

POLYUNSATURATED LIPID SPECIES OF HDL ARE MOST STRONGLY AFFECTED BY GENETIC APOLIPOPROTEIN A-I DEFICIENCY

Emile Zakiev^{ac}, Fabiana Rached^b, Marie Lhomme^a, Carlos V. Serrano Jr.^b, Raul D. Santos^b, John Chapman^a, Alexander Orekhov^{cd}, Anatol Kontush^a

a – UMR-ICAN 1166, National Institute for Health and Medical Research (INSERM), Paris, France

b – University of Sao Paulo , Sao Paulo, Brazil

c – Institute of General Pathology and Pathophysiology, Moscow, Russia

d – Institute for Atherosclerosis Research, Moscow, Russia



INSTITUTE FOR
ATHEROSCLEROSIS RESEARCH
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- Authors declare that they have no conflict of interest to disclose

Rationale



- Both circulating levels of **high-density lipoprotein** (HDL) and its functionality are under genetic control
- **Anti-atherogenic** properties of HDL depend on the **molecular composition** of its **lipidome** and proteome
- Kindreds with genetically determined low HDL-cholesterol are characterised by markedly perturbed lipoprotein metabolism

Working hypothesis



- Hereditary familial hypoHDLemia features distinct alterations in molecular species of the HDL lipidome

