PATRICK CHERRY

💌 pcherry [at] pm dot me | 📞 upon request | 🛍 Senior Scientist | Genomics | 🖫 Twist Bioscience | 🛍 San Francisco, California | 😚 pdcherry.github.io | 😱 github.com/pdcherry | in linkedin.com/in/p-cherry | 😇 United States Citizen | Updated on 2024-07-30

PhD scientist and professional critical thinker, problem-solver, and science communicator. At the interface of data science, genomics, and statistics; skilled at data visualization, bioinformatics, next-generation seguencing (NGS), and tool-building. I've coded reproducible pipelines for high-throughput experimental designs and genomic analyses. I've launched best-in-class oncology reference standards and analyzed NGS data from Illumina, MGI, and Element. Originally trained in Molecular Biology, I am passionate about advancing data science and bioinformatics to improve human health and to understand biology.



May 2019 Aug 2013

PhD

University of Colorado School of Medicine

Aurora/Denver. Colorado

- · Ph.D. in Molecular Biology. Advisor: Jay Hesselberth, PhD
- Thesis: RNA terminus chemistry impact decay events that target HAC1 mRNA during the Unfolded Protein Response

May 2013 Aug 2009

BA

Hendrix College

Conway, Arkansas

- Biochemistry and Molecular Biology, with Distinction. Advisor: Andres Caro, PhD
- · Senior Capstone Project showing key stress response gene expression changes to oxidative stress in liver cells
- · Minor in Mathematics; PI: Lars Seme; Project: Newton's method as a fractal chaotic dynamical system



INDUSTRY EXPERIENCE

Apr 2024 Nov 2021

Senior Scientist, Genomics

Twist Bioscience

South San Francisco, California

- Produced \$1M+ in new revenue as Tech Lead on invention and launch of: Pan-cancer cfDNA (v1 & v2), CNV Controls, RNA Fusion Controls, Fragmentome Controls, and RNA-seq
- Answered biological and business questions with DoE and data analysis using R and the libraries: dplyr, dbplyr, DBI, purrr, ggplot2, Bioconductor, DEseq2, seurat, tidymodels, glm, nls, lme4, AlgDesign, and more
- · Edited and maintained production QC pipelines with automated reporting using Python and the libraries: NumPy, pandas, polars, seaborn, statsmodels, biopython, pybedtools, pysam, vcfpy, scanpy, and others
- · Maintained production code (qit version control with code review) with CD (github actions); operated in highperformance compute environment aws s3 and Databricks using Unix shell / bash tools, like STAR-fusion
- · Authored internal R package "twistcolorpal" on Github with help files to help scientists style plots to Twist branding

Apr 2024 Jan 2021

Scientist, Genomics

Twist Bioscience

South San Francisco, California

- Earned +\$1M in revenue in year 1 as Tech Lead of Twist Pan-Cancer Reference Standard, an ISO-13485 synthetic control with 458 variants among 84 cancer-associated genes at six QC'd VAFs, plus a WT control, launched Nov. 2021
- · Identified and optimized compatible ddPCR assays using web scraping in R with rvest for cfDNA quality control
- Automated data / QC reports with Python packages Jinja2 (with CSS) and WeasyPrint, documented on Jira
- · Managed data and ran bioinformatic analyses on Illumina RNA & DNA seq using bash tools like: awk, grep, wget, ssh, GATK, BaseSpace CLI, bwa, bedtools, samtools, vcftools, UMI-tools/fgbio; Confluence documentation
- · Answered production and business questions with original SQL queries for Snowflake-based database

Jan 2021 Apr 2019

Scientist I, NGS Core & Strain-Build Process Development

Zymergen, Inc.

Emeryville, California

- Boosted NGS Core genotyping success by 45% using DoE experimentation, data-driven decision-making, and teaching; Guided demanding and diverse internal customers on complex NGS experiments
- · Built and disseminated Rmarkdown and JMP statistical templates for autonomous NGS data exploration
- Coded plasmid and strain build and QC experiments using Zymergen's alembic Python API to LIMS database
- Rendered strain build and plasmid build reports from LIMS SQL database in MySQL Workbench

SELECTED INTELLECTUAL PROPERTY Libraries for mutational analysis Apr 2021 South San Francisco, California Twist Bioscience US Prov. Pat. Ref No 823.101 · Configuration and fabrication of synthetic DNA & RNA reference standards and synthetic variant sequences Method for counterselection in microorganisms Mar 2021 Emeryville, California Zymergen, Inc. · US 2021 0087586 A1 • Demonstration of novel counterselection mechanism in non-model Bacillus microbe for genetic modification SELECTED PUBLICATIONS Twist cfDNA Pan-Cancer Reference Standard v2 Technical Guidance Jan 2024 South San Francisco, California Twist Bioscience Product Sheet & FAQ · Patrick Cherry, Lydia Bonar, & Mike Bocek Characteristics and specificity of the wild-type / 0% VAF reference material Apr 2022 South San Francisco, California Twist Bioscience · Patrick Cherry & Mike Bocek Multiple decay events target HAC1 mRNA during splicing to regulate the unfolded protein response Mar 2019 Aurora/Denver, Colorado eLife · Cherry, P., Peach, S., & Hesselberth, J. Genetic bypass of essential RNA repair enzymes in budding yeast Dec 2017 Aurora/Denver, Colorado **RNA** · Cherry, P., White, L., York, K., & Hesselberth, J. ♣ SELECTED PRESENTATIONS & TALKS Twist pan-cancer reference standard V2: Enhanced precision and reduced errors in ctDNA analysis Feb 2024 Orlando, Florida Advances in Genome Biology and Technology, Twist Bioscience Twist pan-cancer synthetic RNA fusion control for assay development Apr 2023 Orlando, Florida American Association for Cancer Researchers, Twist Bioscience Use of synthetic CNV fragments to mimic copy number alterations for ctDNA reference standards Feb 2023 Hollywood, Florida Advances in Genome Biology and Technology, Twist Bioscience Twist pan-cancer synthetic reference materials for cell-free DNA (cfDNA) assay development Apr 2022 New Orleans, Louisiana American Association for Cancer Researchers, Twist Bioscience R use at Zymergen Jun 2020 Emeryville, CA Z-Tech Talk Data-driven troubleshooting of NGS experiments Apr 2020 Emeryville, CA Data Science Talk RNA processing regulates the unfolded protein response Aug 2017 Ocld Spring Harbor, New York CSHL: mRNA Processing Meeting Talk