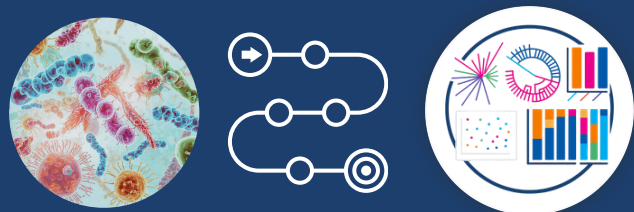


# Working with Microbiome Data?



If you work with **microbial communities**, you likely face these challenges:

- Which microbes are present in my samples – and in what abundance?
- How do community compositions shift across conditions or treatments?
- Which metabolic functions and pathways are enriched or depleted?
- How can I generate reproducible, publication-ready visualizations at scale?

This flyer shows which de.NBI tools can help analyse your microbiome data, step by step.

**de.NBI** Services in this flyer:

**MG-tk**



## de.NBI – Trusted Tools for Analysing Life Science Data

### Get started:

Training



Services



E-Learning



### What we Offer

- ✓ Curated tools for genomics, pangenomics, metagenomics, proteomics and more
- ✓ Training, documentation and expert support
- ✓ Access to the de.NBI Cloud
- ✓ Free of charge for academic users

[www.denbi.de](http://www.denbi.de)  
✉ [contact@denbi.de](mailto:contact@denbi.de)



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de.NBI – German Network for Bioinformatics Infrastructure  
Coordination Office: Forschungszentrum Jülich (FZJ)

## Microbiome Analysis Made Easy – with de.NBI Tools & Training

Free for academic life scientists

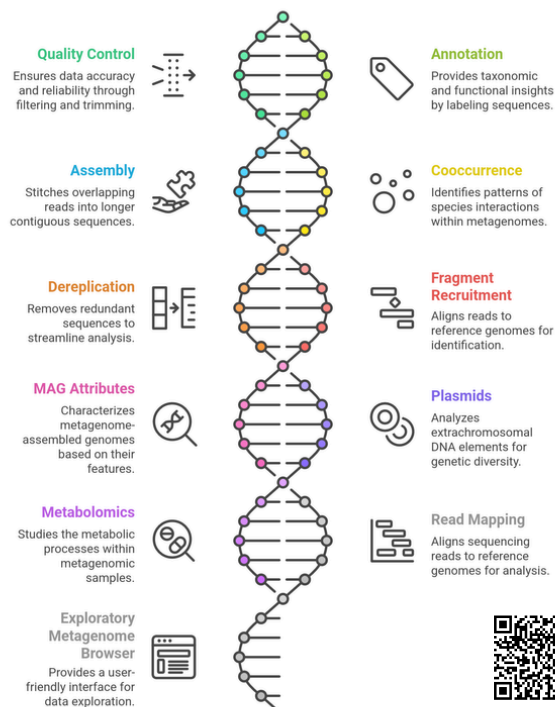


**de.NBI**  
GERMAN NETWORK FOR BIOINFORMATICS INFRASTRUCTURE

# Metagenomics-Toolkit

## MG-tk

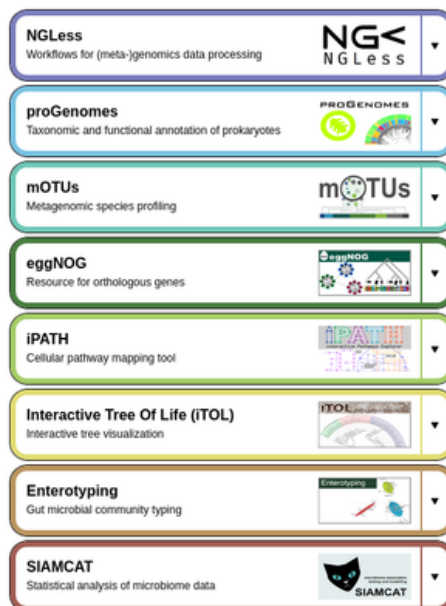
The Metagenomics-toolkit combines novel and established methods for taxonomic and functional microbiome profiling in a reproducible and modular workflow suite.



# Microbiome Analysis Toolbox



Tools and resources for analysing and interpreting microbiome data. From identifying and quantifying microorganisms to exploring cellular pathways, phylogenetic relationships and microbial community structure.



# Training Opportunities

## Metagenomics Training Courses

Hands-on workshops covering the full workflow from raw reads to biological insights and best practices for reproducible microbiome research.

## E-learning: Metagenomics Media Library

Self-paced learning with videos, practical examples and workflow guides to strengthen your skills anytime.

## Learn from the experts

All courses are developed and taught by researchers who build and maintain de.NBI tools – ensuring accuracy and relevance.

## Build your network

Training events offer direct exchange with fellow microbiome researchers across Germany, providing a platform for collaboration and peer support.



# microGalaxy



Browser-based workflows for microbial genomics and microbiome analysis – no installation needed, fully powered by the de.NBI Cloud.

- Easy access: Run analyses directly in the browser.
- Reproducible & scalable: Standardized workflows on de.NBI Cloud infrastructure.



Find a training course for microbiome analysis here:



To the de.NBI E-learning library:

