Natalie DeForest, PhD

Bioinformatics Scientist & Statistical Geneticist

EMAIL > nataliedeforest@gmail.com WEBSITE > nataliedef.github.io

PHONE \$ (510) 378-1641

EDUCATION

PhD Biomedical Sciences, Specialization in Bioinformatics, University of California San Diego Dissertation title: "Leveraging human genetics and functional genomics to investigate insulin resistance disorders"

BS Pharmaceutical Chemistry, Minor in Bioinformatics, University of California Davis June 2018 summa cum laude. Major/Minor GPA: 3.9

RELEVANT EXPERIENCE

Senior Scientist - Computational Genomics, *Fauna Bio* (Emeryville, CA)

Feb 2024 – Present

Identify and prioritize translatable genetic targets responsible for protection in animal models of disease resistance using a variety of omics data (i.e. bulk/single-cell RNA-seq, large-scale human genomics datasets including UK Biobank)

Postdoctoral Scholar, *Majithia Laboratory, School of Medicine, UC San Diego* (La Jolla, CA)

Nov 2023 – Jan 2024

Graduate Student Researcher, Majithia Laboratory, School of Medicine, UC San Diego (La Jolla, CA) June 2019 – Nov 2023 Integrate high-throughput genomic screens, diverse omics datasets, and large-scale human genetic databases to prioritize novel therapeutic targets for prevalent metabolic disorders such as diabetes and cardiovascular disease.

Research Intern, Gilead Sciences (Foster City, CA)

June 2018 – Sept 2018

Research Intern, Cytokinetics (South San Francisco, CA)

June 2017 - Sept 2017

Undergraduate Honors Researcher, UC Davis Dept. of Chemistry (Davis, CA) Aug 2016 - June 2018 Honors Thesis title: Chemoenzymatic Synthesis of Sialyl Lewis X, A Biologically Important Tetrasaccharide, and the Cloning and Characterization of Enzymes for Carbohydrate Synthesis

Clinical Data Intern, Pharmacyclics, an AbbVie Company (Sunnyvale, CA)

June 2016 – Aug 2016

SKILLS

Technical:

General: Statistical / population genetics (GWAS, omicQTLs,

fine-mapping, colocalization, burden testing, Mendelian Randomization, PheWAS)

- Next generation sequencing & bioinformatics workflows (bulk/single-cell RNA-seq, nextflow)
- Scripting languages (R, Python), Linux and command line interfaces (bash), high performance and cloud computing (AWS, GCP), and git version control
- Mining and analyzing relevant biological databases (GTEx, ENCODE, Ensembl, GEO, SRA)
- Familiar with machine learning models and applications, adept in **prompt engineering** for utilizing large language model tools

- Industry experience with drug discovery and **development** in both early and late stage companies
- Experienced in leading and collaborating with multi-disciplinary teams comprised of computational data scientists, experimentalists, and clinicians
- Adept in perusing scientific literature and understanding emerging studies
- Strong track record of scientific publications, conference presentations, and mentoring
- Driven, creative problem solver

SELECTED PUBLICATIONS & PRESENTATIONS

- **DeForest N.** et al. Genome-wide discovery and integrative genomic characterization of insulin resistance loci using serum triglycerides to HDL-cholesterol ratio as a proxy. *Nature Communications*. Sept 14, 2024. 10.1038/s41467-024-52105-y
- **DeForest N.** et al. Human gain-of-function variants in HNF1A confer protection from diabetes but independently increase hepatic secretion of atherogenic lipoproteins. Cell Genomics. May 30, 2023.10.1016/j.xgen.2023.100339.
- **DeForest N.** Activation of PPARG in skeletal muscle and visceral adipose tissues ameliorate NASH biomarkers in humans: implications for therapeutic targeting. Presented at NASH Keystone Conference, Whistler, Canada, Aug 2022
- **DeForest N**, Majithia AR. Genetics of Type 2 Diabetes: Implications from Large-Scale Studies. *Current Diabetes* Reports. Mar 19, 2022. 10.1007/s11892-022-01462-3.
- Du X, DeForest N, Majithia AR. Human Genetics to Identify Therapeutic Targets for NAFLD: Challenges and Opportunities. Frontiers in Endocrinology. Dec 7, 2021. 10.3389/fendo.2021.777075.

CERTIFICATIONS / AWARDS

- Machine Learning Specialization, DeepLearning.AI + Stanford University, Coursera
- T32 National Research Award, National Institute for General Medical Sciences (NIGMS)

Dec 2024

June 2020