

Schedule for PhD course “Introduction to nutritional metabolomics”

6 to 10 July 2020

Date/Time	Subject	Lecturer	Lecture room
6 July Monday <i>Theme: Introduction to basic concepts in LC-MS metabolomics research</i>			
9:00--9:30	Outline of the course and introduction of participants	<i>LOD</i>	A001 + A009
9:30--10:05	Perspectives and goals of metabolomics <i>Break</i>	<i>HMR</i>	
10:15--10:50	Metabolomics in Nutrition <i>Break</i>	<i>HMR</i>	
11:00--12:00	Overview of the metabolomics pipeline From sample to data interpretation	<i>GLB</i>	
<i>Lunch will be provided</i>			
13:00--13:30	Experimental design <i>Break</i>	<i>LOD</i>	
13:40--14:00	Navigating freely-available software tools for metabolomics analysis <i>Break</i>	<i>JS</i>	
14:10--16:10	LC-MS based metabolomics Subject 1 : Samples and sample preparation in metabolomics (<i>GLB</i>) Subject 2 : Liquid chromatography and mass spectrometry principles (<i>CDG</i>) Subject 3: LC-MS technologies for targeted and untargeted metabolomics (<i>CDG</i>)	<i>CDG, GLB</i>	
16:20--17:00	LC-MS lab tour	<i>CDG, GLB</i>	
17:30--19:30	<i>Optional: Software installation troubleshooting, group-forming and tutorial</i>	<i>JS, CP, MX</i>	
Date/Time	Subject	Lecturer	Lecture room
7 July Tuesday <i>Theme: Metabolomics data preprocessing, normalization and annotation</i>			
9:00--9:30	Data preprocessing in metabolomics: Basic concepts in preprocessing (e.g. feature detection, alignment, gap filling)	<i>JS</i>	
9:30--9:40	Conversion of raw data	<i>JS</i>	
9:40--10:00	A brief introduction to R: Basic syntax, working in Rstudio <i>Break</i>	<i>JS</i>	
10:10--10:40	A brief introduction to R: We import data together, modify it, and export	<i>JS, CP, MX</i>	
10:40--11:30	XCMS walk-through and CAMERA annotation	<i>JS</i>	

11:30--12:00	Presentation of exercise LC-MS dataset: Extraction of food intake markers <i>Exercise: Hands on LC-MS data preprocessing (XCMS)</i>	JS, CP, MX, HMR	A001 + A009
	Lunch will be provided		
13:00--13:50	<i>Exercise: Hands on LC-MS data preprocessing (XCMS)</i> Break	JS, CP, MX, HMR	
14:00--15:00	Introduction to Normalization and transformations Break	CP	
15:10-17:00	<i>Exercise: Hands on data normalization and annotation</i>	JS, CP, MX	
18:00--21:30	Dinner party		TBD
8 July Wednesday Theme: Metabolomics data analysis			
8:30--9:10	Data Analysis I: Univariate Methods Break	JS	
9:20--10:05	<i>Exercise: Hands on univariate analysis using R</i> Break	JS, CP, MX	
10:15--11:15	Data Analysis II: Multivariate Approach - PCA Break	CP	
11:20--12:00	<i>Exercise: Hands on PCA using R</i>	JS, CP, MX, HMR	A001 + A009
	Lunch will be provided		
13:00--13:20	<i>Exercise: Hands on PCA using R</i> Break	JS, CP, MX, HMR	
13:30--14:30	Data Analysis III: Multivariate Approach - PLSDA Break	CP	
14:40-17:00	<i>Exercise: Hands on PLSDA using R</i>	JS, CP, MX, HMR	
17:30-19:30	<i>Optional Tutorial</i>	JS, CP, MX	
Date/Time	Subject	Lecturer	Lecture room
9 July Thursday Theme: Metabolite identification			
9:00--9:50	Metabolite Identification (Molecular formula, Interpretation of MS and MS/MS spectra, compound databases)	GLB	

10:00--12:00	Break <i>Exercise: Fragmentation, adducts, annotation</i>	GLB, JS, CDG	A001 + A009
	Lunch will be provided		
13:00--14:00	Compound and spectral databases, in-silico prediction, levels of identification	GLB	
14:10--15:20	Break <i>Exercise: Fragmentation, adducts, annotation</i>	GLB, JS, CDG	
15:30--17:00	Break <i>Exercise: Hands on - Identification of markers of food intake</i>	GLB, JS, CDG	
17:30--19:30	<i>Optional Tutorial</i>	GLB, JS, CDG	
10 July Friday Theme: Project work and presentation of results			
8:30--11:30	Students presentations	LOD, JS, CDG, HMR, GLB,	A001 + A009
11:30--12:00	Course Evaluation	LOD, JS	

Lecturers

LOD : Lars Ove Dragsted

JS: Jan Stanstrup

CDG : Cristian De Gobba

HMR: Henrik Munch Roager

GLB : Giorgia La Barbera

CP : Ceyda Pekmez

MX : Muyao Xi