TOMAS BENCOMO

tjbencomo@gmail.com tjbe		ncomo.github.io
EDUCATION		
MD/PhD	University of Washington Medical Scientist Training Program Mentors: Paul Nghiem and Manu Setty	2022 - present
BS	Stanford University, Computer Science Concentration in Biocomputation GPA: 3.99	2017 - 2021
HONORS AND A	AWARDS	
	andation Award nt Reward For College Scientist scholarship to help fund PhD training	2024
	Bio-X Summer Research Grant r undergraduate research in the Dermatology department	2020
Tau Beta I Engineerin	Pi Member g honors society at Stanford	2020
	PUE Conference Grant research at Society For Investigative Dermatology Annual Meeting	2020
	ic Summer Undergraduate Research Fellowship lying glioblastoma in the Precision NeuroTheraputics Program	2018 & 2019
TGen Heli High schoo	os Scholar Il intern in the Helios Program at TGen	2016
•	proposium Poster Award Winner or a top 10 poster at the intern symposium	2016
RESEARCH EX	PERIENCE	
Advisors: I Stud	Student, University of Washington & Fred Hutch Cancer Center Paul Nghiem MD, PhD & Manu Setty PhD died mechanisms of immunotherapy resistance using spatial multi-omics technologies reloped novel computational approaches for spatial transcriptomics data analysis	2023 – present
Advisor: C Invo Per	Associate, Lee Lab, Stanford Medicine arolyn Lee MD, PhD estigated mechanisms of perineural invasion using single cell sequencing formed a meta-analysis of publicly available cSCC RNA-Seq datasets died metabolic reprogramming caused by non-coding mutations in OXA1L	2021 – 2022

Undergraduate Researcher, Lee Lab, Stanford Medicine

Advisor: Carolyn Lee MD, PhD

- Characterized novel genes implicated in skin cancers using multi-omics data
- Examined survival data to find biomarkers for squamous cell carcinoma (SCC)
- Analyzed proteomics data to identify novel interactors for melanoma oncogene RAC1

Summer Undergraduate Research Fellow, Swanson Lab, Mayo Clinic

2018-2019

2017 - 2021

Advisor: Kristin Swanson PhD

- Analyzed MRI data to study blood brain barrier disruption in glioblastoma (GBM)
- Trained machine learning models to predict drug distribution in the brain from MRI
- Built prognostic models using radiomics data to predict patient survival

Helios Scholar, Translational Genomics Research Institute

2016

Advisor: Seungchan Kim PhD and Gil Speyer PhD

- Redesigned the lab's EDDY algorithm to use NVIDIA's CUDA API
- Helped secure \$200,000 Compute 4 the Cure grant from NVIDIA
- Achieved 500x speedup; the algorithm can now analyze previously impossible datasets

PUBLICATIONS (GOOGLE SCHOLAR)

Articles

Gene expression landscape of cutaneous squamous cell carcinoma progression

2024

Tomas Bencomo, Carolyn S Lee

British Journal of Dermatology (accepted)

Unravelling the landscape of skin cancer through single-cell transcriptomics

2023

Ankit Srivastava, **Tomas Bencomo**, Ishani Das, Carolyn S. Lee <u>Translational Oncology</u>

MAB21L4 deficiency drives squamous cell carcinoma via activation of RET

2022

Ankit Srivastava, Cristina Tommasi, Dane Sessions, Angela Mah, **Tomas Bencomo**, Jasmine M Garcia, Tiffany Jiang, Michael Lee, Joseph Y Shen, Lek Wei Seow, Audrey Nguyen, Kimal Rajapakshe, Cristian Coarfa, Kenneth Y Tsai, Vanessa Lopez-Pajares, Carolyn S Lee Cancer Research

Mutant collagen COL11A1 enhances cancerous invasion

2021

Carolyn Lee, Zurab Siprashvili, Angela Mah, **Tomas Bencomo**, Lara Elcavage, Yonglu Che, Rajani Shenoy, Sumaira Aasi, Paul Khavari Oncogene

Preprints

ecDNA in cutaneous squamous cell carcinoma is associated with increased nodal disease

Tomas Bencomo, Amarinder Thind, Bruce Ashford, Marie Ranson, Carolyn S. Lee Biorxiv

2023

2024

Gene expression landscape of cutaneous squamous cell carcinoma progression Tomas Bencomo, Carolyn S Lee

Tomas Dencomo, Caroryn S Lec

Biorxiv

Conference Abstracts

Gene expression landscape of cutaneous squamous cell carcinoma progression

T. Bencomo, C. S. Lee

Society For Investigative Dermatology 2024 Annual Meeting

ecDNA in cutaneous squamous cell carcinoma is associated with increased nodal disease

