

# 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.

Hosts: Andrea Telatin and Andrew Page

Register here:

