

# 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 2024-05-27

PhD scientist skilled in data visualization, statistical modeling, bioinformatics, next-generation sequencing (NGS), and tool-building. I've coded reproducible and rigorous pipelines for high-throughput experimental designs and multi-omic analyses for communication to technical and non-technical audiences. 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 understand biology.

## 🎓 EDUCATION

- 2019  
|  
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
- 2013  
|  
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

- 2024  
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2021  
**Senior Scientist, Genomics**  
Twist Bioscience 📍 South San Francisco, California
  - Invented and introduced multiple new products to market yielding millions of dollars in new revenue as *Tech Lead*, including: *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, vcfp, scanpy, 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 STAR-fusion
  - Authored internal R package "twistcolorpal" on Github with help files to help scientists style plots to Twist branding
- 2024  
|  
2021  
**Scientist, Genomics**  
Twist Bioscience 📍 South San Francisco, California
  - Tech Lead of Nov. 2021 launch 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, earning \$1 million in revenue in year 1
  - Identified and optimized compatible ddPCR assays using web scraping in R with rvest for cfDNA quality control
  - Used Python to author automated data / QC reports using Jinja2 (with CSS) and WeasyPrint
  - Managed data and ran bioinformatic analyses on Illumina RNA & DNA seq using bash tools like: awk, grep, scp, wget, Illumina BaseSpace CLI, ssh, bwa, bedtools, samtools, vcftools, UMI-tools/fgbio
  - Wrote original SQL queries for Snowflake-based database to answer production and business questions.
- 2021  
|  
2019  
**Scientist I, NGS & NPI-Build**  
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 notebook for autonomous NGS data exploration
  - Coded build and QC experiments using Zymergen's Drawbridge alembic API Python library to LIMS
  - Rendered strain build and plasmid build reports from LIMS SQL database in MySQL Workbench



## SELECTED INTELLECTUAL PROPERTY

4/9/21



### **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

3/25/21



### **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

1/4/24



### **Twist cfDNA Pan-Cancer Reference Standard v2 Technical Guidance**

Twist Bioscience [Product Sheet](#) & [FAQ](#)

📍 South San Francisco, California

- Patrick Cherry, Lydia Bonar, & Mike Bocek

4/22/22



### **Characteristics and specificity of the wild-type / 0% VAF reference material**

Twist Bioscience

📍 South San Francisco, California

- Patrick Cherry & Mike Bocek

3/19/19



### **Multiple decay events target HAC1 mRNA during splicing to regulate the unfolded protein response**

[eLife](#)

📍 Aurora/Denver, Colorado

- Cherry, P., Peach, S., & Hesselberth, J.

12/6/17



### **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

2/6/24



### **Twist pan-cancer reference standard V2: Enhanced precision and reduced errors in ctDNA analysis**

[Advances in Genome Biology and Technology](#)

📍 Orlando, Florida

4/16/23



### **Twist pan-cancer synthetic RNA fusion control for assay development**

[American Association for Cancer Researchers](#), [Twist Bioscience](#)

📍 Orlando, Florida

2/7/23



### **Use of synthetic CNV fragments to mimic copy number alterations for ctDNA reference standards**

[Advances in Genome Biology and Technology](#)

📍 Hollywood, Florida

4/12/22



### **Twist pan-cancer synthetic reference materials for cell-free DNA (cfDNA) assay development**

[American Association for Cancer Researchers](#)

📍 New Orleans, Louisiana

6/16/20



### **R use at Zymergen**

Z-Tech Talk

📍 Emeryville, CA

4/20/20



### **Data-driven troubleshooting of NGS experiments**

Data Science Talk

📍 Emeryville, CA

8/24/17



### **RNA processing regulates the unfolded protein response**

CSHL: mRNA Processing Meeting Talk

📍 Cold Spring Harbor, New York