T. Bencomo, A. Thind, I. Donohue, B. Ashford, M. Ranson, C. S. Lee

Society For Investigative Dermatology 2024 Annual Meeting

Mast cells orchestrate perineural invasion in aggressive epithelial cancers

A. Srivastava, T. Bencomo, A. Mah, J. Garcia, I. Donohue, A. Tan, L. Seow, A. Nguyen, T. Jiang,

P. Dwivedi, C. M. Rose, R. Brown, S. Aasi, C. S. Lee

Society For Investigative Dermatology 2024 Annual Meeting

Metabolic reprogramming by cancer-associated mutant OXA1L promotes tumor progression and dedifferentiation

A. Srivastava, I. Donohue, A. Tan, A. Nguyen, T. Bencomo, J. Ye, B. Ashford, M. Ranson, D.

Kashatus, C. S. Lee

Society For Investigative Dermatology 2024 Annual Meeting

Chromosomal structural variants but not extrachromosomal DNAs are common in Merkel cell carcinoma

J. D. Bloomstein , J. Luebeck , J. Rose , **T. Bencomo**, D. S. Hippe , S. Jani, V. Bafna , H. Chang , P. Nghiem

Society For Investigative Dermatology 2024 Annual Meeting

Somatic Mutation of the OXA1L 5'UTR enables Cutaneous Squamous Cell Carcinoma

Angela Mah, Jasmine Garcia, Dane Sessions, **Tomas Bencomo**, Ashley Amado, Ankit Srivastava, Carolyn Lee

Plenary Session - Society For Investigative Dermatology 2021 Annual Meeting

Molecular Profiling of Cutaneous C-Group Non-Langerhans Cell Histiocytoses

Rebekah Wieland, Tomas Bencomo, Carolyn Lee, Ryanne Brown

United States and Canadian Academy of Pathology 2021 Annual Meeting

Sex Differences In GBM Patient Survival As A Function of Extent Of Surgical Resection and Cycles of Adjuvant Temozolomide During Standard Of Care Regimens

Julia Lorence, **Tomas Bencomo**, Haylye White, Cassandra Rickertsen, Susan Massey, Kyle Singleton, Andrea Hawkins-Daarud, Sandra Johnston, Alyx Porter, Maciej Mrugala, Bernard Bendok, Leland Hu, Joshua Rubin, Kristin Swanson

Society for Neuro-Oncology 2020 Annual Meeting

Rac1-interacting proteins are prognostic factors for melanoma survival

Marten CG Winge, **Tomas Bencomo**, M. Peter Marinkovich, Carolyn S. Lee Society For Investigative Dermatology 2020 Annual Meeting

Sex differences in GBM treatment: an observational study

Lorence J., **Bencomo T.**, White H., Rickertsen C.R., De Leon, G. Singleton K. W., Daruud-Hawkins A., Bendok B. R., Porter A. B., Mrugala M. M., Rubin J.B., and Swanson K. R.

3rd Annual Celebration of Women's Health Research – Sexx as a Biological Variable

Sex Differences in Tumor Location Incidence in Newly Diagnosed Glioblastoma Patients

Lorence J., **Bencomo T.**, White H., Rickertsen C.R., De Leon G., Singleton K., Bendok B., Porter A. B., Rubin J. B., Swanson K. R.

Society for Neuro-Oncology 2019 Annual Meeting

Modeling the interface between non-invasive imaging and drug distribution

Ranjbar, Sara; **Bencomo, Tomas**; Jackson, Pamela R; Randall, Elizabeth; Regan, Micheal; Abdelmoula, Walid M; Lopez, Begona GC; Massey, Susan Christine; He, Lihong; Macura, Slobodan; Hu, Leland; Agar, Jefferey N; Sarkaria, Jann; Agar, Nathalie; Swanson, Kristin R. CSBC/PS-ON 2019 Annual Investigators Meeting

The Same But Different: Identifying Distinct Imaging Ecologies in Male and Female Glioblastomas

Tomas Bencomo, Andrea Hawkins-Daarud, Kyle Singleton and Kristin R. Swanson. <u>CSBC/PS-ON 2018 Annual Investigators Meeting</u>

GPU-accelerated differential dependency network analysis

Gil Speyer, Juan Rodriguez, **Tomas Bencomo** and Seungchan Kim Euromicro International Conference on Parallel, Distributed and Network-Based Processing 2018

LANGUAGES

English: Native Language

Spanish: Intermediate Proficiency

TECHNICAL SKILLS

Programming: R, Python, Java, C, C++

Biostatistics: Regression modeling, machine learning, survival analysis, experimental design

Bioinformatics: Mutation profiling, bulk and single cell RNA-sequencing, spatial analysis, data visualization

Computing: Snakemake, Docker, Unix, Bash, HPC, scikit-learn, pandas, tidyverse

EXTRACURRICULARS

Tau Beta Pi Honors Society Board – Academic Co-Chair Organize academic development events and lead the Peer Mentorship Program Computer Science Peer Mentor Advise and mentor underclassmen considering a major in Computer Science