

# Joseph C. Mays

🏠 564 1st Ave, Apt 12C, New York, NY 10016

✉ josephcmays@gmail.com

☎ 973.934.5264

🌐 github.com/joeymays

## EDUCATION

---

**New York University School of Medicine**, New York, NY

August 2018–Present

PhD, Cell Biology

Cumulative GPA: 3.87

**Colgate University**, Hamilton, NY

May 2016

Bachelor of Arts, Molecular Biology (with honors) & Classical Studies

Cumulative GPA: 3.59, Magna Cum Laude

## HONORS, AWARDS, & GRANTS

---

2019	Genome Integrity Training Grant, NYU School of Medicine
2016	NIDCD Director's Award, NIH
2016	Beta Beta Beta Biology Honor Society, Colgate University
2016	Eta Sigma Phi Classics Honor Society, Colgate University
2016	N.L. Andrews Prize for Excellence in the Classics, Colgate University
2013–2016	Dean's Award for Academic Excellence, Colgate University
2016	James M. Maury M.D. Scholarship, Colgate University
2016	Marion A. Cincotta Scholarship, Colgate University
2012–2015	North Eastern Roofing Educational Foundation Scholarship, NERCA
2014	Lila & Curtiss Frank Scholarship, Colgate University

## RESEARCH EXPERIENCE

---

**National Institute on Deafness and other Communication Disorders,**

**National Institutes of Health**, Bethesda, MD

*Post-baccalaureate Fellow, Laboratory of Cochlear Development*

July 2016–August 2018

Mentors: Dr. Michael Kelly, PhD; Dr. Matthew Kelley, PhD

- Evaluated the role of histone acetylation in cell identity maintenance and fate plasticity in the developing mammalian cochlea
- Characterized cellular heterogeneity within the mammalian cochlea using several single-cell RNA sequencing platforms
- Explored transcriptomes of mammalian pineal gland cells and their role in circadian rhythm regulation in collaboration with the Nat'l Institute of Child Health and Development
- Applied computational analyses to evaluate and interpret high-dimensional sequencing data
- Mentored undergraduate students in project management and laboratory techniques
- Collaborated with several NIH labs to share single-cell genomics experience and skills
- Managed a large colony of several transgenic mouse lines for use in experiments
- Presented novel research at national and regional hearing research conferences

**National Institute on Deafness and other Communication Disorders,  
National Institutes of Health, Bethesda, MD**

*Intern, Laboratory of Cochlear Development*

*June 2015–December 2015*

Mentors: Dr. Kathryn Ellis, PhD; Dr. Matthew Kelley, PhD

- Identified lineage relationships of cell types in the mammalian cochlea through clonal analysis
- Analyzed gene expression in the cochlea using immunohistochemistry and confocal microscopy over several developmental time points
- Engaged in biology coursework for the Colgate University-NIH off-campus study program

**Colgate University, Hamilton, NY**

*Research Assistant, Department of Biology*

*January 2014–May 2014*

Mentor: Dr. Douglas Guarneri, PhD

- Assisted Prof. Guarneri in continuing research on stress-induced gene expression in mice
- Validated complex plasmid maps through restriction enzyme digestion
- Implemented software solutions to provide virtual models of plasmids, restriction digests, and site-directed mutagenesis to aid future experimental designs

---

**EMPLOYMENT EXPERIENCE**

**Colgate University, Hamilton, NY**

*Technology Assistant, Classics Department*

*February–May 2016*

- Prepared and maintained presentation equipment for Prof. Ammerman's weekly classes

**MedLabs Diagnostics, Cedar Knolls, NJ**

*Intern, Department of Molecular Diagnostics*

*June–August 2014*

- Performed amplification-based diagnostic assays to test samples for bacteria and parasites
- Participated in diagnostic method comparison studies and method validations
- Compiled and edited standard operating procedures for new diagnostic instruments and methods

**Colgate University, Hamilton, NY**

*Lifeguard, Lineberry Natatorium*

*January 2013–May 2015*

---

**LEADERSHIP EXPERIENCE**

**NYU School of Medicine, New York, NY**

*Secretary, Sackler Institute Student Council*

*Spring 2019–Present*

- Manage and execute ongoing council projects
- Coordinate orientation and recruitment events for new and prospective students
- Organize meetings and resources for council members

**Colgate University, Hamilton, NY**

*President, Colgate Classics Society*

*Spring 2015, Spring 2016*

- Coordinated outreach events with Classics Department faculty to broaden interest in the department and its course offerings
- Communicated and collaborated with society members during bi-monthly meetings
- Managed a semester budget for team-building and outreach events

---

**TEACHING EXPERIENCE**

**National Institute on Deafness and other Communication Disorders,  
National Institutes of Health, Bethesda, MD**

*Instructor, EARssentials Mouse Cochlea Dissection Workshop*

*July 2016, July 2017*

- Co-taught laboratory workshop on neonatal mouse cochlea micro-dissection

## ORAL PRESENTATIONS

---

- *Exploration of a multicellular phenotype in cochlear tissue using single-cell RNA-sequencing methods.*  
NIDCD Trainee Talks, NIH, June 2017.
- *Drop-Seq as a low-cost, high-throughput method for single-cell gene expression profiling of cochlear cells.*  
NIDCD Division of Intramural Research Retreat, NIH, May 2017.
- *Using Seurat and Monocle for analysis of single-cell RNA-sequencing data.*  
NIAMS Bioinformatics Interest Group, NIH, April 2017.
- *Lineage tracing in the developing mammalian cochlea.*  
Colgate University Honors Talks, April 2016.
- *Doing science in college. (Invited Speaker)*  
West Milford High School Science Honor Society, March 2016.

## POSTER PRESENTATIONS

---

- *Single-cell pineal gland neuro-transcriptomic analysis reveals cell type-specific day/night changes.*  
NICHD Division of Intramural Research Retreat, NIH, September 2017.
- *Drop-seq as a low-cost, high-throughput method for single-cell gene expression profiling of cochlear cells.*  
Post-baccalaureate Fellow Poster Day, NIH, May 2017.
- *Demonstration of analysis of high-throughput single cell RNA-Seq data using open-source R packages.*  
Pi Day, NIH, March 2017.
- *Drop-seq as a low-cost, high-throughput method for single-cell gene expression profiling of cochlear cells.*  
Association for Research in Otolaryngology Mid-Winter Meeting, February 2017.
- *Lineage tracing in the developing mammalian cochlea.*  
Summer Student Poster Day, NIH, August 2015.

## OTHER CONFERENCES ATTENDED


---

- *Next-Generation Genomics*  
New York University, New York, NY, August 2019.
- *Eastern Auditory Retreat*  
Georgetown University, Washington, DC, June 2017.
- *Single Cell Analysis Investigators Meeting*  
National Institutes of Health, Bethesda, MD, June 2017.

## PUBLICATIONS

---

**Mays JC**, Kelly MC, Coon SL, Holtzclaw L, Rath MF, Kelley MW, and Klein DC. (2018) Single-cell RNA sequencing of the mammalian pineal gland identifies two pinealocyte subtypes and cell type-specific daily patterns of gene expression. PLOS ONE 13(10): e0205883.

 PubMed 30347410 </> Codebook

Coon SL, Fu C, Hartley S, Holtzclaw L, **Mays JC**, Kelly MC, Kelley MW, Mullikin JC, Rath MF, Savastano LE and Klein DC. (2019) Single Cell Sequencing of the Pineal Gland: The Next Chapter. Front. Endocrinol. 10:590.