ALEXANDER FERRENA

Ph.D. Candidate in Cancer Bioinformatics at Albert Einstein College of Medicine

alexander.ferrena@gmail.com

github.com/FerrenaAlexander | linkedin.com/in/alexander-ferrena | ferrenaalexander.github.io Seeking bioinformatics research opportunities in pharmaceuticals and biotech, starting June 2024.

EDUCATION

August 2019 –

Expected Graduation,
May 2024

Ph.D. Candidate, Albert Einstein College of Medicine (AECOM)

Ph.D. in Clinical Investigation, Institute for Clinical and Translational Research
Thesis: Molecular and Clinical Implications of Skp2 inhibition in Osteosarcoma

M.A. Biotechnology, Columbia University
Department of Biological Sciences, Graduate School of Arts and Sciences
Thesis: The Impact of Aging on Lung Adenocarcinoma.

September 2012 –
May 2016

B.A. Biology, New York University
College of Arts and Sciences

EXPERIENCE

June 2020 – Present	Ph.D. Candidate, Albert Einstein College of Medicine Lab of Dr. Deyou Zheng (Genetics Dept.) and Dr. Bang Hoang, Orthopedics Designed and applied analysis workflows for single-cell transcriptomics data from cancer and embryonic development. Developed a pipeline package for scRNA-seq integration and replicate-aware cross-condition comparison.
August 2022 – March 2023	Bioinformatics Consultant, Singulomics Corporation Developed an efficient, modular analysis pipeline for spatial transcriptomics data.
May 2018 – June 2020	Computational Biology Research Assistant, Memorial Sloan Kettering Lab of Dr. Tuomas Tammela, Memorial Sloan Kettering Cancer Center Analyzed high-throughput sequencing data such as scRNA-seq, CITE-seq, and others as a bioinformatician in a dynamic cancer genetics lab.
June 2016 – September 2017	Bioinformatics Internship, New York University (NYU) Lab of Dr. David Gresham, NYU Center for Genomics and Systems Biology

Bulk RNAseq data analysis of differential conditions of yeast culture conditions.

TECHNICAL SKILLS

<u>Programming</u>: Advanced, daily use of **R** for genomic data analysis and scalable package development, including libraries like Seurat, EdgeR, and InferCNV. Daily use of **Unix shell (bash)**. Experience with **Python** for genomics applications including use of Scanpy, scVelo and Cellrank.

<u>**Data analysis**</u>: Broad experience in multivariate statistics and machine learning for big data.

<u>Cloud Computing</u>: Extensive use of High-Performance Clusters via the SLURM and Sun Grid schedulers, and experience with AWS.

PUBLICATIONS

March 2024 scDAPP: a comprehensive single-cell transcriptomics analysis pipeline optimized for cross-group comparisons. In preparation. (In preparation) Ferrena A, Zheng XY, Jackson K, Hoang B, Zheng D. February 2024 Comprehensive single cell transcriptomics analysis of murine osteosarcoma uncovers (In preparation) Skp2 function in metastasis, genomic instability and immune activation and reveals additional target pathways. In preparation. Ferrena A, Zhang R, Wang J, Zheng XY, Zhao H, Schwartz E, Loeb D, Goker B, Borjihan H, Chae S, Yang R, Geller D, Zheng D, Hoang B. February 2024 Facilitate integrated analysis of single cell multiomic data by binarizing gene expression values. In review, Nature Communications. Posted to bioRxiv Feb 2024. (Preprint) Misra R, Ferrena A, Zheng D. Targeted inhibition of SCFSKP2 confers anti-tumor activities resulting in a survival February 2024 benefit in osteosarcoma. Oncogene, February 2024. Wang J*, Ferrena A* (co-first author), Singh S, Zhang R, Viscarret V, Al-Harden W, Aldahamsheh O, Borjihan H, Singla A, Yaguare S, Tingling J, Zi X, Yungtai L, Gorlick R, Schwartz L, Zhao H, Yang R, Geller D, Zheng D, Hoang B. October 2023 SKP2 knockout in Rb1/p53 deficient mouse models of osteosarcoma induces immune infiltration and drives a transcriptional program with a favorable prognosis. Molecular Cancer Therapeutics, October 2023. Ferrena A, Wang J, Zhang R, Karadal-Ferrena B, Al-Hardan W, Singh W, Borjihan H, Schwartz E, Zhao H, Oktay M, Yang R, Geller D, Hoang B, and Zheng D. May 2023 Six3 and Six6 jointly regulate the identities and developmental trajectories of multipotent retinal progenitor cells in the mouse retina. In review, *PLOS ONE*. Preprint posted to *bioRxiv* May 2023. Ferrena A, Zhang X, Shrestha R, Zheng D, Liu W. March 2023 Single cell transcriptomics uncovers a non-autonomous Tbx1-dependent genetic program controlling cardiac neural crest cell deployment and progression. Nature Communications, 2023 March. De Bono C, Liu Y, Ferrena A, Valentine A, Zheng D, Morrow B. October 2022 Systematic comparison of pancreatic ductal adenocarcinoma models identifies a conserved highly plastic basal cell state. Cancer Research, 2022 October. Pitter K, Grbovic-Huezo O, Joost S, Singhal A, Blum M, Wu K, Holm M, Ferrena A, Bhutkar A, Hudson A, Lecomte N, de Stanchina E, Chaligne R, Iacobuzio-Donahue C, Pe'er D, and Tammela T.

April 2021

Geller DS, Hoang B.

The interaction of SKP2 with p27 enhances the progression and stemness of osteosarcoma. Annals of the New York Academy of Sciences, 2021 April.

Wang J, Aldahamsheh O, Ferrena A, Borjihan H, Singla A, Yaguare S, Singh S, Viscarret V, Tingling J, Zi X, Lo Y, Gorlick R, Zheng D, Schwartz EL, Zhao H, Yang R,

FELLOWSHIPS, HONORS, AND AWARDS

May 2020 <u>Fellowship Award</u>, PhD program in Clinical Investigation (PCI)

Applied for and was accepted to this highly competitive fellowship and training program offered to PhD students at Einstein with research proposals focused on biostatistics and clinical research. This program offers advanced, formal

biostatistics training with a highly clinical and translational focus.

September Student Spotlight, Columbia Graduate School of Arts and Sciences

2019 Received distinction for academic and research performance

Poster Presentation distinction, NYU Undergraduate Research Conference

May 2015 Awarded best in group for poster presentation titled *Identification and*

Characterization of Orthodenticle-Dependent Enhancers.

LEADERSHIP EXPERIENCE

January 2021 – Founder and President, Einstein Montefiore Omics Club

Present Founded this student organization to facilitate networking, collaboration, and

education around -omics technologies. Organized the *Intro to Omics Workshop* series attended by over 150 people to teach bioinformatics data analysis. Led a

session on single cell transcriptomics at this workshop.

September 2019 – <u>Vice-President</u>, Einstein Biotech Club

Present Served on the executive board of this club promoting student engagement and

education with biotechnology. Led and organized many networking events.

November 2019 – <u>Elected Councilor</u>, Einstein Diversity and Inclusion (D&I) Council

May 2023 Elected to serve on this committee for diversity enhancement by the student body of

at Albert Einstein College of Medicine.

September 2017 – President, Columbia Biotechnology Association (GRO-Biotech CU)

May 2019 Served as President of this industry networking group. Helped organize the 2019

Graduate Research Organizations Conference at Columbia with >300 attendees.

TEACHING EXPERIENCE

February 2024 Intro to Omics Workshop, Einstein Montefiore Omics Club

Led a workshop session on scRNA-seq analysis and cell annotation.

September 2017 – <u>Teaching Assistant</u>, Intro to Genomic Information and Technology

December 2017 Worked as a teaching assistant for this genomic data science course taught by

Dr. Dimitris Anastassiou at Columbia.

INTERESTS AND HOBBIES

August 2018 – My Aquarium, Founder and Chief Executive Fish-Fanatic

Present A fun and relaxing pastime is to watch the fish and plants in my tropical aquarium.

CONFERENCES

October 2023 <u>Musculoskeletal Tumor Society (MSTS)</u>, Podium presentation

Gave a talk at this orthopedic oncology-focused meeting titled "Single-cell transcriptomics of osteosarcoma murine models with SKP2 modulation predicts

potent synergistic therapies".

April 2023 <u>American Association for Cancer Research (AACR)</u>, Poster presentation

Presented a poster at this cancer-focused conference titled "SKP2 knockout induces macrophage infiltration in p53/Rb1 null transgenic mouse models of osteosarcoma and drives cone expression correlated with improved survival in national."

and drives gene expression correlated with improved survival in patients".

Musculoskeletal Tumor Society (MSTS), Podium presentation

December 2022 Gave a talk at this orthopedic oncology-focused meeting titled "SKP2 knockout in

Rb1/p53 deficient transgenic mouse models of osteosarcoma improves survival via

induction of macrophage infiltration".

International Society for Limb Salvage (ISOLS), Podium presentation

September 2022 Gave a talk at this orthopedic oncology-focused meeting titled "Knockout of SKP2"

in pre-clinical transgenic mouse models of OS induces macrophage infiltration and

improves survival".

NYU Undergraduate Research Conference, Poster presentation

May 2015 Designed and presented a poster titled "Identification and Characterization of

Orthodenticle-Dependent Enhancers." Won an award for best poster in the session.

PROFESSIONAL REFERENCES

Dr. Deyou Zheng (Ph.D.), Primary PhD Thesis Advisor

- **Position**: Professor, Departments of Genetics, Neuroscience, and Neurology
- Location: Albert Einstein College of Medicine
- Contact: https://www.einsteinmed.edu/faculty/10976/deyou-zheng/
- **Relationship**: Deyou is the primary advisor of my Ph.D. He is an expert in genetics, genomics, and bioinformatics. Deyou has a long track record of leading genomics and transcriptomics studies in a variety of contexts including cancer and embryonic development and has mentored many students through the PhD program at Einstein.

Dr. Bang Hoang (M.D.), Secondary PhD Thesis Advisor

- **Position**: Professor, Department of Orthopedic Surgery
- Location: Albert Einstein College of Medicine, Montefiore Medical Center
- **Contact:** https://www.einsteinmed.edu/faculty/14186/bang-hoang/
- **Relationship**: Bang is my Ph.D. co-mentor. He is an orthopedic surgeon and expert in patient care for osteosarcoma and other sarcomas along with directing a research lab. He serves as the Chair of the Research Committee for the Musculoskeletal Tumor Society.