

New technologies to address HIV Research

Metabolomics: Searching for insights to understand Immunological Non-response to ART

What's metabolomics?

The Need for Multi-Omics Biomarker Signatures in Precision Medicine
Overall, this information is more than the use of genetic variation information to inform clinical practice and the fact that the use of multi-omics approaches to achieve precision medicine in the clinic is extremely valuable but still in its infancy.

Applications of multi-omics analysis in human diseases
Integration of different types of omics data can elucidate underlying pathogenic changes of the disease, which can then be verified in further molecular biosamples. By integrating multi-omics, scientists can filter out noise and capture between biomolecules and disease phenotypes, identify relevant signaling pathways, and establish detailed biomarkers of disease. Therefore, the integration of various omics data will facilitate the match of associations between molecular disease and relevant environmental

Criteria for the use of omics-based predictors in clinical trials
Definition of 'omics'
In its report, Evolution of Translational Omics: Lessons Learned and the Path Forward, the Institute of Medicine Committee on the Review of Omics-Based Tests for Predicting Patient Outcomes in Clinical Trials defines 'omics' as the study of related sets of biological molecules in a comprehensive fashion. Examples of omics disciplines include genomics, transcriptomics, proteomics, metabolomics and epigenomics. An omics-based test is defined as "an assay composed of defined biomolecules (genetic, metabolic, proteomic) and translated by a fully specified computational model to produce a clinically actionable result."

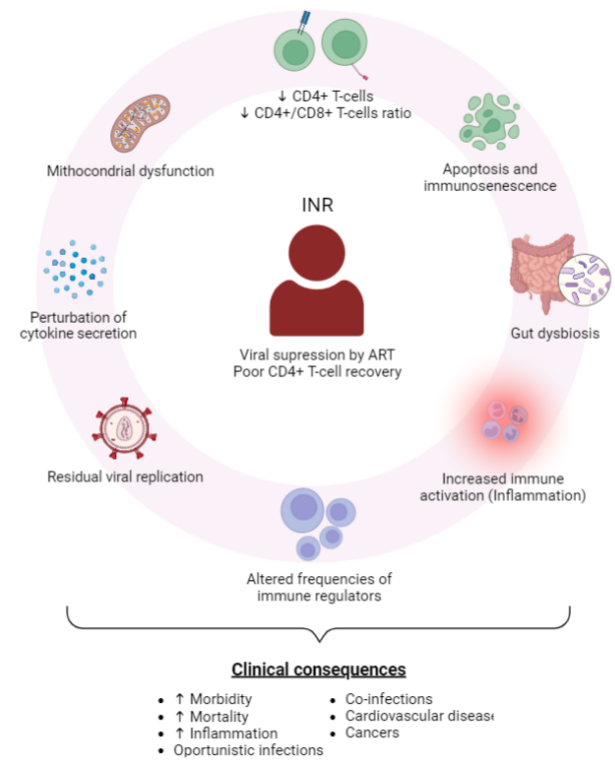
Omics technologies are high throughput techniques that make it possible to gather, in a single experiment, large amounts of data about a specific type of molecules, such as the three billion base pairs of the human genome, the universe of proteins in a given tissue or a large collection of metabolites. Examples of these technologies are next generation sequencing, used for genomics and transcriptomics studies, and metabolomics.

Omics-based biomarkers: current status and potential use in the clinic
Hector Quintado¹, Ana Laura Guzmán-Ortiz², Hugo Díaz Sánchez¹, Ricardo Valle-Rivera¹, Jesús Aguirre-Hernández¹

Practicing precision medicine with intelligently integrative clinical and multi-omics data analysis

Human Genomics

potential distinct results for patient care from highly expressed genes and disease-causing variants [9, 39]. Understanding how genetic variations contribute to health is one important aspect of precision medicine, where additional approaches involve measuring levels of proteins and metabolic products. By harnessing the power of metabolomics, we need to profile a patient's metabo-



Why is metabolomics important in the INR study?

Metabolomics involves the study of...?

- Lipids
- Alkaloids
- Amino acids
- Polymers

Sample preparation involves the following steps

- Extraction
- Processing
- Interpretation

To measure sets of metabolites, you can use

- Targeted metabolomics
- Untargeted metabolomics

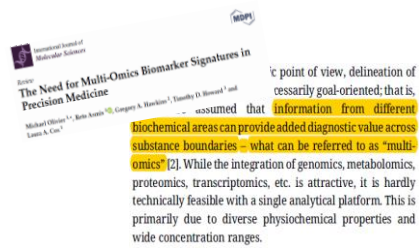
NMR is more sensitive than MS

- True
- False

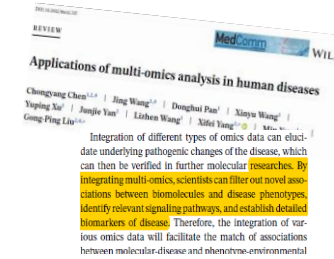
Metabolomics can help to improve healthcare

- True
- False

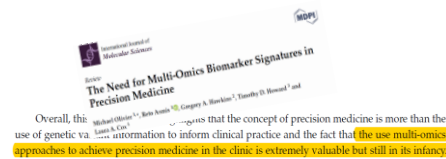
What's metabolomics?



How is metabolomics used?



Why is metabolomics important?



Omics technologies are high throughput techniques that make it possible to gather, in a single experiment, large amounts of data about a specific type of molecules, such as the three billion base pairs of the human genome, the universe of proteins in a given tissue or a large collection of metabolites. Examples of these technologies are next generation sequencing, used for genomics and transcriptomics studies, and metabolomics studies.



What's next for clinical applications of metabolomics?

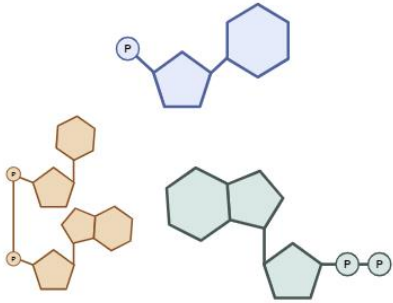
potential indistinct results for patient care from highly expressed genes and disease-causing variants [9, 39]. **Understanding how genetic variations contribute to health is one important aspect of precision medicine, where additional approaches involve measuring levels of proteins and metabolic products.** By harnessing the power of metabolomics, we need to profile a patient's metabo-

How to design a metabolomics study?

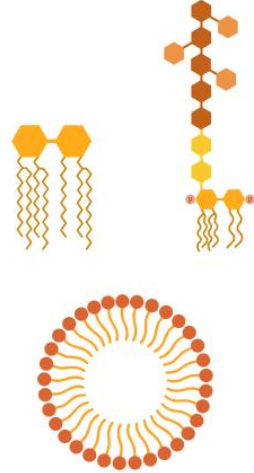
Omics-based biomarkers: current status and potential use in the clinic
Hector Quezada^{1*}, Ana Laura Guzmán-Ortiz^{1*}, Hugo Díaz-Sánchez², Ricardo Valle-Rios^{1*}, Jesús Aguirre-Hernández^{1*}

What's metabolomics?