Experimental strategy

recruitment



Subjects: Nonsense mutation at
codon -2, Q[-2]X in APOA1

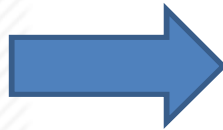
♂=5 ♀= 1

Controls: Healthy normolipidemic
controls

♂=9 ♀=0

Experimental strategy

recruitment



Blood sampling

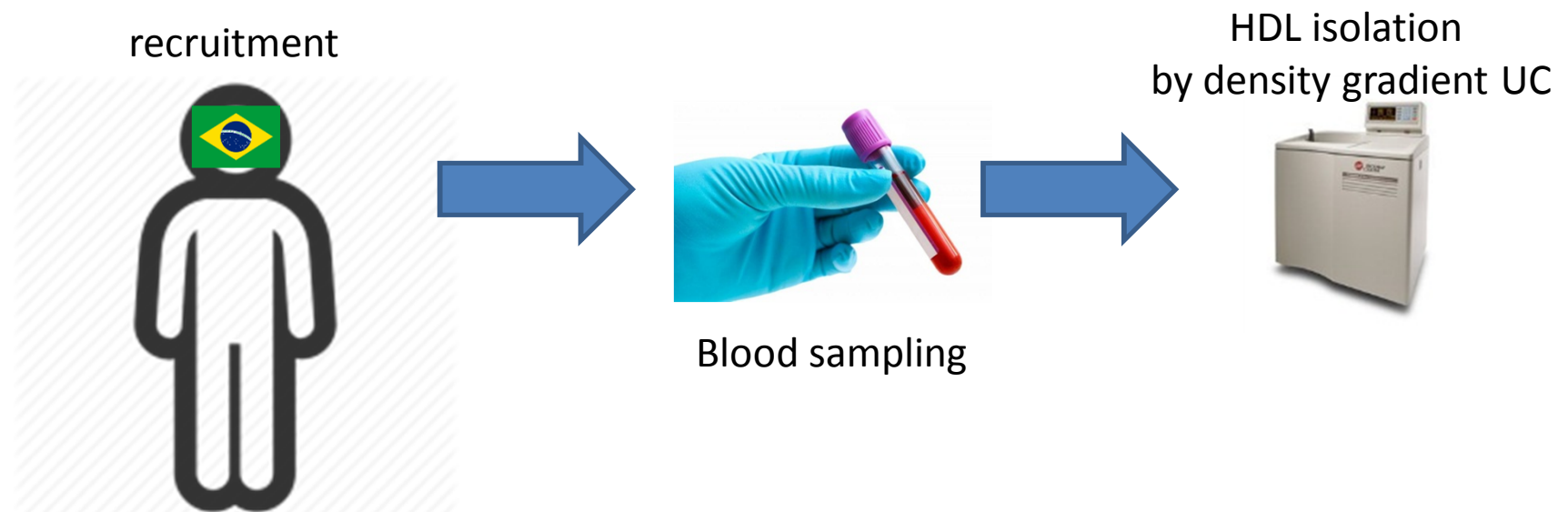
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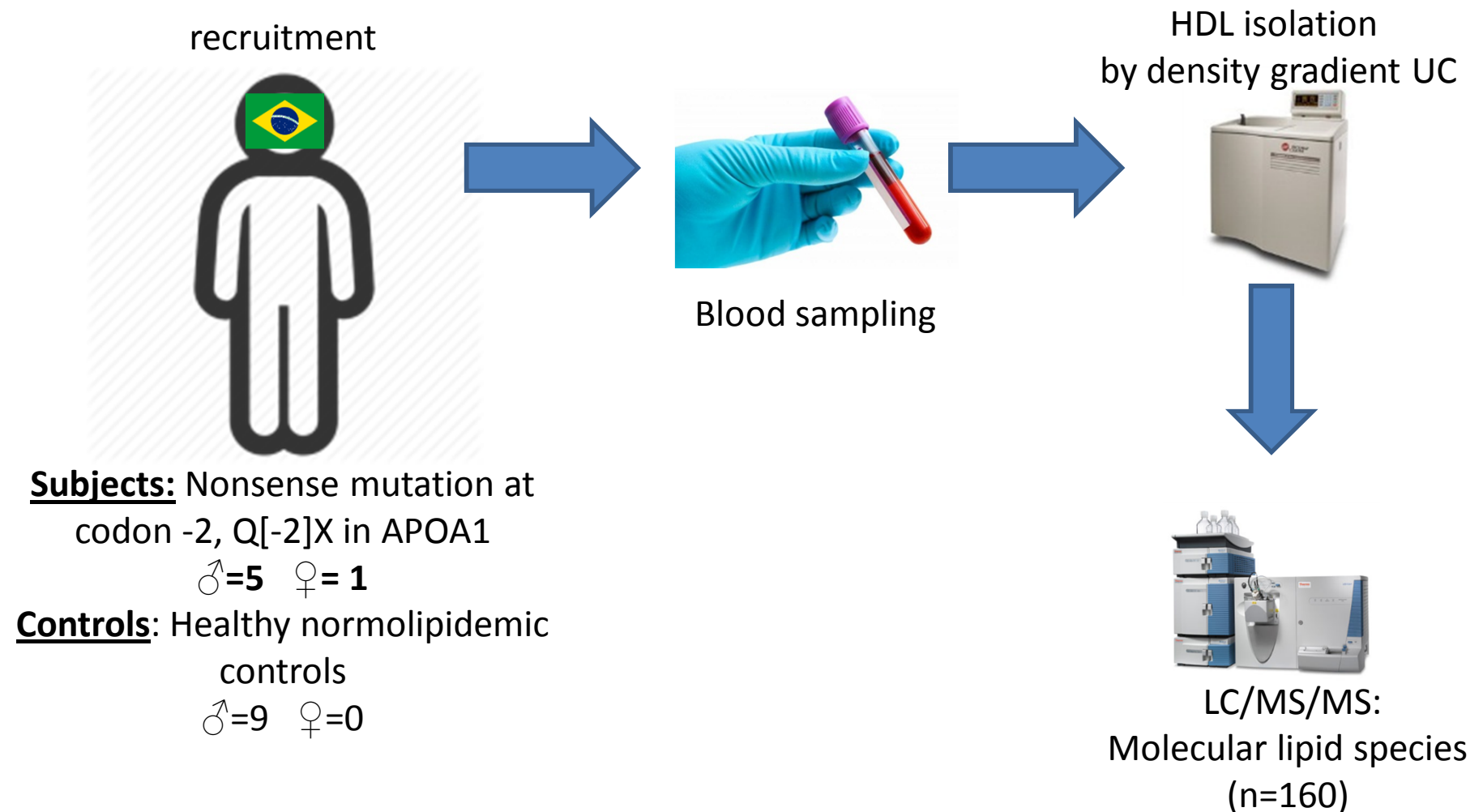
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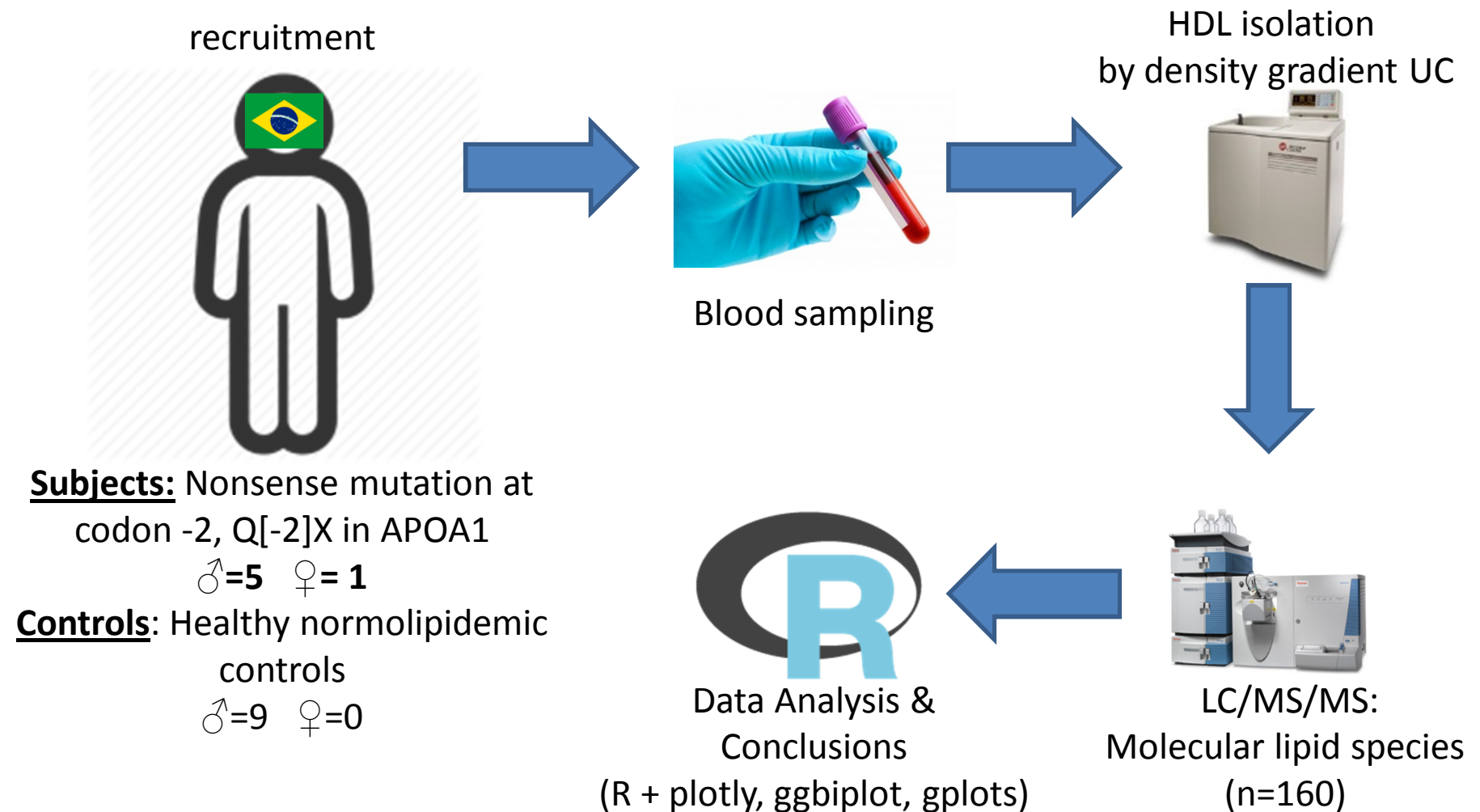
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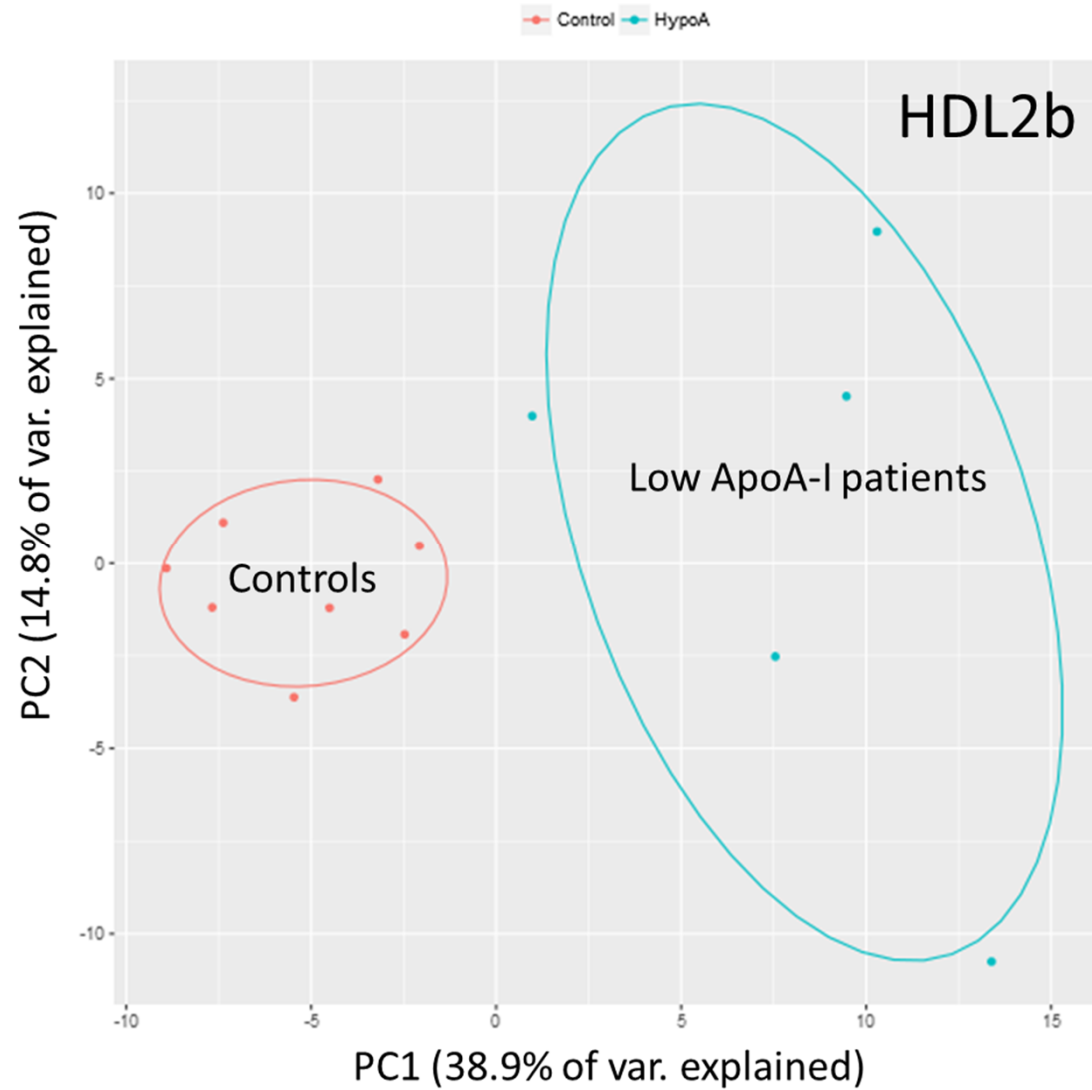
Experimental strategy



