

PATRICK CHERRY

PhD scientist skilled in bioinformatics, biological data science, statistical modeling, next-generation sequencing (NGS), and tool-building. I've coded high-throughput experimental designs and genomic analyses, launched best-in-class oncology reference standards, and invented new molecular methods for DNA and microbe manipulation. Originally trained in Molecular Biology, I'm interested in taking my knowledge and practice of data science and bioinformatics to the next level.

EDUCATION

2019
|
2013

PhD

University of Colorado School of Medicine

📍 Aurora/Denver, Colorado

- Ph.D. in Molecular Biology
- PI: Jay Hesselberth, PhD.
- Thesis on RNA Terminus chemistry affecting the decay events that target HAC1 mRNA during the Unfolded Protein Response

2013
|
2009

BA

Hendrix College

📍 Conway, Arkansas

- Biochemistry and Molecular Biology, with Distinction
- PI: 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
|
2022

Senior Scientist

Twist Bioscience

📍 South San Francisco, California

- Tech Lead of multiple reference control NPIs, custom OEM projects, and commercial release OEM. Includes the *Pan-cancer RNA Fusion Controls*, *Fragmentome Calibration Standards*, *CNV Controls*, *Pan-cancer cfDNA v2*, and RNA-seq.
- Mentored a direct report from Senior Research Associate to Scientist to serving as a Tech Lead on new product introductions
- Original research led to multiple outside presentations and applications for patent protection of product configuration and biochemical methods.
- Custom analysis demonstrated proof-of-concept design and QC success of the Pan-cancer RNA Fusion Controls; designed and implemented the production approach; used public data and feedback from alpha testers to design configuration of fusions RNAs
- Led technical evaluation of a new NGS platform with custom experiments and bioinformatic analyses to enable faster gene QC in Production. Also led ancillary experiments to speed up production. Coded, implemented, and distributed on company GitHub an internal package, *twistcolorpal*, that automatically adds Twist-brand colors to ggplot2 plots and sets up database connectors to SQL / Snowflake for parameterized dbplyr querying.

CONTACT

✉ pcherry [at] pm.me
☎ upon request
🏢 Senior Scientist | Genomics
🏢 Twist Bioscience
📍 San Francisco, California
🐙 github.com/pdcherry
🌐 linkedin.com/in/p-cherry

I currently split my time between wet lab and computational activities. I have worked in a variety of roles ranging from HTP strain onboarding to genomics scientist. I like collaborative environments where I can learn from my peers and in turn teach others.

Last updated on 2023-12-12.

*Data-driven résumé made in R using
pagedown.*

Current
|
2021



Scientist

Twist Bioscience

📍 South San Francisco, California

- Tech Lead of *Pan-Cancer Reference Standard*, an ISO-13485 synthetic positive control with 458 unique variants among 84 cancer-associated genes at six QC'd VAFs, plus a WT control, which launched in Nov of 2021
- Designed, implemented, & validated primer removal procedure for DNA standards *that is compatible with methylation*
- Devised precise high-throughput DNA quantification process for accurate pooling
- Multiple iterations of custom NGS analysis refined the QC approach and thresholds for ensuring a contamination-free production process.
- Made extensive use of UMI sequencing and novel method to precisely quantify conversion efficiency to evaluate product and potential secondary sources

2021
|
2019



Scientist I

Zymergen, Inc.

📍 Emeryville, California

- Designed and implemented an automated high-throughput genotyping assay
- Designed & carried out complex experiments on automation with and without LIMS sample tracking
- Supported a company-wide NGS core under high demand from diverse groups with complex needs using data-driven decision making and teaching
- Used statistical methods to screen and optimize a genetic engineering protocol for newly-on-boarded microbe; delivered robust process while working on New Product Introduction team
- Built hundreds of plasmids using modern molecular cloning techniques like Gibson and Golden Gate



RESEARCH EXPERIENCE

2019
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2014



Doctoral Research

University of Colorado School of Medicine

📍 Aurora/Denver, Colorado

- Wrote, revised, & published two academic papers on RNA repair & yeast genetics
- Engineered and characterized genetic bypass of essential genes in budding yeast; on-boarded CRISPR/Cas9 for efficient and precise gene knock-in
- Expressed, purified, and used wild-type and mutant recombinant protein in *E. coli* to carry out an RNA modification enzymatic assay
- Optimized custom RNA-seq library protocol; independently planned, executed, troubleshooted RNA modification detection
- Routinely conducted northern blotting, targeted depletion, primer extension, splinted ligation, and other esoteric DNA and RNA experiments

I worked on a few projects during my PhD, and the RNA repair project led me to custom 5'-OH RNA-seq libraries, which inspired my fascination with transcriptomics and bioinformatics.



INTELLECTUAL PROPERTY

3/7/23



Methylation-mediated adapter removal on nucleic acid sequences

Twist Bioscience

📍 South San Francisco, California

- US 63/317,466

11/12/21





Expansion of cfDNA for Libraries

Twist Bioscience

📍 South San Francisco, California


- US Prov. Pat. Ref No 823.102

Working at Twist and Zymergen on new product research requires confidentiality, but public evidence of accomplishments often comes in patent applications. The Legal teams know me well for being a helpful expert in the process.

- 4/9/21 ● **Libraries for mutational analysis**
Twist Bioscience  South San Francisco, California
• US Prov. Pat. Ref No 823.101
- 3/25/21 ● **Method for counterselection in microorganisms**
Zymergen, Inc.  Emeryville, California
• US 2021_0087586 A1










SELECTED PUBLICATIONS

- 12/21/21 ● **Twist Pan-cancer synthetic reference materials technical guide**
[Twist Bioscience](#)  South San Francisco, California
• Patrick Cherry & Mike Bocek
- 2019 ● **Multiple decay events target HAC1 mRNA during splicing to regulate the unfolded protein response**
[eLife](#)
• Cherry, P., Peach, S., & Hesselberth, J.
- 2018 ● **Genetic bypass of essential RNA repair enzymes in budding yeast**
[RNA](#)
• Cherry, P., White, L., York, K., & Hesselberth, J.



SELECTED PRESENTATIONS & TALKS

- 11/19/23 ● **High sensitivity detection of specific ultra low-frequency somatic mutations for minimal residual disease (MRD) monitoring**
International Society of Liquid Biopsy Annual Congress  Madrid, Spain
- 4/16/23 ● **Twist pan-cancer synthetic RNA fusion control for assay development**
American Association for Cancer Researchers  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
- 7/13/21 ● **Molecular Methods Meet the Standards: Or how I learned to stop worrying and love UV-quantification**
Twist R&D Meeting  South San Francisco, CA
- 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

I communicate my results clearly, both in writing and in live presentations. I enjoy writing research papers, but my career has required more tech notes and app notes recently.

I give audience-centered presentations by adapting on the fly and over time to the venue and occasion. I like to *transfer knowledge* by giving methods/best practices talks whose slides can also serve as documentation.