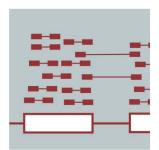




REGISTRATION AWARDS AVAILABLE

RNA-seq Analysis



Date

June 17-19, 2024 - Toronto, Ontario

Faculty

Malachi Griffith, Obi Griffith

Course Description

High-throughput sequencing of RNA libraries (RNA-seg) has become increasingly common and has become the industry standard for transcriptome profiling. When processed appropriately, RNA-seq data has the potential to provide an exceptionally richview of the transcriptome. The CBW has developed a 3-day course providing an introduction to bulk RNA-seg data analysis followed by integrated tutorials demonstrating the use of popular RNA-seq analysis packages. The tutorials are designed as self-contained units that include example data (Illumina paired-end RNA-seq data) and detailed instructions for installation of all required bioinformatics tools (HISAT, StringTie, Kallisto, etc.).

Sponsors











For registration details visit our website:

www.bioinformatics.ca