# Sedat Demiriz

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## Experience

#### Computational Biologist - Sequence Bio Inc.

Lead the development of GWAS and ROH analysis pipelines with a focus on scalability and reproducibility Derived disease associations by analyzing data from company's Newfoundland population cohort Combined public and company datasets to derived insights into founder population genetics signature

#### Research Programmer - BC Genome Sciences Center

Performed quality control of incoming clinical oncopanel wet lab data by NGS data quality metrics Streamlined existing quality control and visualization scripts and configured for reusability Assisted graduate students with applying computational approaches to research questions

#### Biosensor R&D Co-op Programmer - CiBER Lab, SFU

Automated sensor response data processing and generation of easily digestible visualizations

Performed collection of datafrom hundreds of sensor instances during development

Presented and implemented metrics for determining sensor instance success based on response pattern

#### Genetics Co-op Student - Istanbul University Genetics Institute

Carried out clinical PCR, DNA and RNA isolation and sample quality control procedures Worked routinely in cell culture laboratory to grow five varieties of tumnor cell populations for lab use Lead nad supported four other co-op students during their first month with the lab

### Skills

Experience implementing robust genomics data processing workflows using pipeline utilities

Experience working with general population cohort data, identifying phenotype-genotype associations

Fluent in Python, R, and Bash for data analysis, transformation and visualization tasks

Applying good software development practices: version control, unit testing, documentation

Querying SQL databases and unstructured public health data sources

#### Education

#### BSc Molecular Biology and Biochemistry & Computing Science

Computing Science: Molecular Biology: Statistics:

#### **Publications**

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