

# Microbiome Sequencing

## from the 16S rDNA (V1-V9 regions)

### Number of samples

From 10 to 2 500 samples per run

### Depth

From 10 000 to 100 000 reads per sample

### Type of samples

gut, stool, soil, blood, urine, saliva, milk, water, cells, tissues

### Fast and robust results

6 weeks for 150 samples

OTUs classification

Statistic and diversity analysis

Analysis complete report

Data via a FTP server or by USB Drive

### Other options

Fungal Microbiome (ITS region)

