

## Microbiome–Metabolome Integration & GNPS Bridges — Hands-on

Learn how to connect microbiome profiles with metabolomics features to build mechanistic, pathway aware stories. This module covers data structures for integration, GNPS and feature based molecular networking, linking taxa to metabolites, and summarising joint microbiome metabolome patterns for manuscripts and reports.

# Microbiome–Metabolome Integration & GNPS Bridges

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### Session 1

**Fee: Rs 8800** [Apply Now](#)

## Integration Concepts & Data Preparation

Why integrate microbiome and metabolome data

[ecology and function in one view](#) [complementary resolution of pathways](#) [case studies from gut and environment](#)

Data structures for multi omics integration

[feature tables \(ASVs, species, OTUs\)](#) [MS feature](#)

**tables and peak lists** **sample and batch metadata alignment**

Pre processing for each layer before integration

**normalisation and scaling principles** **batch effects and drift correction overview** **ensuring paired and matched samples**

### **Session 2**

**Fee: Rs 11800** Apply Now

## **Metabolomics Tables, GNPS & Molecular Networking**

From raw metabolomics to feature tables

**peak detection and alignment overview** **MS1 and MS2 feature information** **intensity matrices and metadata joins**

GNPS and feature based molecular networking concepts

**spectral libraries and cosine scores** **molecular families in networks** **exporting network and annotation tables**

Visualising molecular networks and feature clusters

**Cytoscape style layouts overview** **colouring by class or source** **overlying sample group information**

### **Session 3**

**Fee: Rs 14800** Apply Now

## **Linking Taxa, Features & Pathways**

Correlation and association between taxa and metabolites

**Spearman and partial correlations** **multiple testing and sparsity awareness** **simple network style visualisations**

Pathway mapping and enrichment with integrated data

**mapping features to KEGG or other resources**

**overlaying microbiome contributions** **interpreting joint pathway level shifts**

Basic multi omics integration approaches

**simple CCA and PLS style concepts** **block PCA and**

**correlation circle ideas** **linking integrated signatures to phenotypes**

#### **Session 4**

**Fee: Rs 18800** Apply Now

### Mini Capstone: Microbiome–Metabolome Story

Designing an integration analysis for one dataset

**guided theory plus practical**

Building joint microbiome metabolome summaries

**correlation heatmaps and networks** **pathway or**

**molecular family plots** **linking signatures to clinical or environmental traits**

Deliverables: integrated figure set and short narrative

**plots connecting taxa and metabolites** **summary**

**tables for key associations** **brief methods and results text for reports**