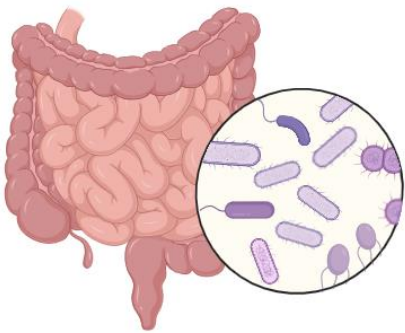
Metabolites



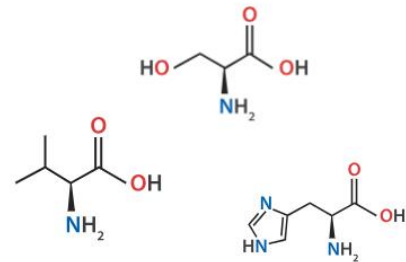
Lipids



Metabolites from microbiota



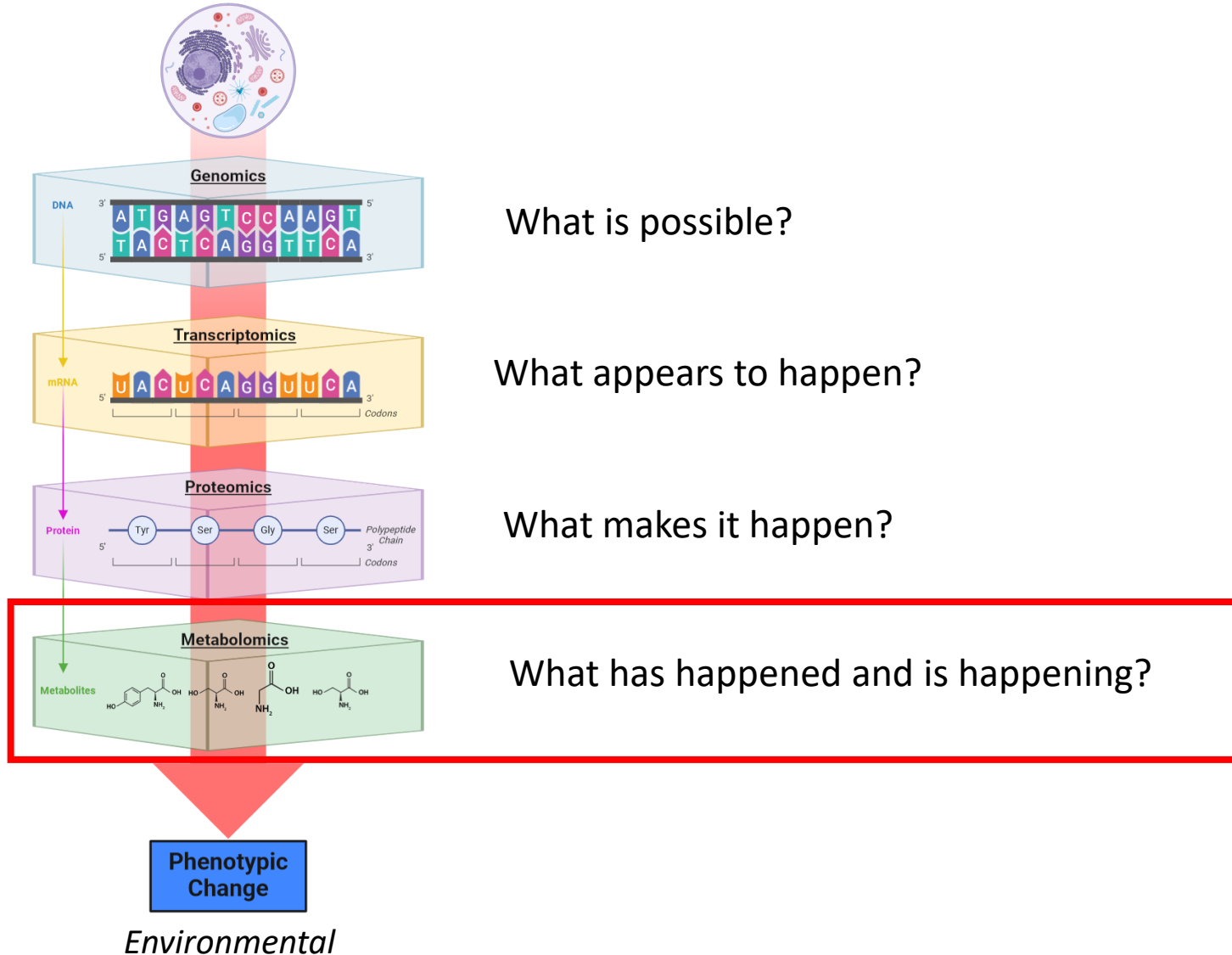
Amino acids



Comprehensive, qualitative, and quantitative study of low-molecular-weight molecules.

Examples of small molecules: sugars, lipids, amino acids, fatty acids, phenolic compounds, alkaloids, vitamins and many other types of molecules which are often the building blocks for larger compounds.

Why is metabolomics important?



What is possible?

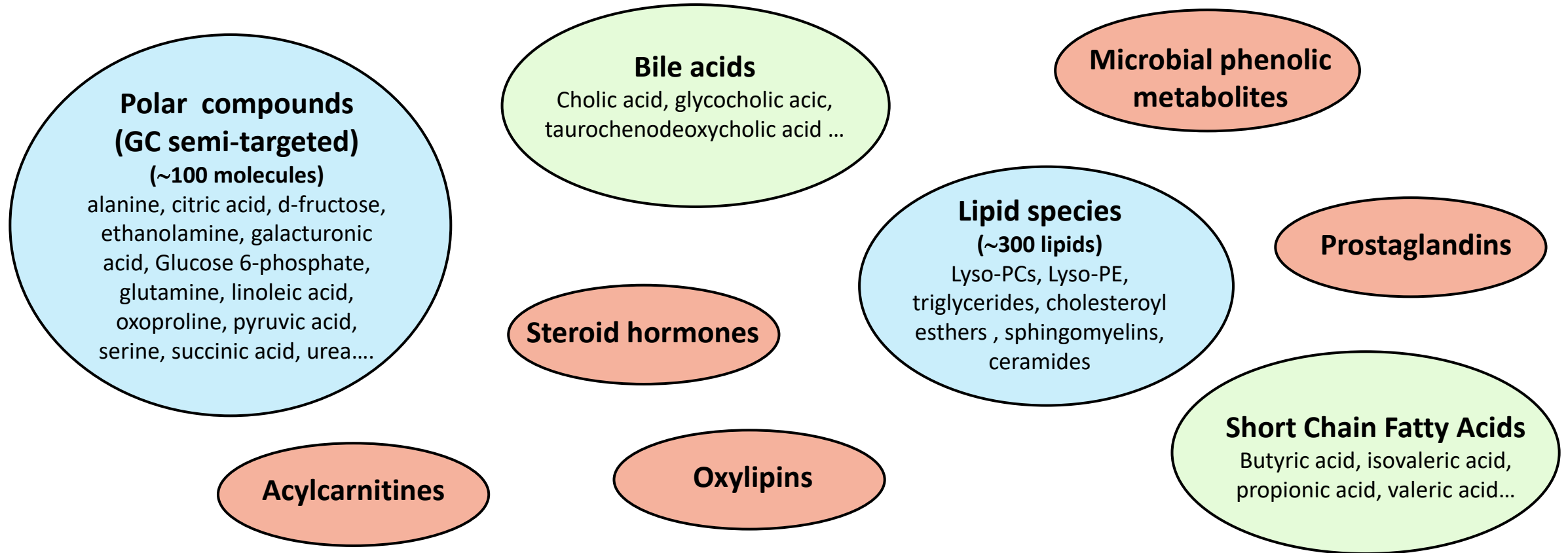
What appears to happen?

What makes it happen?

What has happened and is happening?

Snapshot of the physiology of the cell most closely related to the phenotype.

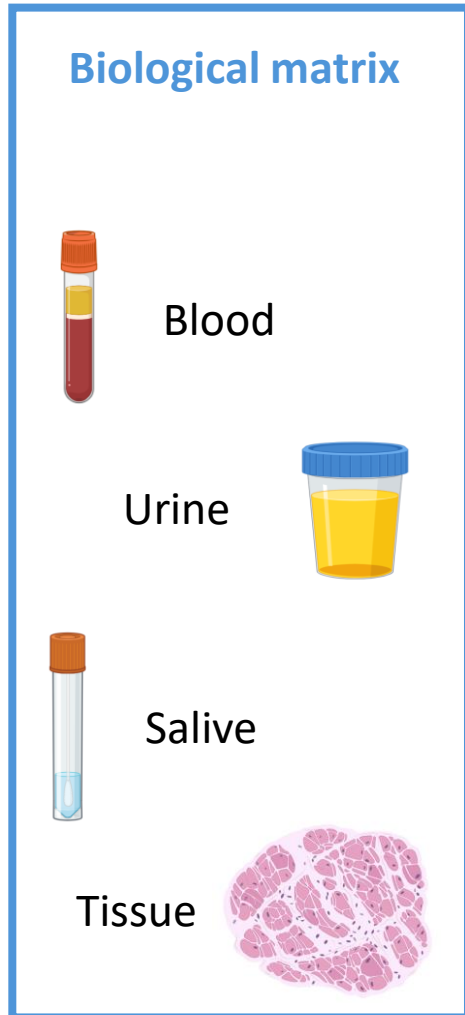
How to design a metabolomics study ?



Untargeted studies to identify a wide range of metabolites (profiling) to generate a hypothesis.

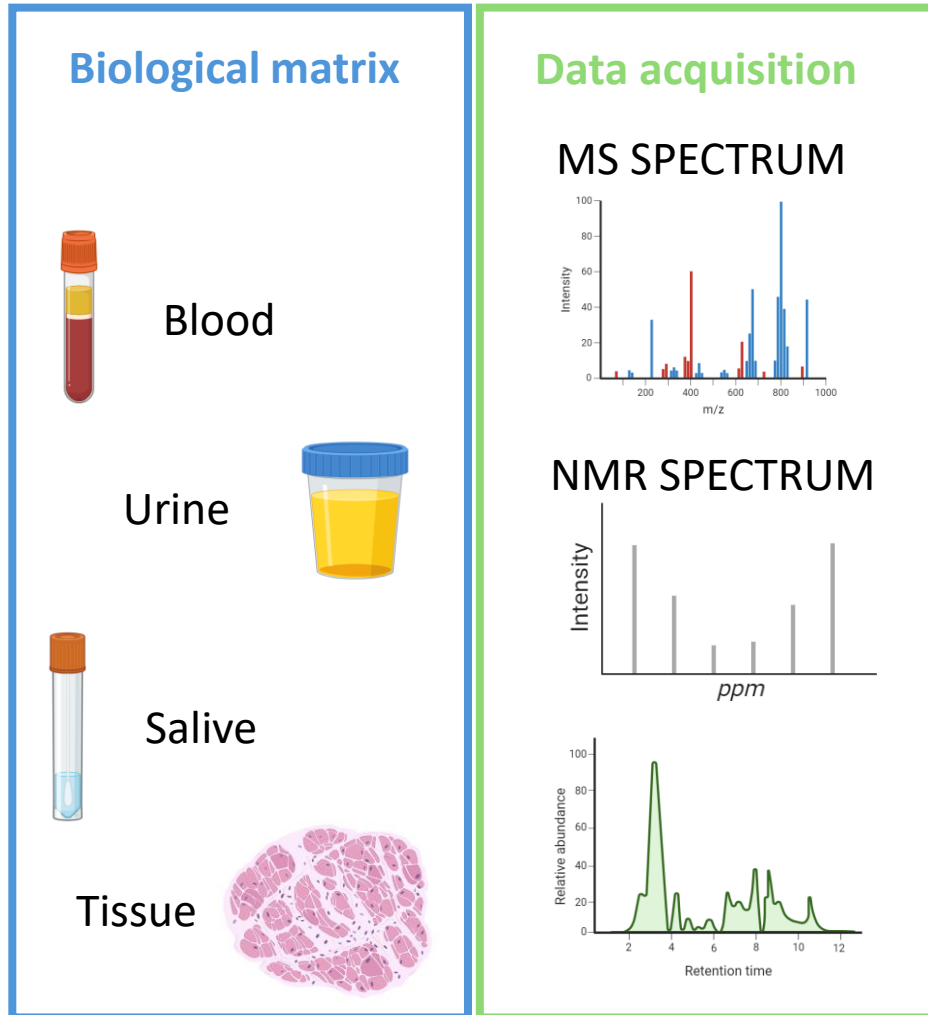
Targeted studies focus on a few specific metabolites when a hypothesis is already postulated.

How to design a metabolomics study?

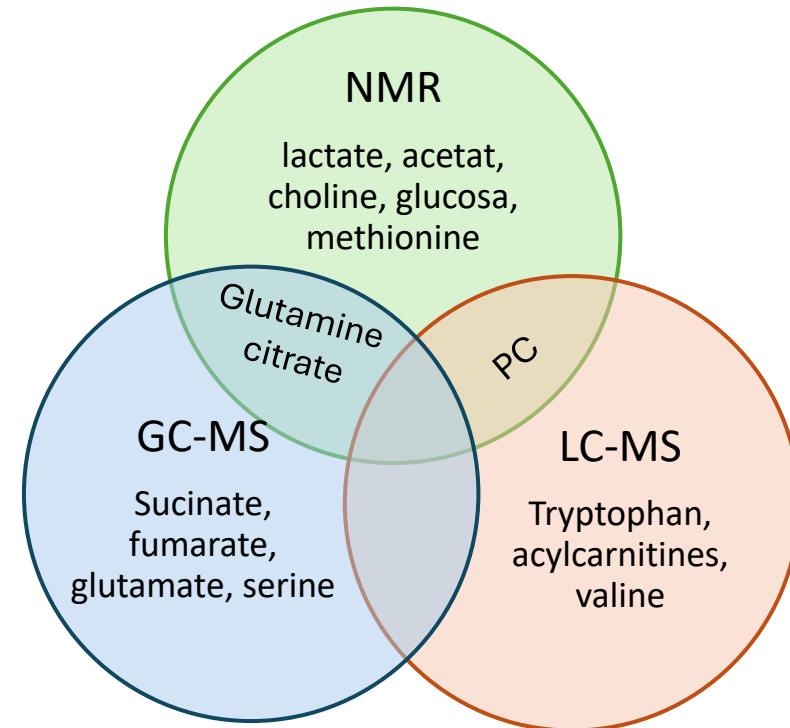


Similar pipeline for Targeted and Untargeted metabolomics studies but different handling and preparation depending on the study design (approach selected and technique required).

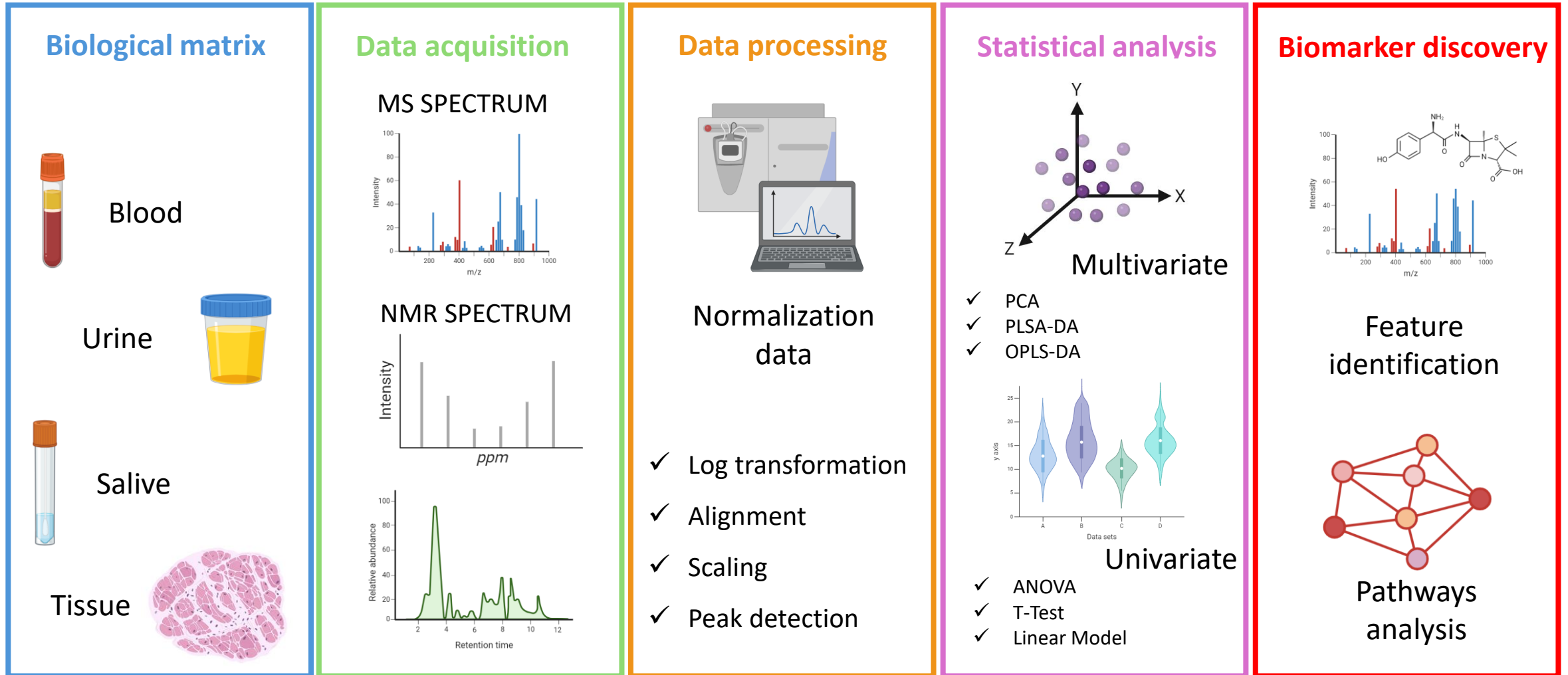
How to design a metabolomics study?



MS more sensitive and accurate and offer a high mass resolution than NMR



How to design a metabolomics study?



How is metabolomics used?

For diseases

- ✓ Risk assessment
- ✓ Screening
- ✓ Diagnosis
- ✓ Treatment
- ✓ Prognosis
- ✓ Monitoring

Technologies for identification

- ✓ NGS, Genome annotation
- ✓ Transcriptomics
- ✓ Proteomics: MS, Protein chips....
- ✓ **Metabolomics: NMR, MS....**

Biomarker contribution

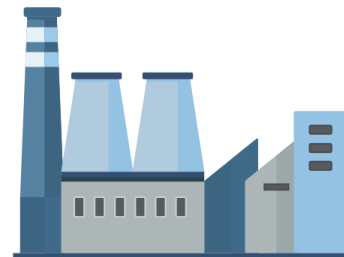
- ✓ Identification of events
- ✓ Develop a dose-response
- ✓ Determine variability and effects
- ✓ Correlated events with disease

Biomarker evaluation

- ✓ Safe and easy to measure
- ✓ Low cost of follow-up test
- ✓ Proven to treatment to modify the biomarker



ACADEMIC



INDUSTRY



CLINICAL

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Sample preparation involves the following steps

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To measure sets of metabolites you can use

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NMR is more sensitive than MS

- True
- False

Metabolomics can help to improve healthcare

- True
- False

Metabolite is a low molecular weight organic compound, typically involved in a biological process as a substrate or product

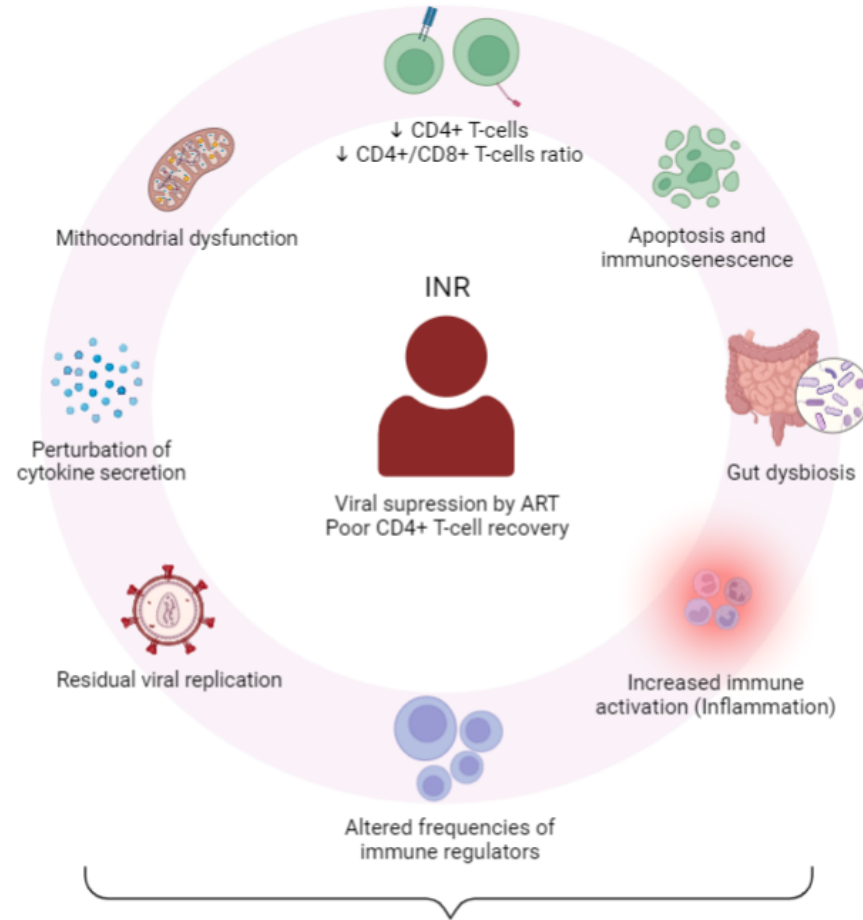
Sample preparation usually includes collection, storage, extraction and preparation

Untargeted measures as many metabolites as possible from a range of biological samples.

MS techniques are incredibly sensitive and accurate and offer a high mass resolution

To develop early-detection systems.

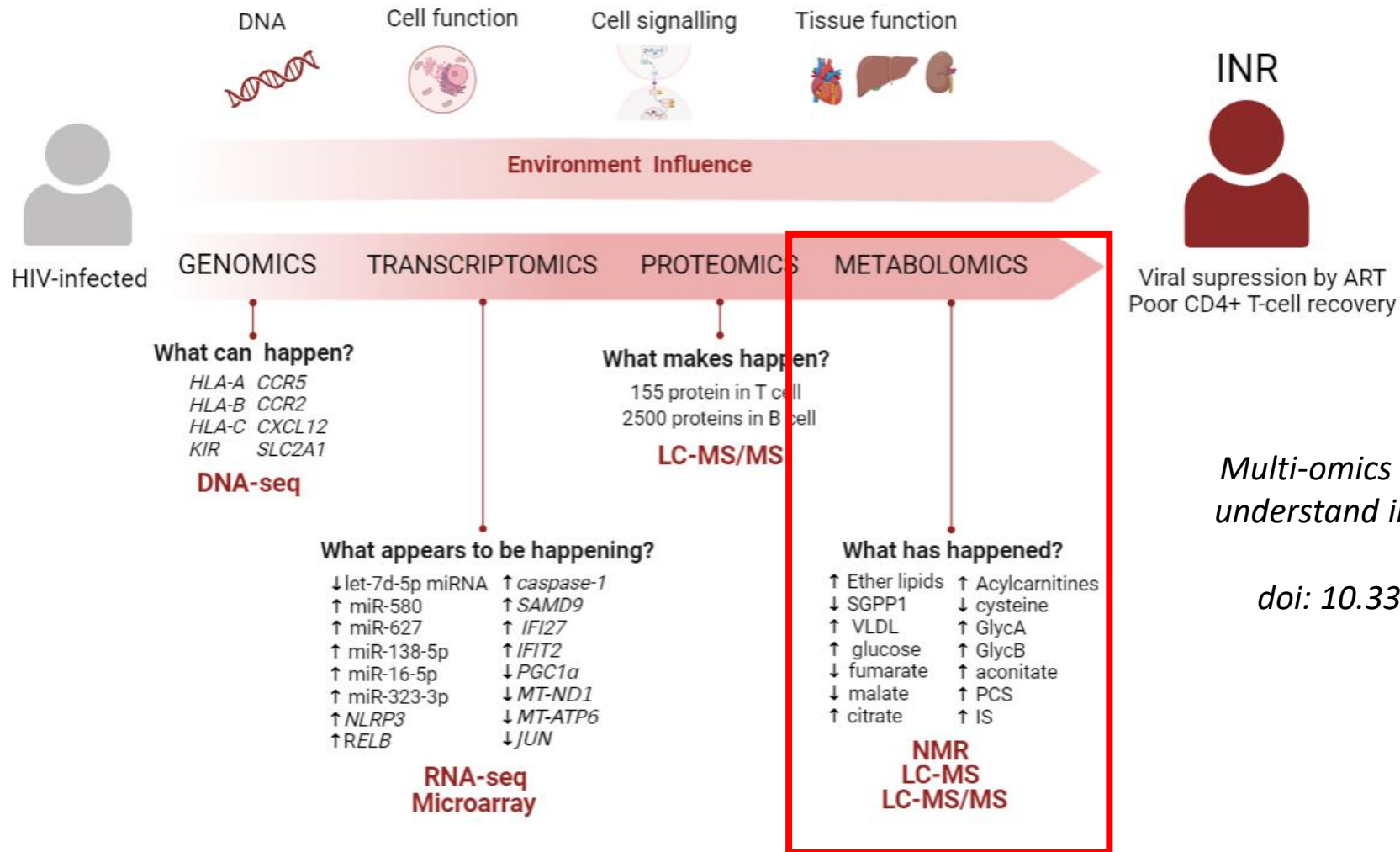
Metabolomics in PLHIV Immunological Non-Responders (INR)



Clinical consequences

- ↑ Morbidity
- ↑ Mortality
- ↑ Inflammation
- Opportunistic infections
- Co-infections
- Cardiovascular diseases
- Cancers

