

Early Application Fee Before
MAY 14th
249\$ + Tax

Late Application Fee is
an additional \$100

Limited to **24** participants
per workshop

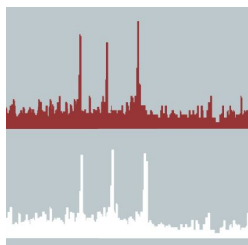


ONLINE

Canadian Bioinformatics Workshop Series

Registration Awards Available

Epigenomic Data Analysis



Dates

October 22 - 23, 2020 - Live Online starting at 10:00 AM EDT

Faculty

Guillaume Bourque, Martin Hirst, David Bujold

Course Objectives

High-throughput sequencing of Chromatin-Immunoprecipitated libraries (ChIP-seq) and of bisulfite converted DNA (WGBS) have become increasingly common and have largely supplanted microarrays for chromatin and DNA methylation profiling. When processed appropriately, ChIP-seq data provides base-pair resolution representations of transcription factor DNA-binding events and nucleosome (histone) modifications genome-wide. Similarly, WGBS can provide a quantitative genome wide profile of cytosine methylation.

Sponsors



Canadian Centre for
Computational
Genomics

MiCoM McGill initiative in
Computational Medicine



compute | **calcul**
canada | canada

Join our mailing list:

To be notified when registration
are open. (~end of January)

<https://bit.ly/34ictnl>

www.bioinformatics.ca

