

Lecture 11. Lipidomics.

Learning outcomes:

1. Give the definition to the terms “lipids”, “lypoproteins”, “lipidome”, “lipidomics”.
2. Analyze the different types of lipids by their chemical structure and function, give the specific examples.
3. Explain the methods of lipidomic research.
4. Explain different disturbances of lipid metabolism and methods of their diagnostics and treatment, give the specific examples.

Lipidomics is the large-scale study of pathways and networks of cellular lipids in biological systems. The word "**lipidome**" is used to describe the complete lipid profile within a cell, tissue, organism, or ecosystem and is a subset of the "**metabolome**" which also includes the three other major classes of biological molecules: proteins/amino-acids, sugars and nucleic acids. Lipidomics is a relatively recent research field that has been driven by rapid advances in technologies such as **mass spectrometry (MS)**, **nuclear magnetic resonance (NMR) spectroscopy**, **fluorescence spectroscopy**, **dual polarisation interferometry** and **computational methods**, coupled with the recognition of the role of lipids in many metabolic diseases such as obesity, atherosclerosis, stroke, hypertension and diabetes. This rapidly expanding field complements the huge progress made in genomics and proteomics, all of which constitute the family of systems biology.

Lipids are a diverse and ubiquitous group of compounds which have many key biological functions, such as acting as structural components of cell membranes, serving as energy storage sources and participating in signaling pathways. Lipids may be broadly defined as **hydrophobic** or **amphipathic** small molecules that originate entirely or in part from two distinct types of biochemical subunits or "building blocks": ketoacyl and isoprene groups. The huge structural diversity found in lipids arises from the biosynthesis of various combinations of these building blocks. For example, **glycerophospholipids** are composed of a glycerol backbone linked to one of approximately 10 possible headgroups and also to 2 fatty acyl/alkyl chains, which in turn may have 30 or more different molecular structures. In practice, not all possible permutations are detected experimentally, due to chain preferences depending on the cell type and also to detection limits - nevertheless several hundred distinct glycerophospholipid molecular species have been detected in mammalian cells. **Plant chloroplast thylakoid membranes** however, have unique lipid composition as they are deficient in phospholipids. Also, their largest constituent, **monogalactosyl diglyceride or MGDG**, does not form aqueous bilayers. Nevertheless, dynamic studies reveal a normal lipid bilayer organisation in thylakoid membranes.

Most methods of lipid **extraction** and **isolation** from biological samples exploit the **high solubility of hydrocarbon chains in organic solvents**. Given the diversity in lipid classes, it is not possible to accommodate all classes with a common extraction method. The traditional Bligh/Dyer procedure uses **chloroform/methanol**-based protocols that include phase partitioning into the organic layer. These protocols work relatively well for a wide variety of physiologically relevant lipids but they have to be adapted for complex lipid chemistries and low-abundance and labile lipid metabolites. When organic soil was used, **citrate buffer** in the extraction mixture gave higher amounts of lipid phosphate than **acetate buffer, Tris, H₂O or phosphate buffer**. The simplest method of **lipid separation** is the use of **thin layer chromatography (TLC)**. Although not as sensitive as other methods of lipid detection, it offers a rapid and comprehensive screening tool prior to more sensitive and sophisticated techniques. **Solid-phase extraction (SPE) chromatography** is useful for rapid, preparative separation of crude lipid mixtures into different lipid classes. This involves the use of prepacked columns containing silica or other stationary phases to separate glycerophospholipids, fatty acids, cholesteryl esters, glycerolipids, and sterols from crude lipid mixtures. **High-performance**

liquid chromatography (HPLC or LC) is extensively used in lipidomic analysis to separate lipids prior to mass analysis. Separation can be achieved by either **normal-phase (NP) HPLC or reverse-phase (RP) HPLC**. For example, NP-HPLC effectively separates glycerophospholipids on the basis of headgroup polarity, whereas RP-HPLC effectively separates fatty acids such as eicosanoids on the basis of chain length, degree of unsaturation and substitution. For global, untargeted lipidomic studies it is common to use both RP and NP or **Hydrophilic Interaction Liquid Chromatography (HILC)** columns for increased lipidome coverage. The application of **nano-flow liquid chromatography (nLC)** proved thereby to be most efficient to enhance both general measurement sensitivity and lipidome coverage for a global lipidomics approach. **Chromatographic (HPLC/UHPLC) separation** of lipids may either be performed offline or online where the eluate is integrated with the ionization source of a **mass spectrometer**. The progress of modern lipidomics has been greatly accelerated by the development of **spectrometric methods** in general and soft ionization techniques for mass spectrometry such as **electrospray ionization (ESI)**, **desorption electrospray ionization (DESI)**, and **matrix-assisted laser desorption/ionization (MALDI)** in particular. "Soft" ionization does not cause extensive fragmentation, so that comprehensive detection of an entire range of lipids within a complex mixture can be correlated to experimental conditions or disease state. In addition, the technique of **atmospheric pressure chemical ionization (APCI)** has become increasingly popular for the analysis of nonpolar lipids.

The questions for self - control:

1. What are the "lipids", "lypoproteins", "lipidome", "lipidomics"?
2. Chemical structure and functions of the different types of lipids.
3. Methods of lipidomic research.
4. Diseases connected with the disturbances of lipid metabolism and methods of their diagnostics and treatment.

Recommended readings:

1. Wenk MR (July 2005). "The emerging field of lipidomics". *Nat Rev Drug Discov.* 4 (7): 594–610. doi:10.1038/nrd1776. PMID 16052242. S2CID 83931214.
2. Watson AD (October 2006). "Thematic review series: systems biology approaches to metabolic and cardiovascular disorders. Lipidomics: a global approach to lipid analysis in biological systems". *J. Lipid Res.* 47 (10): 2101–11. doi:10.1194/jlr.R600022-JLR200. PMID 16902246.
3. "Lipidomics". *The Lipid Chronicles*. 2011-12-15. Retrieved 2012-01-08.
4. Han X (2007). "Neurolipidomics: challenges and developments". *Front. Biosci.* 12: 2601–15. doi:10.2741/2258. PMC 2141543. PMID 17127266.
5. Fahy E, Subramaniam S, Brown HA, et al. (2005). "A comprehensive classification system for lipids". *J. Lipid Res.* 46 (5): 839–61. doi:10.1194/jlr.E400004-JLR200. PMID 15722563.
6. YashRoy R.C. (1990) Magnetic resonance studies on dynamic organisation of lipids in chloroplast membranes. *Journal of Biosciences*, vol. 15(4), pp. 281-288. https://www.researchgate.net/publication/225688482_Magnetic_resonance_studies_of_dynamic_organisation_of_lipids_in_chloroplast_membranes?ev=prf_pub
7. Bligh EG, Dyer WJ; Dyer (August 1959). "A rapid method of total lipid extraction and purification". *Can J Biochem Physiol.* 37 (8): 911–7. doi:10.1139/o59-099. PMID 13671378. S2CID 7311923.
8. Krank J, Murphy RC, Barkley RM, Duchoslav E, McAnoy A; Murphy; Barkley; Duchoslav; McAnoy (2007). Qualitative analysis and quantitative assessment of changes in neutral glycerol lipid molecular species within cells. *Meth. Enzymol. Methods in Enzymology.* 432. pp. 1–20. doi:10.1016/S0076-6879(07)32001-6. ISBN 978-0-12-373895-0. PMID 17954211.

9. Ivanova PT, Milne SB, Byrne MO, Xiang Y, Brown HA; Milne; Byrne; Xiang; Brown (2007). Glycerophospholipid identification and quantitation by electrospray ionization mass spectrometry. *Meth. Enzymol. Methods in Enzymology*. 432. pp. 21–57. doi:10.1016/S0076-6879(07)32002-8. ISBN 978-0-12-373895-0. PMID 17954212.
10. Deems R, Buczynski MW, Bowers-Gentry R, Harkewicz R, Dennis EA; Buczynski; Bowers-Gentry; Harkewicz; Dennis (2007). Detection and quantitation of eicosanoids via high performance liquid chromatography-electrospray ionization-mass spectrometry. *Meth. Enzymol. Methods in Enzymology*. 432. pp. 59–82. doi:10.1016/S0076-6879(07)32003-X. ISBN 978-0-12-373895-0. PMID 17954213.
11. McDonald JG, Thompson BM, McCrum EC, Russell DW; Thompson; McCrum; Russell (2007). Extraction and analysis of sterols in biological matrices by high performance liquid chromatography electrospray ionization mass spectrometry. *Meth. Enzymol. Methods in Enzymology*. 432. pp. 145–70. doi:10.1016/S0076-6879(07)32006-5. ISBN 978-0-12-373895-0. PMID 17954216.
12. Garrett TA, Guan Z, Raetz CR; Guan; Raetz (2007). Analysis of ubiquinones, dolichols, and dolichol diphosphate-oligosaccharides by liquid chromatography-electrospray ionization-mass spectrometry. *Meth. Enzymol. Methods in Enzymology*. 432. pp. 117–43. doi:10.1016/S0076-6879(07)32005-3. ISBN 978-0-12-373895-0. PMID 17954215.
13. Sullards MC, Allegood JC, Kelly S, Wang E, Haynes CA, Park H, Chen Y, Merrill AH; Allegood; Kelly; Wang; Haynes; Park; Chen; Merrill Jr (2007). Structure-specific, quantitative methods for analysis of sphingolipids by liquid chromatography-tandem mass spectrometry: "inside-out" sphingolipidomics. *Meth. Enzymol. Methods in Enzymology*. 432. pp. 83–115. doi:10.1016/S0076-6879(07)32004-1. ISBN 978-0-12-373895-0. PMID 17954214.
14. Å. Frostegård, A. Tunlid & E. Bååth (August 1991). "Microbial biomass measured as total lipid phosphate in soils of different organic content". *Journal of Microbiological Methods*. 14 (3): 151–163. doi:10.1016/0167-7012(91)90018-L.
15. Kaluzny MA, Duncan LA, Merritt MV, Epps DE; Duncan; Merritt; Epps (January 1985). "Rapid separation of lipid classes in high yield and purity using bonded phase columns". *J. Lipid Res*. 26 (1): 135–40. PMID 3973509.
16. Malavolta M, Bocci F, Boselli E, Frega NG; Bocci; Boselli; Frega (October 2004). "Normal phase liquid chromatography-electrospray ionization tandem mass spectrometry analysis of phospholipid molecular species in blood mononuclear cells: application to cystic fibrosis". *J. Chromatogr. B*. 810 (2): 173–86. doi:10.1016/j.jchromb.2004.07.001. PMID 15380713.
17. Nakamura T, Bratton DL, Murphy RC; Bratton; Murphy (August 1997). "Analysis of epoxyeicosatrienoic and monohydroxyeicosatetraenoic acids esterified to phospholipids in human red blood cells by electrospray tandem mass spectrometry". *J Mass Spectrom*. 32 (8): 888–96. Bibcode:1997JMSp...32..888N. doi:10.1002/(SICI)1096-9888(199708)32:8<888::AID-JMS548>3.0.CO;2-W. PMID 9269087.
18. Danne-Rasche, Niklas; Coman, Cristina; Ahrends, Robert (2018). "Nano-LC/NSI MS Refines Lipidomics by Enhancing Lipid Coverage, Measurement Sensitivity, and Linear Dynamic Range". *Analytical Chemistry*. 90 (13): 8093–8101. doi:10.1021/acs.analchem.8b01275. ISSN 1520-6882. PMID 29792796.
19. Z. Takáts; J.M. Wiseman; B. Gologan; R.G. Cooks (2004). "Mass Spectrometry Sampling Under Ambient Conditions with Desorption Electrospray Ionization". *Science*. 306 (5695): 471–473. Bibcode:2004Sci...306..471T. doi:10.1126/science.1104404. PMID 15486296. S2CID 22994482.
20. Fuchs B, Schiller J; Schiller (2008). MALDI-TOF MS analysis of lipids from cells, tissues and body fluids. *Subcell. Biochem. Subcellular Biochemistry*. 49. pp. 541–65. doi:10.1007/978-1-4020-8831-5_21. ISBN 978-1-4020-8830-8. PMID 18751926.

21. Byrdwell WC (April 2001). "Atmospheric pressure chemical ionization mass spectrometry for analysis of lipids". *Lipids*. 36 (4): 327–46. doi:10.1007/s11745-001-0725-5. PMID 11383683. S2CID 4017177.