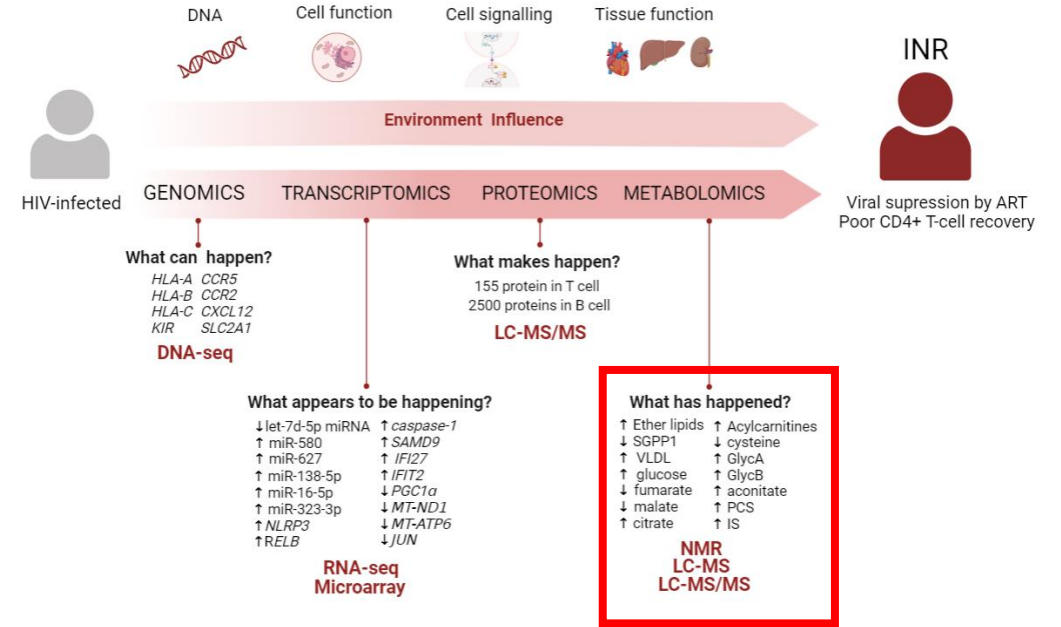
Metabolomics in PLHIV Immunological Non-Responders (INR)



Multi-omics in HIV: searching insights to understand immunological non-response in PLHIV.

doi: 10.3389/fimmu.2023.1228795

METABOLOMICS				
Scarpellini et al.	PLHIV with fail to increase CD4 ⁺ T-cell count by at least 30%	Targeted MS/MS	40 acylcarnitines 19 proteinogenic amino acids, ornithine and citrulline, 19 biogenic amines, the sum of hexoses, 76 phosphatidylcholines, 14 lyso-phosphatidylcholines 15 sphingomyelins	↑ Ether lipids ↓ SGPP1 β-oxidation
Rodriguez-Gallego et al.	pre-ART low nadir (<200 cells/μl) and CD4 ⁺ T-cell count lower than 250 cells/μl at 36 months on ART (baseline study).	NMR	HDL HDL-Cholesterol HDL- TGs VLDL VLDL-Cholesterol VLDL- TGs LDL/HDL	non-HDL lipoprotein particle ↑ VLDL particles ('medium' subclass) ↑ glucose
Masip et al.	pre-ART low nadir (<200 cells/μl) and CD4 ⁺ T-cell count lower than 250 cells/μl at 36 months on ART (longitudinal study).	NMR	HDL HDL-Cholesterol HDL- TGs VLDL VLDL-Cholesterol VLDL- TGs LDL/HDL	↑ large HDL-P ↑ small HDL-P (increased from baseline levels, Rodriguez-Gallego et al)
Qian et al.	CD4 ⁺ T-cell count rise after 2 years of <100 or >300 cells/μl of ART.	PLC-MS/MS-ESI RP/UPLC-MS/MS-ESI+ RP/UPLC-MS/MS-ESI-	125 lipids 68 amino acids 7 peptides 14 carbohydrates 12 cofactors and vitamins 9 nucleotides 6 energy metabolites	↑ Acylcarnitines (MC, PC, OC, and SC) associated with INR
Ferrari et al.	CD4 ⁺ T-cells <350/μL receiving ART for 2 or more years	UPLC-MS/MS	125 metabolites	↑ citrate, aconitate, linolenate ↓ nicotinamide, fumarate, malate and phospholipids ↓ amino acids (isoleucine, alanine, glycine...)
Nyström et al.	pre-ART low nadir (<200 cells/μl) and rise in CD4 ⁺ T-cells <50 cells/year in the first 2 years following suppressive ART.	LC-MS technique	200 metabolites	↓ levels of cysteine could be associated with poor CD4 ⁺ T-cell recovery
Malo et al.	CD4 ⁺ T-cell count rises after 2 years of <100 or >300 cells/μl of ART.	NMR	Plasma glycoprotein profiles	↑ levels of GlycA and GlycB associated with a worse immunological state. ↑ levels of baseline glycoprotein concentrations tend to respond less to ART.



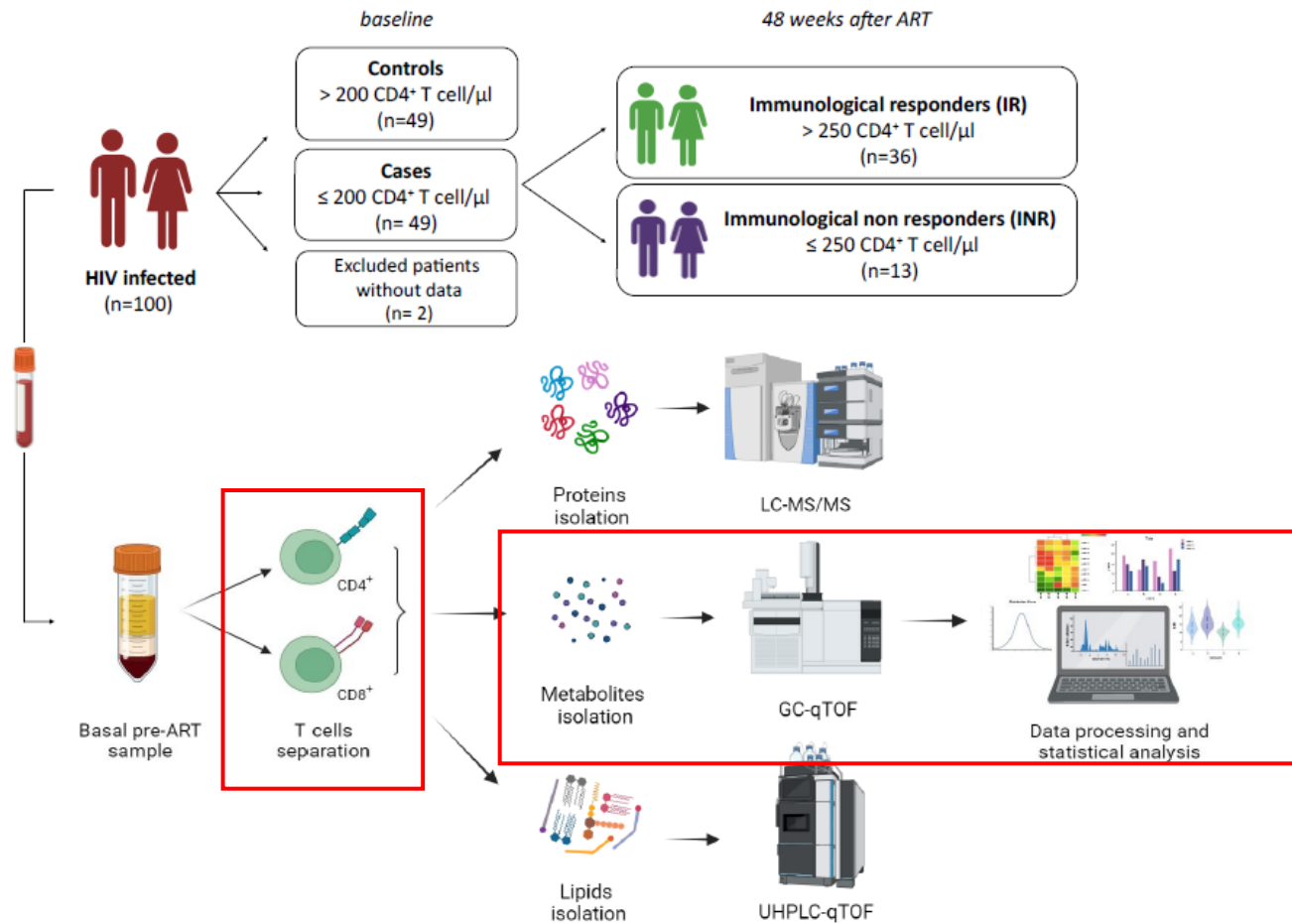
7 articles

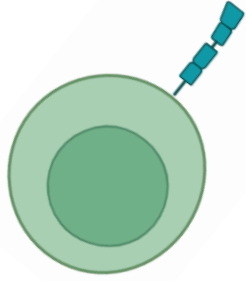
- 4MS
- 3 NMR

- ✓ Plasma/Serum samples
- ✓ Identification of possible biomarkers

Metabolomics in PLHIV Immunological Non-Responders (INR)

