Ryan Najac

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Experience

Columbia University – Institute for Cancer Genetics

Senior Research Technician, Palomero Lab

- Manage flexible, high-performance cloud-based computational infrastructure for bioinformatic analyses • Build automated workflows for multi-omic sequencing data analysis, enabling batch processing of up to 1,000
- samples simultaneously to accelerate the discovery of clinically-relevant biomarkers in leukemia and lymphoma • Prepare bulk and single-cell sequencing libraries (RNA-seq, ChIP-seq, ATAC-seq), perform quality control assays, and operate Illumina sequencers to generate high-quality data for downstream analyses.
- Develop a tiered data compression system using SAM-BAM-CRAM pipelines integrated with Amazon Web Services to optimize storage efficiency and reduce long-term archival costs of over 200 TB of sequencing data

Hvannis Port Research (HPR)

Software Engineer Intern

- Implemented a low-latency parser for the Nasdaq TotalView-ITCH data feed using the C standard library
- Developed simulated data feeds and utilized profiling tools to validate parser functionality, optimize performance, and resolve memory leaks, resulting in improved error handling and elimination of bottlenecks
- Collaborated with network engineers to develop a custom TCP/IP protocol that can recover corrupted data packets

Columbia University – Institute for Cancer Genetics

Research Technician, Iavarone/Lasorella Labs

- Designed a standard operating procedure for single-nucleus sequencing of archival glioblastoma samples
- Performed sub-cranial tumor implantations in mouse models to evaluate the efficacy of novel therapeutic agents
- Created and validated a high-throughput batch analysis pipeline for fluorescence microscopy data

Education

Columbia University

Bachelor of Arts, Computer Science

- Relevant Coursework: Operating Systems, Computer Networks, Data Structures and Algorithms, Software Design, Advanced Programming in the Linux Environment, Probability and Statistics
- Notable Projects: C HTTP web server, Linux file system and scheduler implementations, C-like language compiler

Technical Skills

Programming Languages:

- Proficient in C, C++, Python, VimL, and Linux shell script (sh, bash, POSIX)
- Some experience functional programming in Ocaml and object-oriented programming in Java
- Working knowledge of SQL database management and web development with HTML, CSS, JavaScript, and PHP

Frameworks and Software:

- Cloud computing and data management using Amazon Web Services (AWS): EC2, S3, Batch, and Lambda
- Software testing and debugging with Unity, GoogleTest, pytest, GDB, Valgrind, and Wireshark
- Data analysis and visualization using R, pandas, ggplot2, and Matplotlib
- Technical documentation and scientific writing using Jupyter notebooks, Rmarkdown, and LATEX

Publications

- "Integrative multi-omics networks identify PKC δ and DNA-PK as master kinases of glioblastoma subtypes and guide targeted cancer therapy." Migliozzi, S. et al. Nature Cancer, 4, 181-202, 2023. DOI
- "Pathway-based classification of glioblastoma uncovers a mitochondrial subtype with therapeutic vulnerabilities." Garofano, L. et al. Nature Cancer, 2, 141-156, 2021. DOI

New York, NY September 2022 - Present

> New York, NY May 2024

Needham, MA

May 2023 - August 2023

New York, NY

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