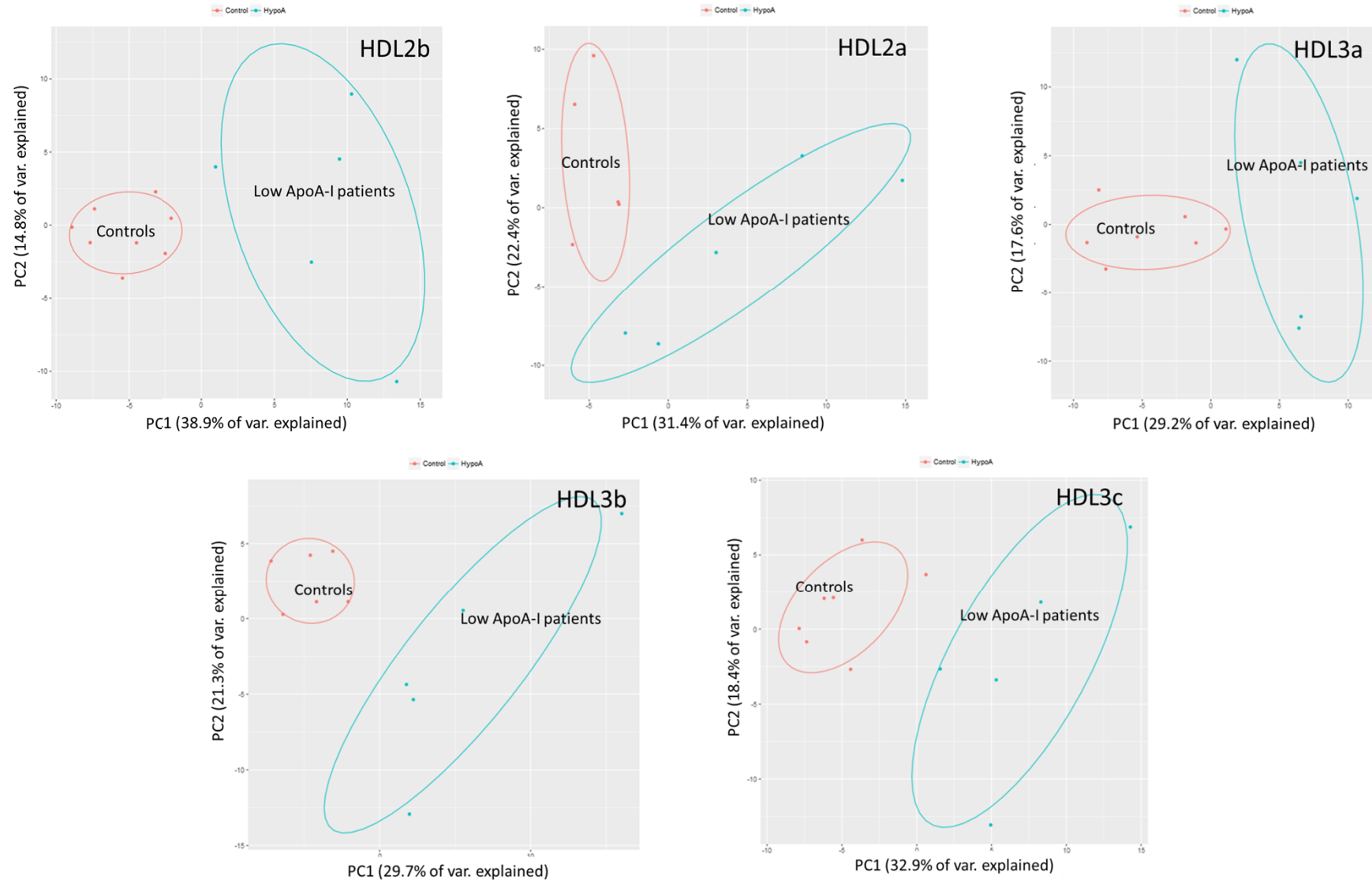
Experimental strategy

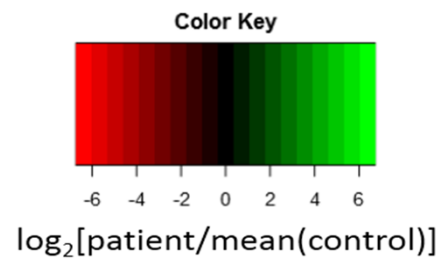


Principal Component Analysis: Distinct lipidome of all HDL subpopulations in apoA-I deficiency

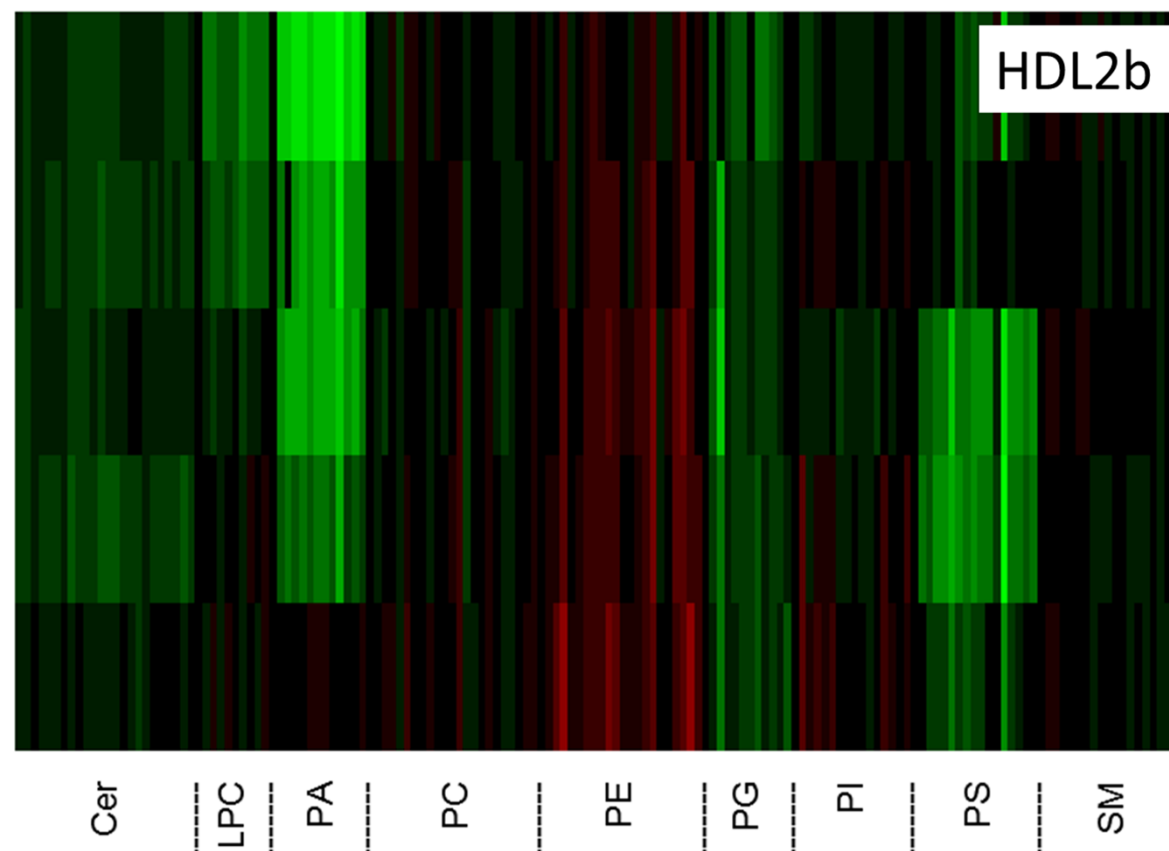


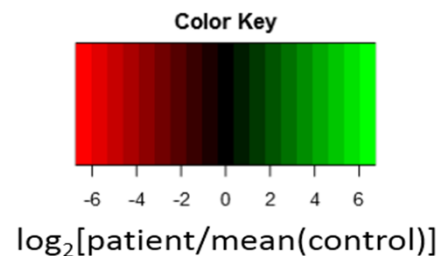
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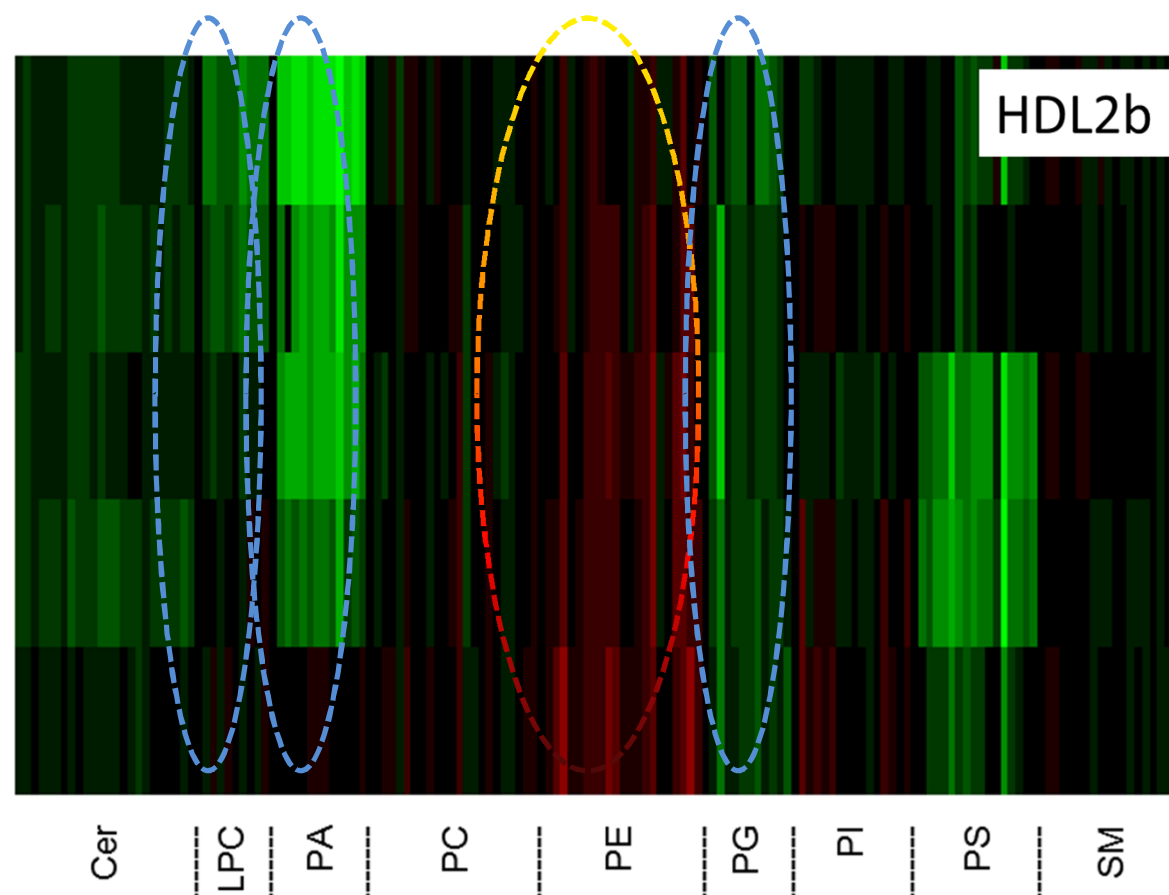


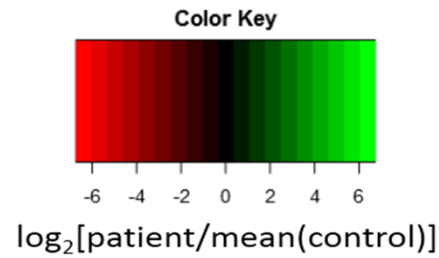
Heatmaps: Systematic alterations of molecular species of LPC, PA, PG and PE across HDL subpopulations in apoA-I deficiency



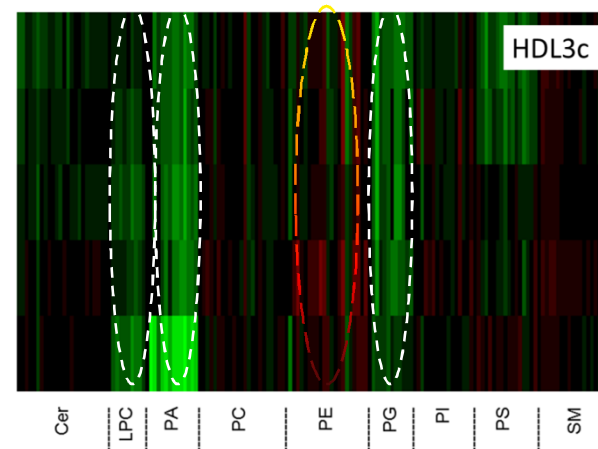
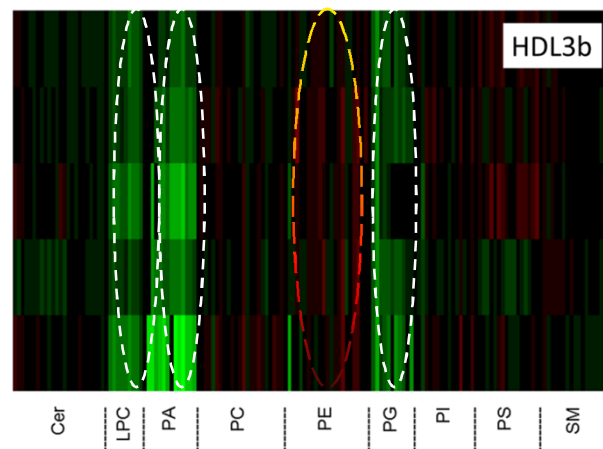
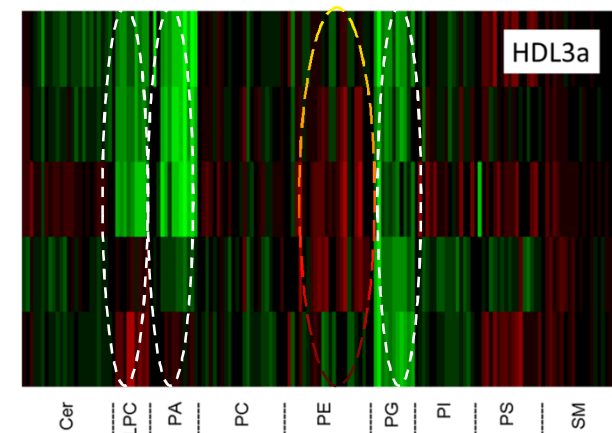
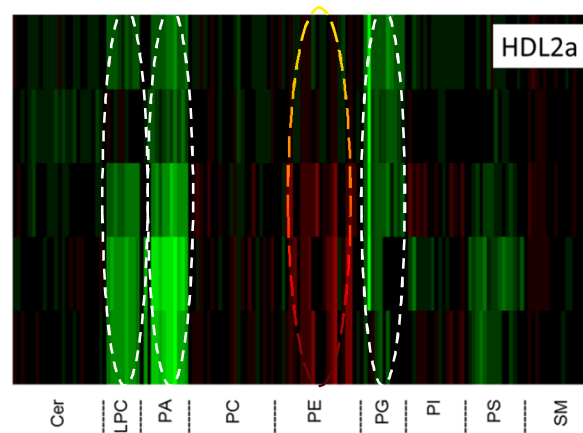
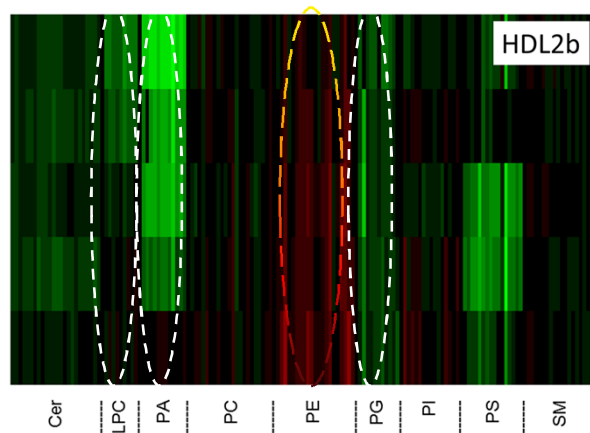


Heatmaps: Systematic alterations of molecular species of LPC, PA, PG and PE across HDL subpopulations in apoA-I deficiency

