

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



From 10 to 2 500 samples per run

## Depth

From 10 000 to 100 000 reads per sample

#### Type of samples

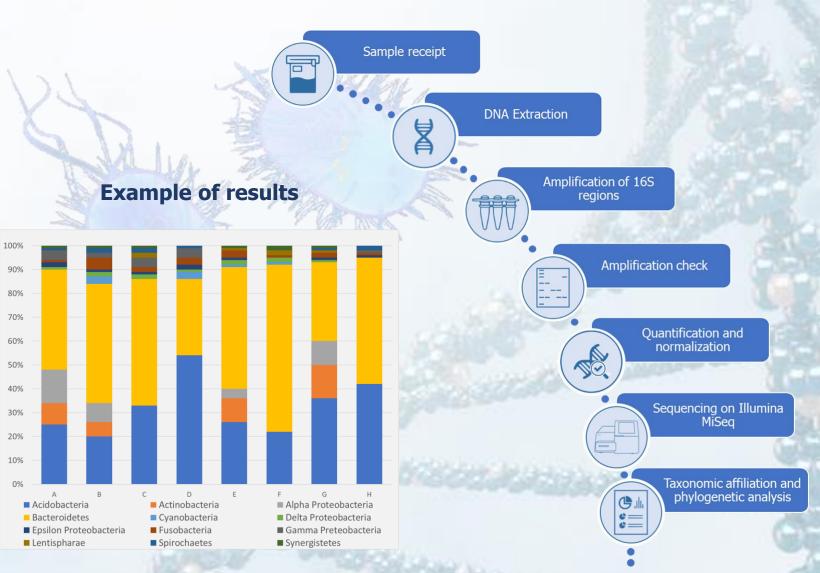
gut, stool, soil, blood, urine, saliva, milk, water, cells, tissues

### 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)



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