



PRESENTS

STATISTICS AND APPLICATION TO METHYLATION

**Application deadline:
16th October 2022**

Date: 27th of October 2022

Time: 12:00-13:00h and 14:00-17:00h

Location: CRG Training Center (PRBB Patio)

Number of onsite participants: 18

Trainer/s: Emanuele Raineri, Bioinformatics Development & Statistical Genomics Team, CRG-CNAG

We will look at the main bioinformatics concepts and software used to analyze DNA methylation data (focusing on Whole Genome Bisulfite Sequencing and Nanopore reads).

During the class we will write a tool to call differentially methylated loci and differentially methylated regions and compare different possible approaches.

Contact

training@crg.eu

More information