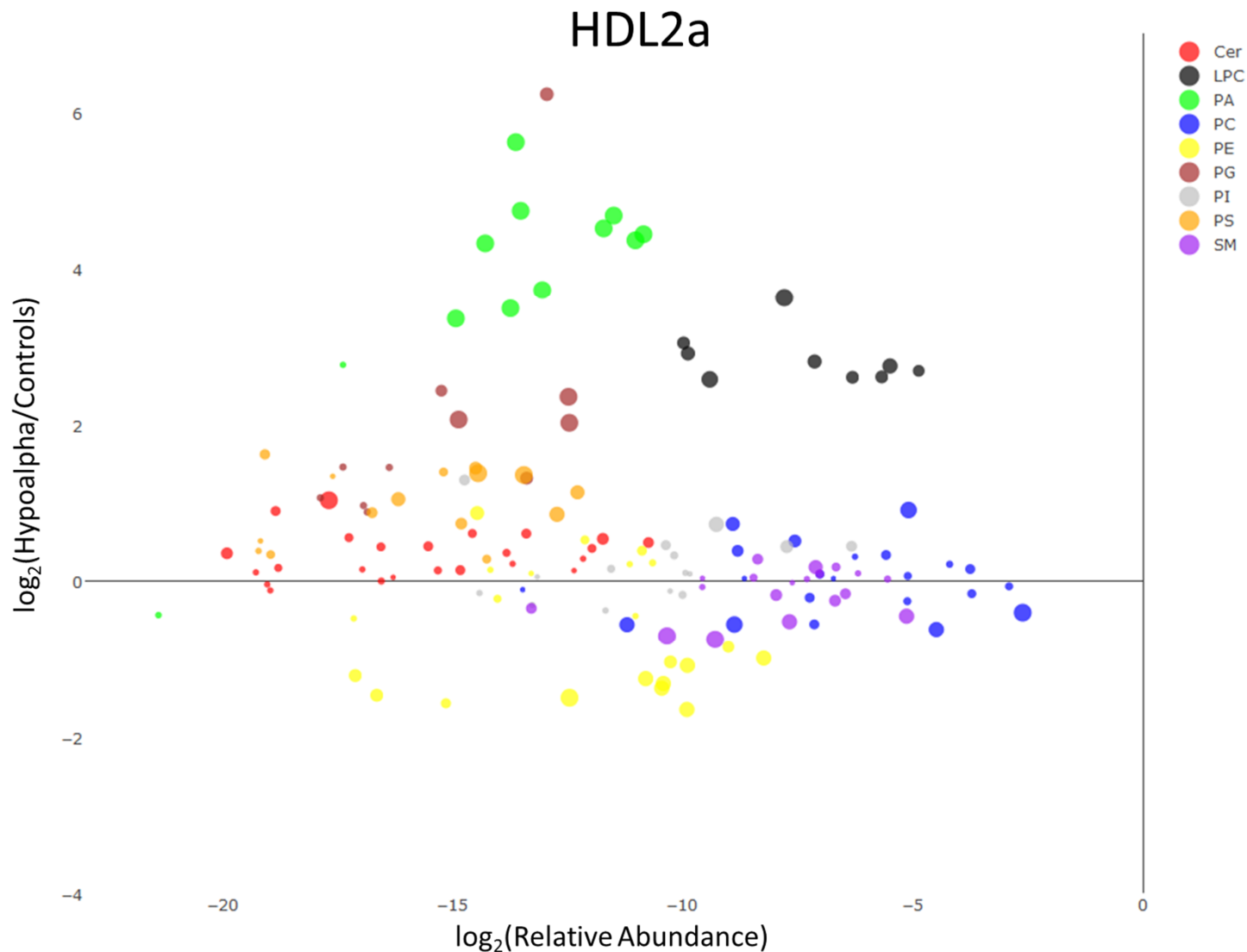




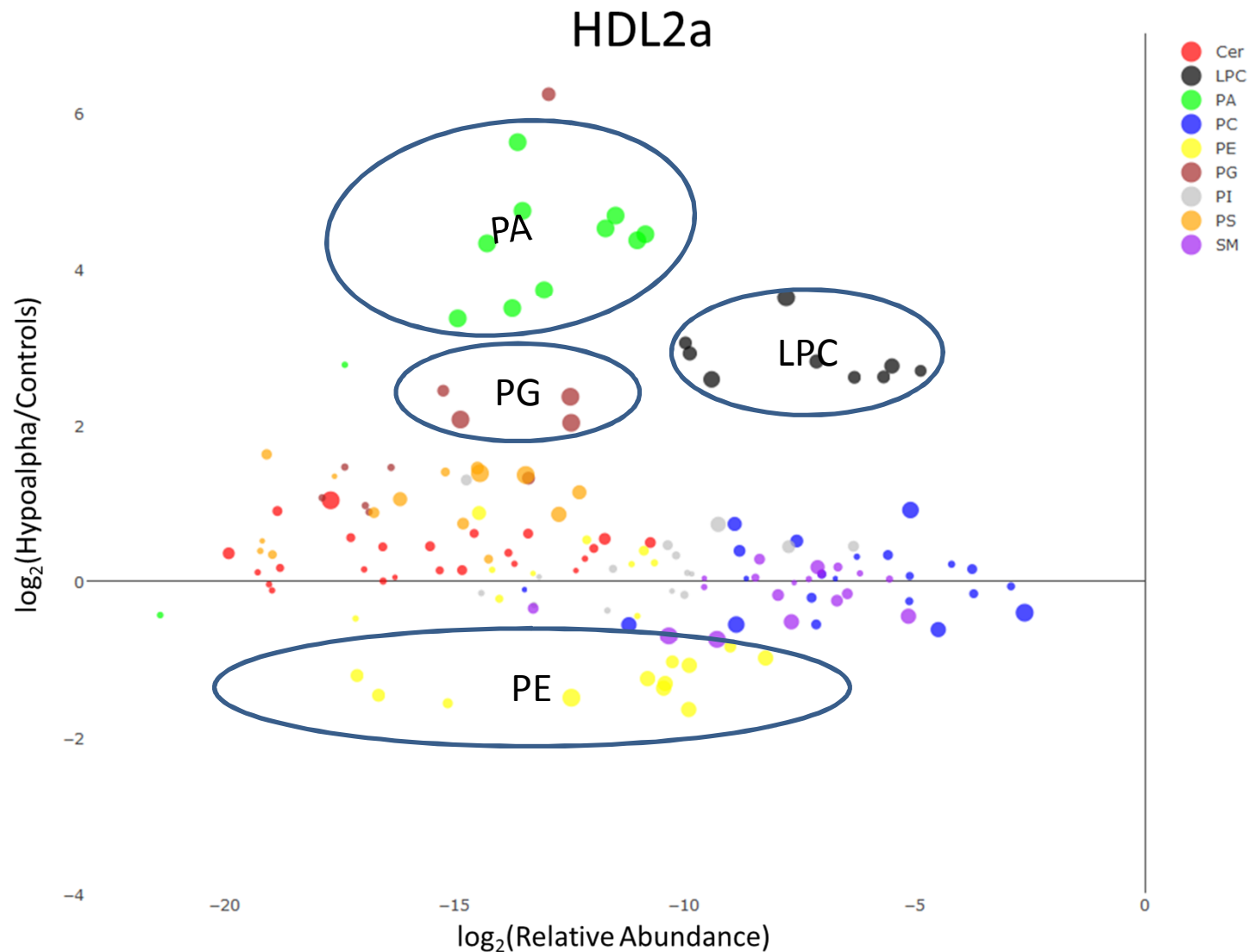
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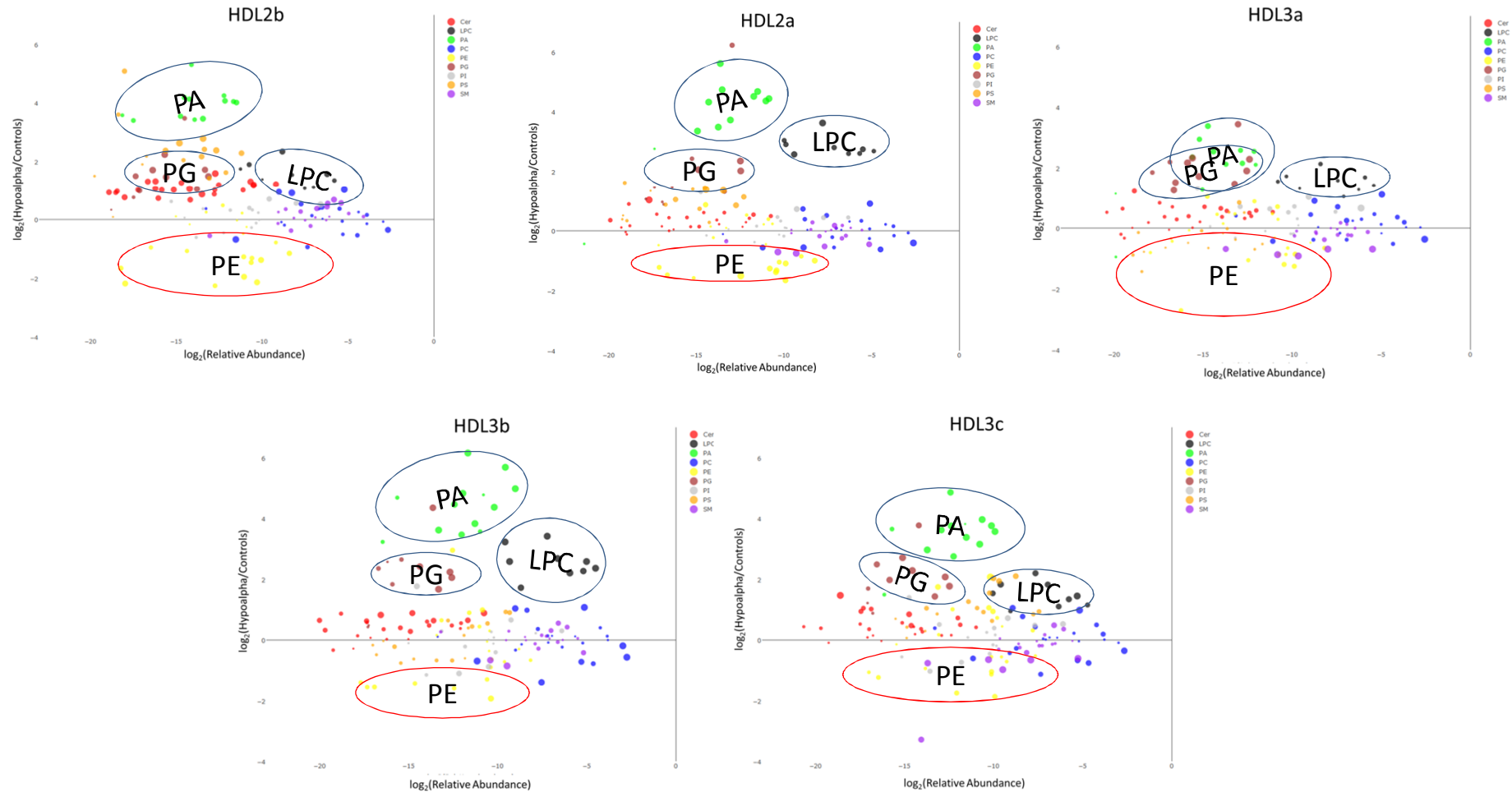
Bubble charts: Altered molecular species of LPC, PA, PG and PE are present in HDL at low-to-moderate abundance