Identification of molecular pathways altered in INR condition (Preliminary results)



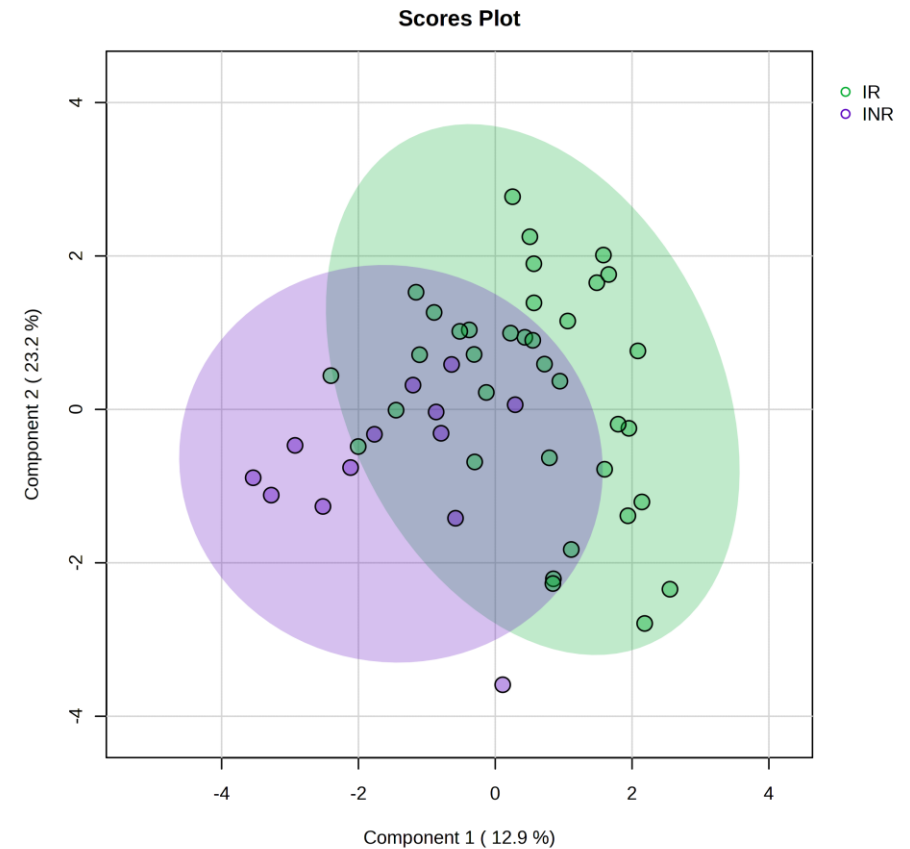
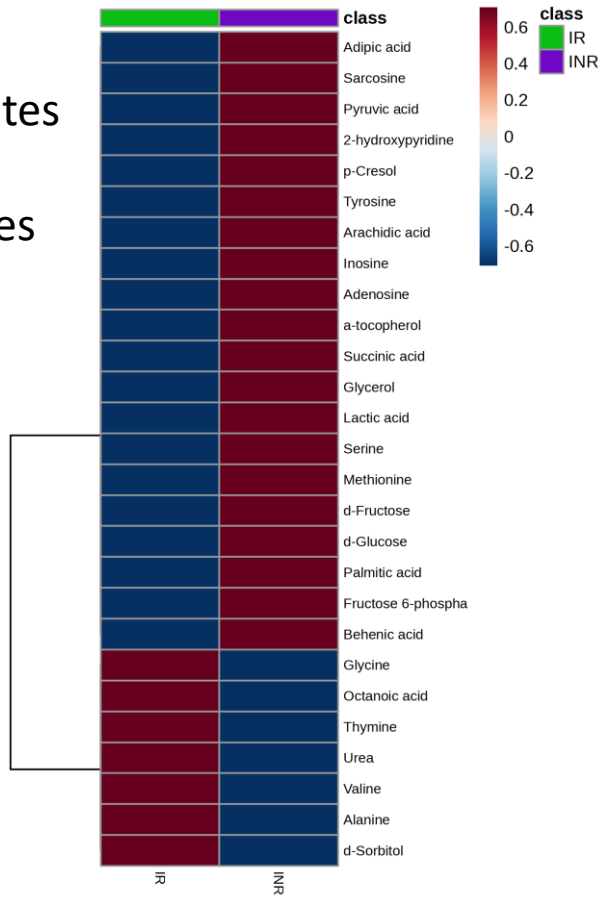


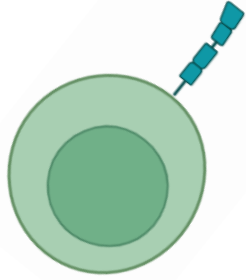
Metabolomics in CD4⁺ T Cells

55 metabolites detected
27 significant metabolites between IR vs INR (Mann-Whitney test)

↑20 metabolites

↓7 metabolites

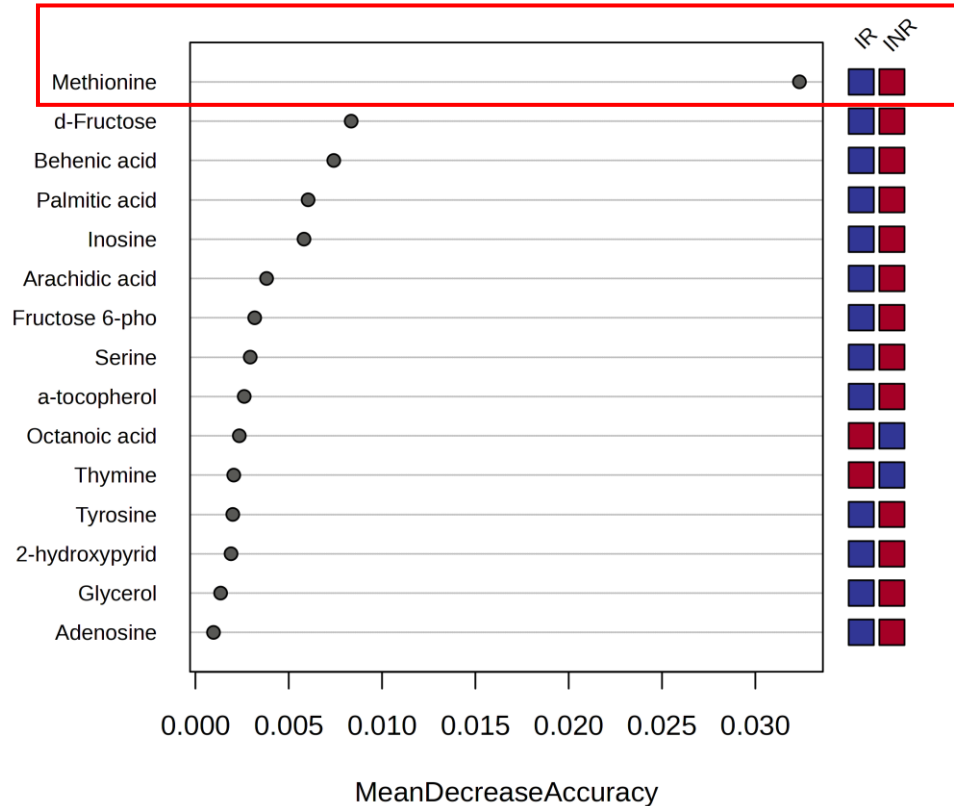




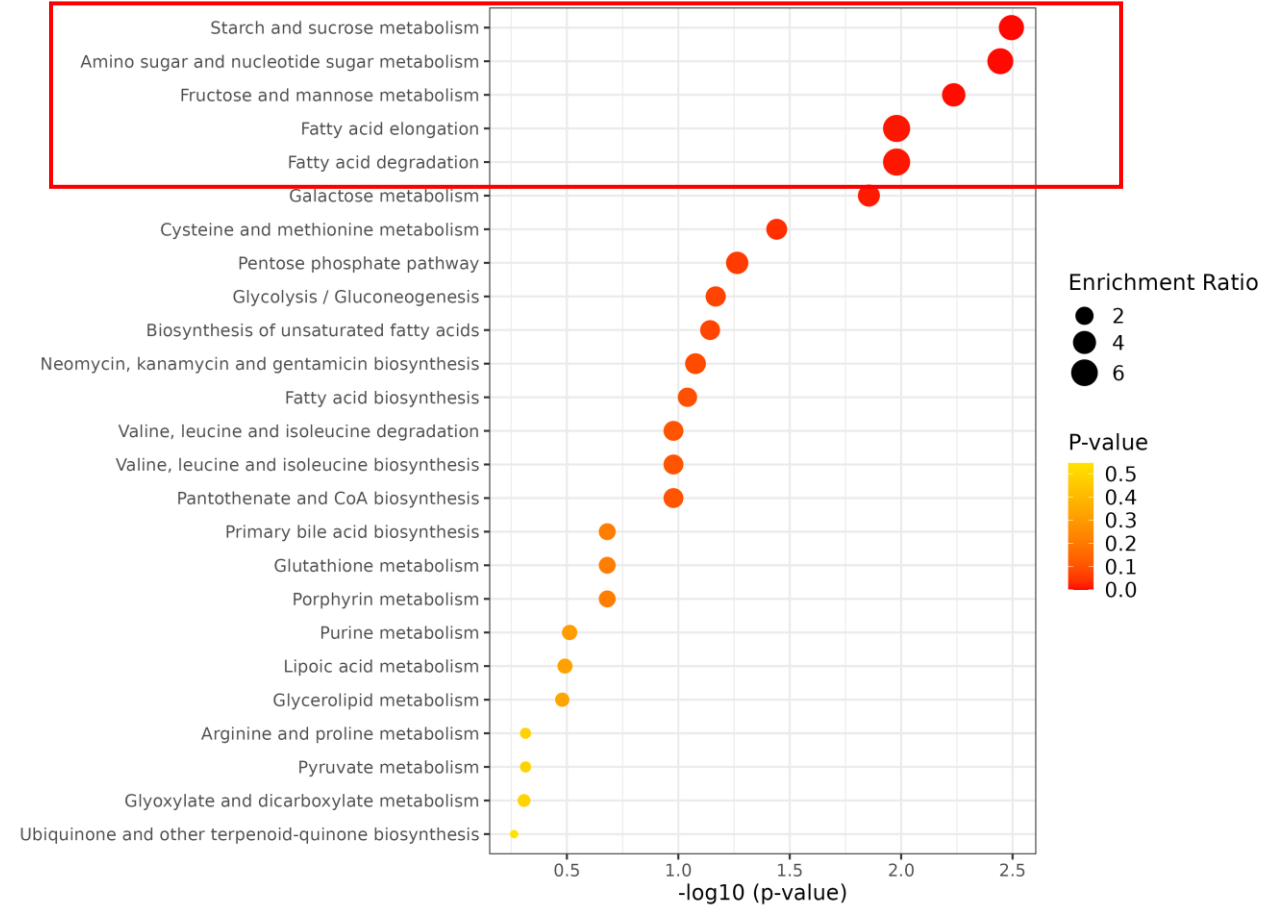
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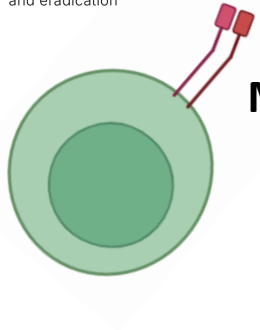
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Overview of Enriched Metabolite Sets (Top 25)



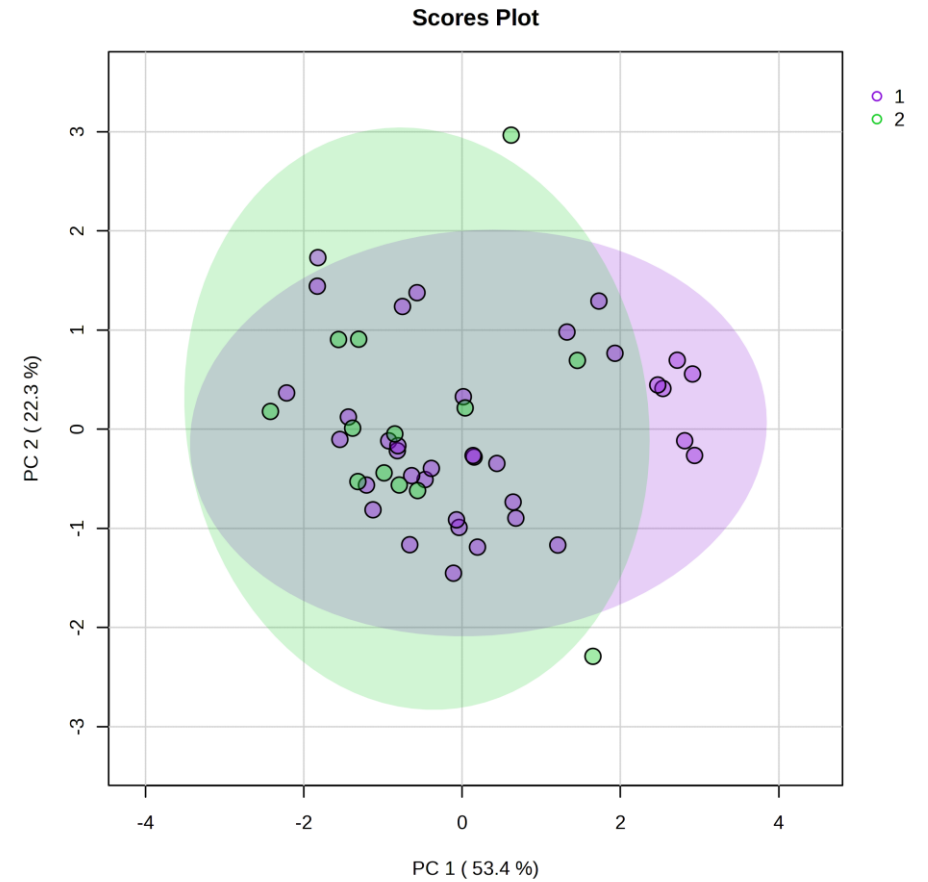
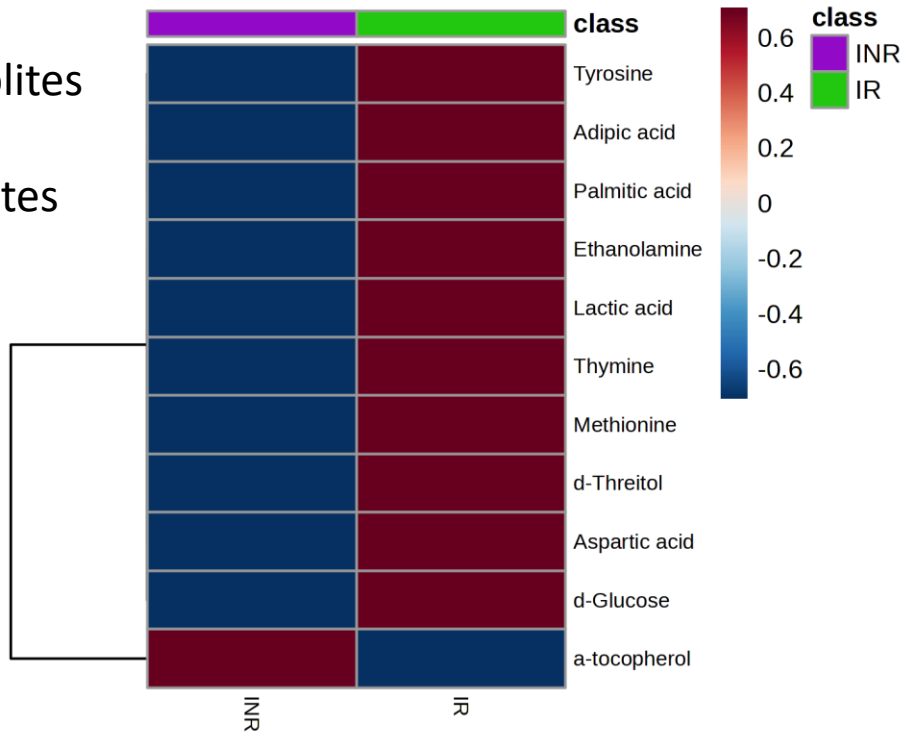


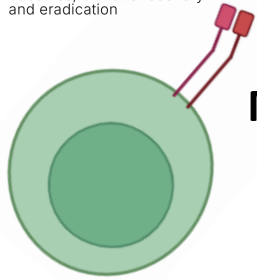
Metabolomics in CD8⁺ T Cells

55 metabolites detected
11 significant metabolites between IR vs INR (Mann-Whitney test)

↑10 metabolites

↓1 metabolites

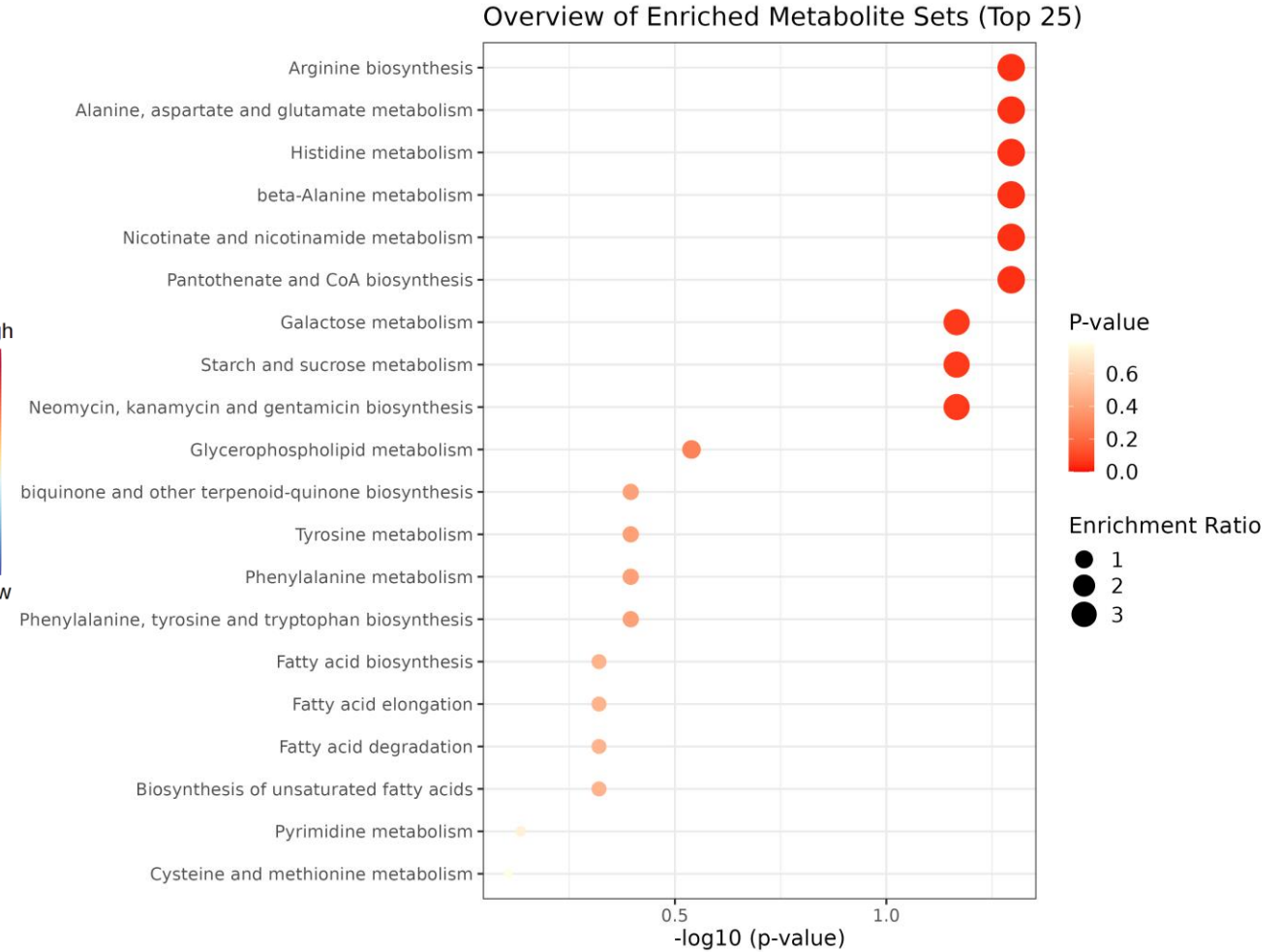
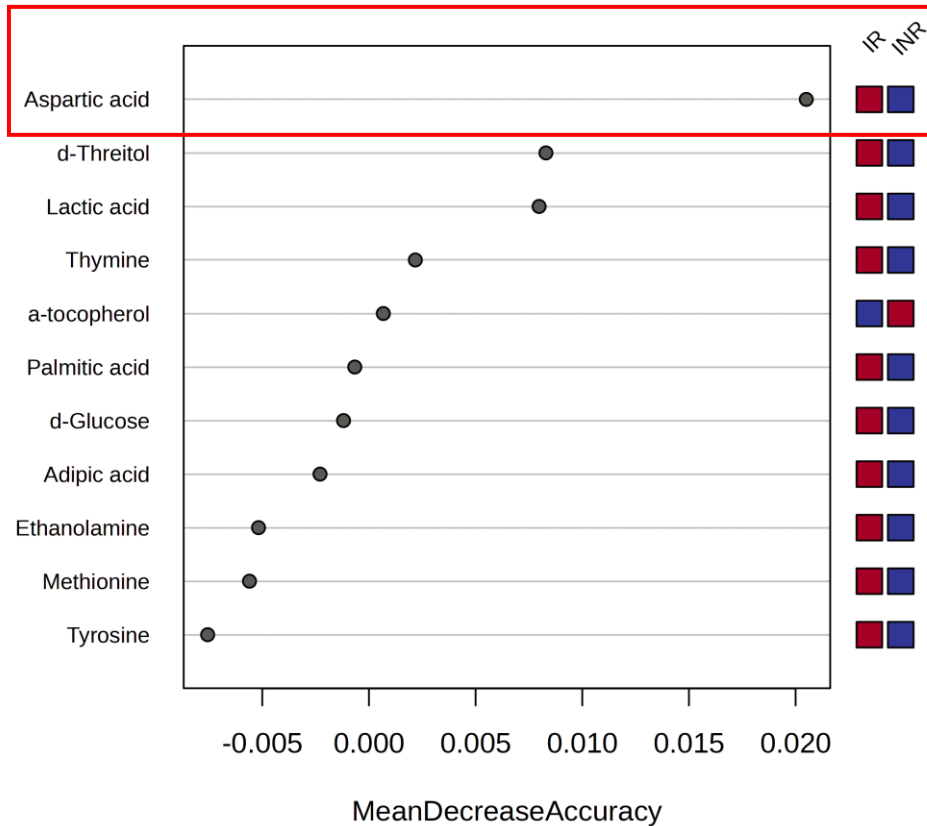




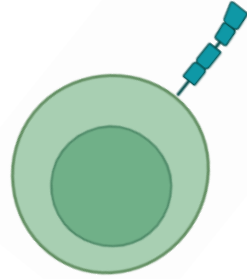
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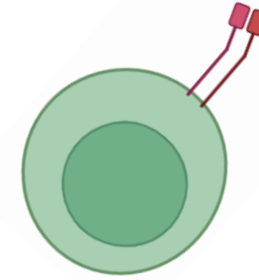


Conclusions



CD4⁺ T Cells

- ✓ Variability in CD4⁺ T cell response
- ✓ Sugar metabolism and fatty acid metabolism alteration



CD8⁺ T Cells

- ✓ Less differences in CD8⁺ T cell metabolic profile (11/55 metabolites)
- ✓ Amino acid metabolism alteration

Impaired mitochondrial activity

Take home message

- ✓ Metabolomics is the comprehensive, qualitative, and quantitative study of low-molecular-weight molecules.
- ✓ Metabolomics can be targeted or untargeted depending on the aim of the study.
- ✓ Applications: Clinical biomarkers, identification of molecular pathways....
- ✓ Metabolomics is a snapshot of the physiology of the cell most closely related to the phenotype.



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Funding



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