stable Cell Line: Troubleshooting Restriction Cloning of a CRISPR Vector For Sequential Genome Insertion.</i>	Cambridge, MA July 2023

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**OTHER EXPERIENCES**


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<b>Quantitative Methods Workshop</b> Massachusetts Institute of Technology	Cambridge, MA January 2023
<ul style="list-style-type: none"> <li>• Employed principal component analysis (PCA) and k-means clustering to differentiate cells of a scRNA-Seq PBMC data set.</li> <li>• Used machine learning pipelines (CryoSPARC) to generate 3D protein structures from high-framerate movies of CryoEM data.</li> </ul>	
<b>Scripting Team Leader</b> AvatarMC	Remote May 2017 — June 2022
<ul style="list-style-type: none"> <li>• Developed both command-line &amp; GUI-based automation tools to assist workflows of other teams.</li> <li>• Designed quality assurance testing plans for experimental features.</li> <li>• Assisted in writing &amp; maintaining documentation of in-house scripting language.</li> <li>• Wrote design documents for deliverable requests/feature improvement tasks.</li> <li>• Led projects using professional management software (Phabricator).</li> <li>• Directed team meetings &amp; effectively delegated tasks.</li> </ul>	

KNOWLEDGE & SKILLS

Biological Research

Restriction Cloning, Flow Cytometry, Electroporation, Polymerase Chain Reaction (PCR), Gel Electrophoresis, Agarose Gel Extraction, Immunoprecipitation, Cell Culture (Mammalian & Bacterial), Murine Model Handling, Miniprep/Maxiprep

Programming Languages

Python	Professional.
R	Professional.
C++	Professional.
Bash/Shell	Advanced.
Java	Advanced.
RegEx	Intermediate.

Software

LaTeX	Professional
Git	Professional.
Slurm	Advanced.
SnapGene	Advanced.
Docker	Elementary.
PyMol	Elementary.

Operating Systems

Linux/UNIX	Professional.
HPC-Clusters	Advanced.

Natural Languages

English	C2 level.
Latin	A2 level.
Spanish	A2 level.
Yoruba	A1 level.

COMMUNITY INVOLVEMENT

Society for the Advancement of Chicanos/Hispanics and Native Americans in Science (SACNAS) | *Vice President*

University of Central Florida

Orlando, FL  
April 2024 — Present

- Assisted president in managing club and leading officer meetings.
- Held general body meetings discussing tips/advice to help underrepresented students get started in research and the journey to graduate school.

Active Minds | *Secretary*

University of Central Florida

Orlando, FL  
December 2023 — Present

- Advocated for mental health and the importance of fighting against the negative stigmas associated with them.
- Organized club records, took meeting minutes & performed general administrative duties.