

# PATRICK CHERRY

✉ pcherry [at] pm dot me | 📞 upon request | 🏢 Senior Scientist | Genomics | 🏠 Twist Bioscience | 🏡 San Francisco, California |  
🌐 [pdcherry.github.io](https://pdcherry.github.io) | 🐙 [github.com/pdcherry](https://github.com/pdcherry) | in [linkedin.com/in/p-cherry](https://www.linkedin.com/in/p-cherry) | 🇺🇸 United States Citizen | Updated on 2024-12-10

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 sequencing (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.

## 🎓 EDUCATION



- 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

- Current  
|  
Sep 2024
- **Senior Scientist, Oncology**  
BillionToOne 📍 Menlo Park, CA
    - Generated 500+ automated ddPCR primer / probe designs with pipeline in snakemake using s3 database of variant data and primer3 primer selection
    - Informed key product decisions with historical analyses using PostgreSQL / ssh and s3 data sources and linear mixed-effects modeling
- Apr 2024  
|  
Jan 2021
- **Senior Scientist, Genomics**  
Twist Bioscience 📍 South San Francisco, California
    - Invented and introduced multiple new products to market, yielding \$1M+ in new revenue as *Tech Lead*, including: *Pan-cancer cfDNA* (v1 & v2), *CNV Controls*, *RNA Fusion Controls*, *Fragmentome Controls*, RNA-seq, and primer / adapter removal methods for highly-multiplexed dsDNA pools
    - 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, Jinja2 (with CSS), WeasyPrint, and others
    - Maintained production code (git version control with code review) with CD (github actions); operated in high-performance compute environment aws s3 and Databricks using Unix shell / bash tools, like ssh, STAR-fusion, GATK, BaseSpace CLI, bwa, bedtools, samtools, vcftools, UMI-tools/fgbio; Confluence documentation
    - Authored internal R package "[twistcolorpal](#)" on Github with help files to help scientists style plots to Twist branding
    - 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 R markdown 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

- Apr 2021 • **Libraries for mutational analysis**  
Twist Bioscience  South San Francisco, California
- US Prov. Pat. Ref No 823.101
  - Configuration and fabrication of synthetic DNA & RNA reference standards and synthetic variant sequences
- Mar 2021 • **Method for counterselection in microorganisms**  
Zymergen, Inc.  Emeryville, California
- US 2021\_0087586 A1
  - Demonstration of novel counterselection mechanism in non-model *Bacillus* microbe for genetic modification










## SELECTED PUBLICATIONS

- Jan 2024 • **Twist cfDNA Pan-Cancer Reference Standard v2 Technical Guidance**  
Twist Bioscience [Product Sheet](#) & [FAQ](#)  South San Francisco, California
- Patrick Cherry, Lydia Bonar, & Mike Bocek
- Apr 2022 • **Characteristics and specificity of the wild-type / 0% VAF reference material**  
Twist Bioscience  South San Francisco, California
- Patrick Cherry & Mike Bocek
- Mar 2019 • **Multiple decay events target HAC1 mRNA during splicing to regulate the unfolded protein response**  
[eLife](#)  Aurora/Denver, Colorado
- Cherry, P., Peach, S., & Hesselberth, J.
- Dec 2017 • **Genetic bypass of essential RNA repair enzymes in budding yeast**  
[RNA](#)  Aurora/Denver, Colorado
- Cherry, P., White, L., York, K., & Hesselberth, J.



## SELECTED PRESENTATIONS & TALKS

- Feb 2024 • **Twist pan-cancer reference standard V2: Enhanced precision and reduced errors in ctDNA analysis**  
[Advances in Genome Biology and Technology](#), Twist Bioscience  Orlando, Florida
- Apr 2023 • **Twist pan-cancer synthetic RNA fusion control for assay development**  
[American Association for Cancer Researchers](#), Twist Bioscience  Orlando, Florida
- Feb 2023 • **Use of synthetic CNV fragments to mimic copy number alterations for ctDNA reference standards**  
[Advances in Genome Biology and Technology](#), Twist Bioscience  Hollywood, Florida
- Apr 2022 • **Twist pan-cancer synthetic reference materials for cell-free DNA (cfDNA) assay development**  
[American Association for Cancer Researchers](#), Twist Bioscience  New Orleans, Louisiana
- Jun 2020 • **R use at Zymergen**  
Z-Tech Talk  Emeryville, CA
- Apr 2020 • **Data-driven troubleshooting of NGS experiments**  
Data Science Talk  Emeryville, CA
- Aug 2017 • **RNA processing regulates the unfolded protein response**  
CSHL: mRNA Processing Meeting Talk  Cold Spring Harbor, New York