

Miniworkshop on NGS Data Analysis



27 November (UG44A)

10.30 • Andrea Telatin

An introduction to 16S metabarcoding analysis

14.30 • Perla Troncoso Rey

Introduction to RNA-Seq bioinformatics analyses

28 November (UG55A)

10.30 • Prof. Nicola Vitulo

Interpreting data from metabarcoding experiments

15.30 • Prof. Chiara Romualdi

Notes on statistical methods for RNA-seq

27th - 28th November 2019

Chiara Romualdi (University of Padua) is a biostatistician with a decade long experience in statistical methods for transcriptome analysis.

Nicola Vitulo (University of Verona) is a bioinformatician currently working on a range of metagenomics projects.

Host: Andrea Telatin (Gut Microbes and Health)

Register here:

