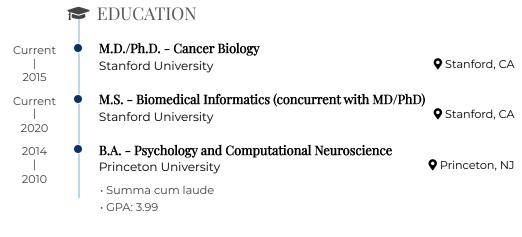
TIMOTHY KEYES

I am a data scientist, bioinformatician, and cancer biologist. In my work, I develop statistical and machine learning algorithms for analyzing high-dimensional single-cell data and predicting clinical outcomes in cancer patients.

I am currently searching for a position at the intersection of biomedical data science, machine learning, and medicine where I can use data to solve problems relevant to human health.



EMPLOYMENT

Current | 2022 Data Science Mentor - Posit Academy Posit, PBC (formerly RStudio, PBC)

Stanford, CA

2022

Graduate Intern - Oncology Bioinformatics, gRED

Genentech. Inc

South San Francisco, CA

■ SELECT PUBLICATIONS

2022

{tidytof}: A user-friendly framework for scalable and reproducible high-dimensional cytometry data analysis.

Under review (copy available upon request)

- · Keyes TJ, Koladiya A, Lo YC, Nolan GP, Davis KL.
- · Project website: https://keyes_timothy.github.io/tidytof/

2022

CytofIn enables Integrated Analysis of Public Mass Cytometry Datasets using Generalized Anchors

Nature Communications

· Lo YC, **Keyes TJ**, Jager A, Sarno J, Domizi P, Majeti R, Sakamoto KM, Lacayo N, Mulligan CG, Waters J, Sahaf B, Bendall SC, Davis KL

2020

A cancer biologist's primer on machine learning applications in highdimensional cytometry

Cytometry

· Keyes TJ, Domizi P, Lo YC, Nolan GP, and Davis KL

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PROGRAMMING

R / tidyverse / tidymodels
Python
git / GitHub
C++
Bash
HTML
CSS

Q DATA ANALYSIS

Exploratory data analysis
Data visualization (e.g. ggplot2)
Data cleaning (e.g. dplyr, pandas)
Deep Learning (Keras, TF)
Machine learning (e.g. Factor
Analysis, GLMs, SVMs, Tree-based
models)

LITERATE CODING

R Markdown
Quarto
Jupyter Notebooks
LaTex
flexdashboards
blogdown / bookdown

Last updated on 2022-12-04.