



¹³C-FLUXOMICS IN CANCER CELLS

The objective is to acquire theoretical and practical knowledge for the analysis of metabolic systems in cancer cells using ¹³C-fluxomics approaches

TARGET STAFF

- Have an initiated or intermediate level in metabolism and cancer
- Have an ongoing project or submitted around the understanding of metabolism in cancer cells
- Target audience: PhD students, post-docs, researchers, engineers or technical staff in oncology

PROGRAM

Day 1

- General Introduction
- Introduction : Cancer and Metabolism
- Introduction: Metabolic systems

Day 2

- Module 1 « Experimental design and sampling » (Theoretical and practical courses) :
 - Polar molecules
 - Lipides

Day 3

- Module 2 « Analysis and data treatment»
- Module3 « Flux maps »
- Module 4 « Data analysis for metabolic fluxes»

Day 4

- Module 4 : Metabolic flux modelization: cells scale»
- Feedback & round table
- Conclusion and evaluation of the training

Responsible :

Maud Heuillet

Research Engineer INSA GSO

Mass spectrometry and isotopic analysis

Speakers :

Lindsay Peyriga

Engineer assistant INRA

Co-manager of MetaToul- Metabolics Network platform

Jean-Charles Portais

Professor UPS in Biochemistry and metabolism

Scientific director of MetaToul platform

Jean-Emmanuel Sarry

Researcher INSERM

Manager of FlexAML team in CRCT

Justine Bertrand-Michel

Research Engineer INSERM

Co-director of MetaToul platform and responsible of MetaToul-Lipidomics platform

Nathalie Poupin

Researcher INRA

Network analysis and bioinformatics

Floriant Bellvert

Engineer CNRS

Cco-manager of MetaToul- Metabolics Network platform

Edern Cahoreau

Research Engineer CNRS

NMR, isotopic analysis and fluxomics

Fabien Jourdan

Research director INRA

Network analysis and bioinformatics

INEOS

from 08 to 11 October
2019

Duration :

4 days – 30 hours

Location: INSA Toulouse

Price: 800 €

Academic GSO price : 620€

Information & Registration :

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