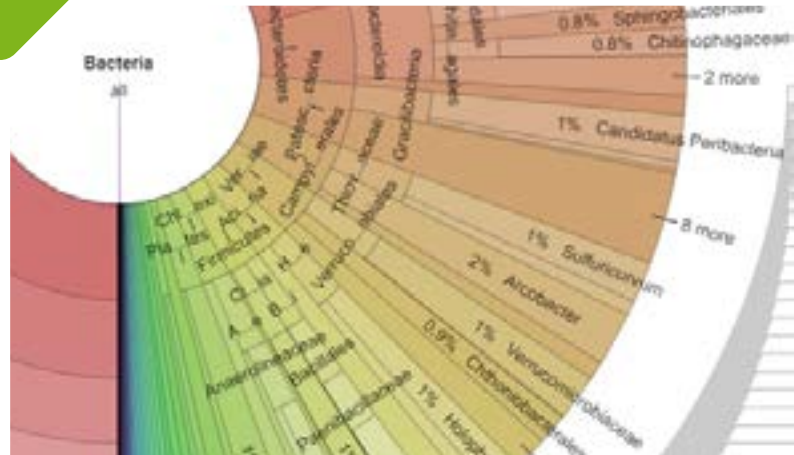




Microbiome and Metagenome Data analysis workshop



 1.–5. 4. 2019

 08:30–17:00

 České Budějovice
Czech Republic

 1 250 € without VAT

 English

 Seats available: 25

You will be briefly introduced into the Linux operating system and in this environment you will analyze sequencing data using various publicly available tools. All workshop participants will perform all steps of data analysis tasks themselves!

Both community analysis of amplicon sequencing data as well as analysis of whole genome shotgun sequencing data will be covered including methods for statistical comparisons of multiple metagenomic samples.

Who should attend?

This workshop is aimed at life scientists and bioinformaticians working or being interested in the field of metagenomics / microbiomics. No prior knowledge of metagenomic / microbiomics sequences analysis or specific programming skills are required.

Join the workshop

