

Bioinformatics Core and Consultation Services

Overview

Bioinformatics rapidly field growing is seeks apply computational and data that to science tools to the management, analyses, processing, and storage of biological data and health records. As the amount of genetic and genomic data continues to doable approximately year; additionally, the variety every experimental methods that yield large volumes of data in need of analysis, increases as well. Absolute Genomics is here to provide end-to-end analysis a wide variety of genetics and solutions for genomics projects.

Data Transfer

For secure HIPAA compliant data transfer AWS Genomics services provides secure data transfer from you institution to our company

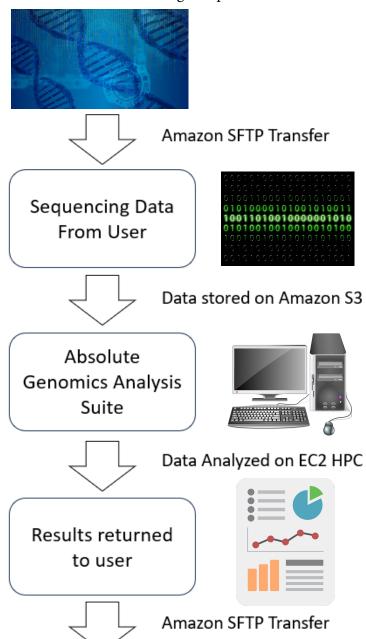
Analyses

cloudemploying and scalable By secure (AWS) resources offer based we variety of analyses routines to meet your research needs, including, but not limited to, analysis pipelines for:

- WGS variant calling and annotation
- WEX variant calling and annotation
- TEX variant calling and annotation
- 16S microbiome sequencing (16S) phylogenetics
- (Differential) ChIP-Seq occupancy analysis
- ATAC-/MNase-Seq nucleosome occupancy analysis
- (differential) RNA-Sequencing (RNA-Seq) analysis
- Chromosome Conformation Capture Analyses
 - -3C, 4C, 5C, HiC
- Single cell RNA/RNA+ATAC sequencing analyses
 - -Clustering
 - -Differential Expression
 - -Trajectory Analysis
 - -Cell Type Annotation
- DNA methylation and MicroArray Analyses

1/2 the Cost in 1/2 the Time* - Results Within 24 Hours

*over leading competitors



Consultation

Our lead Bioinformatician, Dr. Mariani, has extensive experience in computational genomics, bioinformatics, statistical genetics, and computer programming. In addition to our standard workflows, consultation services are available. We are here to meet your bioinformatics and computational genomics needs!