

## Microbiome Network Ecology & Co Occurrence — Hands-on

Learn how to build and interpret microbiome co occurrence and interaction style networks from amplicon and metagenomic data. This module walks through correlation and graphical models, community detection, keystone taxa and stability concepts so you can convert microbiome tables into ecological network insights for clinical, environmental and industrial studies.

# Microbiome Network Ecology & Co Occurrence

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### Session Index

[Session 1 — Microbiome Network Ecology Foundations](#) [Session 2 — Co Occurrence Network Construction Workflows](#) [Session 3 — Community Structure, Keystone Taxa & Stability](#) [Session 4 — Mini Capstone: Network Analysis of a Microbiome Cohort](#)

### Session 1

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## Microbiome Network Ecology Foundations

Networks in microbiome and microbial ecology studies

**nodes as taxa, functions and samples** **edges as associations and candidate interactions** **use cases in health, environment and industry**

Types of microbiome association networks

**correlation and covariance networks** **graphical**

**models and sparse inverse covariance** **time ordered**  
**and bipartite style networks**

Compositionality, sparsity and confounding in networks

**why naive correlations can mislead** **sparsity, zero**  
**inflation and prevalence filters** **batch effects and**  
**shared environment issues**

## **Session 2**

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### **Co Occurrence Network Construction Workflows**

Preparing feature tables for network analysis

**choosing taxonomic or functional resolution**  
**prevalence and abundance filtering rules**  
**normalisation and transform options**

Correlation and composition aware association methods

**Spearman and Pearson baselines** **SparCC style and**  
**related methods** **graphical lasso or SPIEC style**  
**approaches**

Thresholding, multiple testing and edge confidence

**p values, FDR and effect size trade offs** **symmetric vs**  
**directed edge choices** **exporting networks for**  
**downstream tools**

## **Session 3**

**Fee: Rs 14800** Apply Now

### **Community Structure, Keystone Taxa & Stability**

Network topology and community structure metrics

**degree, centrality and connectivity patterns**  
**community detection and modules** **hub, connector**  
**and peripheral nodes**

Identifying keystone and driver taxa candidates

**degree and betweenness based views** **module core species and connectors** **linking candidates to function and metadata**

Network stability, robustness and scenario style thinking

**simulated node removal and perturbations** **what if views for interventions and shifts** **bridging networks back to ecological narratives**

#### **Session 4**

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### Mini Capstone: Network Analysis of a Microbiome Cohort

End to end network analysis on a real microbiome dataset

**theory plus guided practical from feature table**

Comparing networks across groups or time points

**case vs control or treatment vs baseline** **topology changes and key edge differences** **simple visual summaries for stakeholders**

Deliverables: network files, plots and ecological summary note

**network edge and node tables for reuse** **graph visualisations and key metric tables** **PDF or HTML network ecology mini report**