

## NMR Metabolomics — Acquisition, Processing & Annotation — Hands-on

Develop a complete, practice ready workflow for NMR based metabolomics. This module walks you from basic <sup>1</sup>H NMR concepts and experiment setup through acquisition, spectra processing, bucketing or peak picking, and metabolite annotation using curated spectral libraries and databases for robust quantitative and qualitative readouts.

### NMR Metabolomics — Acquisition, Processing & Annotation

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#### Session 1

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#### NMR Basics, Experiments & Metabolomics Context

Principles of NMR for metabolomics applications

[chemical shift & spin systems](#) [relaxation & line width](#)  
[sensitivity & dynamic range](#)

<sup>1</sup>H NMR experiments commonly used in metabolomics

[1D proton](#) [CPMG / T2 filter \(overview\)](#) [NOESY presat /  
water suppression](#)

Matrices, sample tubes and basic preparation for NMR

**biofluids (plasma, serum, urine)** **buffers & pH control**  
**deuterated solvents & references**

### **Session 2**

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## **Acquisition Parameters & QC for NMR Runs**

Setting up NMR acquisition for metabolomics

**spectral width & offset** **number of scans & relaxation**  
**delay** **temperature control**

Shimming, locking, tuning and water suppression checks

**lock optimization** **manual vs automated shimming**  
**water peak management**

Run QC and system suitability for NMR metabolomics

**reference compounds** **reproducibility metrics**  
**instrument drift awareness**

### **Session 3**

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## **Spectral Processing & Data Reduction**

Core processing steps from FID to spectrum

**zero filling & apodization** **Fourier transform** **phase &**  
**baseline correction**

Alignment, reference and solvent region handling

**chemical shift referencing** **alignment approaches**  
**(overview)** **water and urea region treatment**

Data reduction: binning vs peak picking strategies

**fixed width and adaptive binning** **peak lists** **export to**

**multivariate tools**

**Session 4**

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## Metabolite Annotation & Reporting

Annotation workflows with spectral libraries and databases

**HMDB / BMRB (concepts)** **in house libraries**  
**confidence levels**

Relative and absolute quantitation considerations in NMR

**integrals vs peak heights** **internal standards** **dilution**  
**and linearity checks**

Deliverables: processed dataset, annotation table & methods  
note

**bucket / feature matrix (CSV)** **metabolite annotation**  
**sheet** **NMR metabolomics methods summary**