Director: Prof. Konstantinos Krampis
Email: agbiotec@gmail.com
Office: Rm. 467F Belfer Research Building
Phone: (212) 396-6930
Fax: (212) 650 3565

Facility Manager: Carlos Lijeron
Email: clijeron@hunter.cuny.edu
Office: Rm. 400 Belfer Research Building
Phone: (212) 396-7018

Bioinformatician: Baekdoo Kim
Email: Baekdoo.Kim59@myhunter.cuny.edu
Office: Rm. 400 Belfer Research Building
Bioinformatics and Sequencing Facility - Biology

Bioinformatician: Thahmina Ali
Email: Thahmina.Ali62@myhunter.cuny.edu
Office: Rm. 400 Belfer Research Building

Network Advisor Main Campus
Dr. Lloyd Williams
Email: williams@genectr.hunter.cuny.edu

Description of the Facility

Background Overview
The Hunter College/CTBR Bioinformatics resources is located on the 4th floor of the Belfer
Research Building at 69th Street and York Ave. The facility affords access to researchers and faculty, a high-performance computer cluster with a large range of bioinformatics software and data analysis pipelines. The facility provides cutting-edge bioinformatics technology for translational and basic research on health disparities. We also host a web-accessible bioinformatics platform based on Galaxy, (http://galaxy.hunter.cuny.edu:8080) to support genomic sequencing analysis. Additionally, the facility offers Illumina Sequencing using the Illumina MiSeq sequencing platform and Nanopore sequencing using Oxford Nanopore MinIon sequencer. Both these instruments are capable of sequencing entire complement of DNA, or genome, of many animal, plant, and microbial species for basic biological and medical research. A detailed description of our services and available equipment is given below

**Services**

- RNAseq and variation discovery
- small RNAs sequencing
- de novo bacterial genomes
- RNAseq Analysis
- Targeted amplicon sequencing
- Computational Capacity
- Scalable Storage

**Bioinformatics and Sequencing Resources and Equipment**
**Illumina MiSeq**

MiSeq desktop sequencer:
Allows narrowly focused applications such as targeted gene sequencing, metagenetics, metagenomics, small genome and transcriptome sequencing, targeted gene expression, and amplicon sequencing.

**Oxford Nanopore MinIon sequencer**

Nanopore (real-time sequencing):
MinIon portable sequencer: provides a rapid and portable, real-time sequencing platform that includes sequencing of full length transcripts with long reads, haplotype sequencing, metagenomic and 16S sequencing.

**Agilent Technologies, 2100 Electrophoresis Bioanalyzer**

The Agilent 2100 Bioanalyzer is a microfluidics-based platform that provides sizing, quantitation and quality control of DNA, RNA, proteins and cells.
Two assay principles - electrophoresis and flow cytometry

**Galaxy Web-accessible Bioinformatics Platform**
We are running an installation of Galaxy, a web-based platform for data intensive biomedical research.

http://galaxy.hunter.cuny.edu:8080

**Silicon Mechanics, HPC Cluster System**

The high performance computing cluster provides 800 CPU cores, 3TB of high-speed RAM, a GPU Node, and dynamic resource allocation.

- Redundant Head Node, 12 CPU Cores, 64 GB RAM
- 10 Compute Nodes, 20 CPU Cores each, 128 GB RAM
- 1 Medium Memory Node, 32 CPU Cores, 512 MB RAM
- 1 High Memory Node, 32 CPU Cores, 1 Terabyte RAM
- 1 GPU Node, K80, 2 CPU Hyper-Threaded / 128 GB RA
Seagate Lustre CS1500
ClusterStor 1500 solutions feature scale-out storage building blocks, the Lustre® parallel filesystem and:
- 362TB of parallel storage
- 5GB/s throughput
- Seagate Enterprise Lustre
- Parallel based storage

Belfer E-Box
The Belfer E-box provides storage for data backup and project archiving.
- 200TB of high availability storage
- 5GB/s throughput

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