ALEX RAJEWSKI

Experienced bioinformatics data scientist with experience in oncology, drug design, single-cell sequencing, spatial transcriptomics, epigenetics, genome assembly, and phylogenetics. Also skilled in pipeline development, including Nextflow, Docker, Singularity, and Git. I am an excellent science communicator to both public and professional audiences.



☐ WORK EXPERIENCE

current | 2023

Senior Computational Biologist

Tempus Al

2023 | 2022

Bioinformatics Scientist

Tempus AI (fmr. SEngine Precision Medicine)

- · Development of high-throughput drug screening analysis pipelines in R
- · Consultation on experimental design and statistical analysis

2022 | 2021

Research Bioinformatician II

Cedars-Sinai Genomics Core

- Advanced analysis of single-cell RNA-seq, spatial transcriptomics, WES/WGS, CUT&Taq, ATAC-seq
- · Pipeline development with Nextflow, Docker, and Git

2012

Congress-Bundestag Fellow & Research Intern

Universität Rostock

- · Vaccine production in transgenic plants
- $\boldsymbol{\cdot}$ Developed plant transformation methods in lupine, pea, to bacco, and potato

2011 | 2010

Research Assistant

Pioneer Hi-Bred (Corteva)

 Developed SQL database to integrate phenotypic and genotypic measurements

Aug 2009 | May

2009

NSF Research Intern

North Carolina State University

· Validated viral protein production in novel hosts

CONTACT

✓ AlexCRajewski@gmail.com

github.com/rajewski

rajewski.github.io

in in/alexrajewski

LANGUAGE SKILLS

R
German
Docker/Singularity
Shell
Nextflow
Python
SQL
汉语

EDUCATION

2020 • PhD Bioinformatics & Plant Biology

University of California, Riverside

- · Genome assembly/annotation, RNA-seq, phylogenetics, high-throughput CRISPR screening
- · NGS library prep, PCR, tissue culture, in situ hybridization, microscopy

2015 • MS Horticulture

University of Georgia

· Population genetics, Microsatellites/SSR, phylogenetics

2010 • BS Biochemistry, Cell, and Molecular Biology

Drake University

■ SELECTED PUBLICATIONS

• SOX9 switch links regeneration to fibrosis at the single-cell level in mammalian kidneys

Science

2024

S. Aggarwal; Z. Wang; D Rincon Fernandez Pacheco; A Rinaldi; **A. Rajewski**; J. Callemeyn; E. Van Loon; B. Lamarthée; A. Ester Covarrubias; Jean Hou et al.

2023 • Stacking the odds: Multiple sites for HSV-1 latency

Science Advances.

Shaohui Wang, Xueying Song, Alex Rajewski, Chintda Santiskulvong, and Homayon Ghiasi

2022 • Multispecies Transcriptomes Reveal Core Fruit Development Genes

Frontiers in Plant Science

Alex Rajewski, Dinusha Maheepala, Jessica Le, Amy Litt

Cell therapy attenuates endothelial dysfunction in hypertensive rats with heart failure and preserved ejection fraction

Heart and Circulatory Physiology

G. de Couto, T. Mesquita, X. Wu, **A. Rajewski,** F. Huang, A. Akhmerov, N. Na, D. Wu, Y. Wang, L. Li, M. Tran, P. Kilfoil, E. Cingolani, E. Marbán

Datura Genome Reveals Duplications of Psychoactive Alkaloid Biosynthetic Genes and High Mutation Rate Following Tissue Culture

BMC Genomics

Alex Rajewski, Derreck Carter-House, Jason Stajich, and Amy Litt

in Vitro Plant Regeneration and Agrobacterium tumefaciens-mediated Transformation of Datura stramonium

Applications in Plant Science

Alex Rajewski, Kevan Elkins, Ashley Henry, Joyce Van Eck, and Amy Litt

2018

Classification and phylogenetic analyses of the Arabidopsis and tomato G-type lectin receptor kinases

BMC Genomics

Marcella A. Teixeira, **Alex Rajewski**, Jiangman He, Olenka G. Castaneda, Amy Litt, and Isgouhi Kaloshian