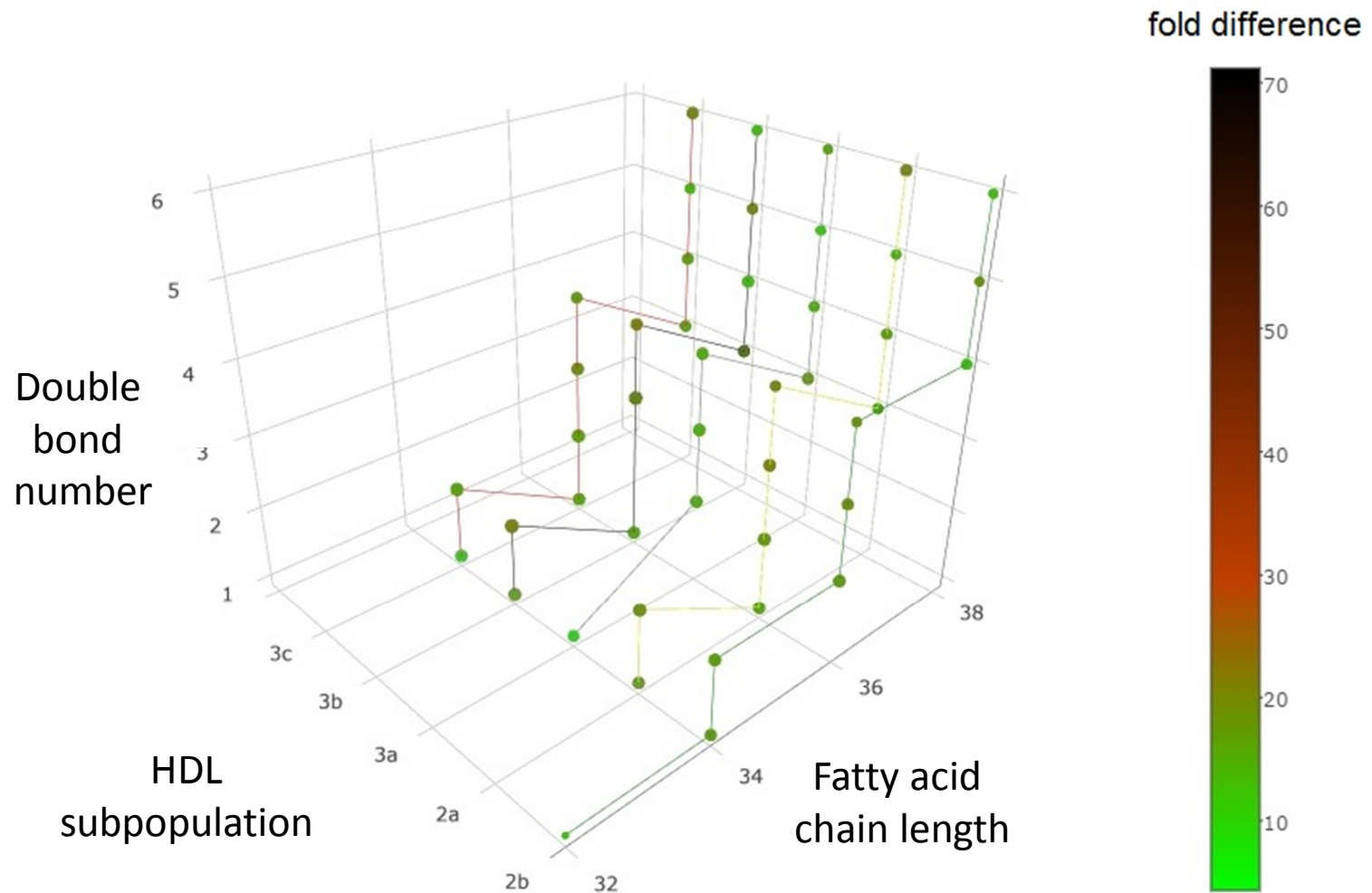
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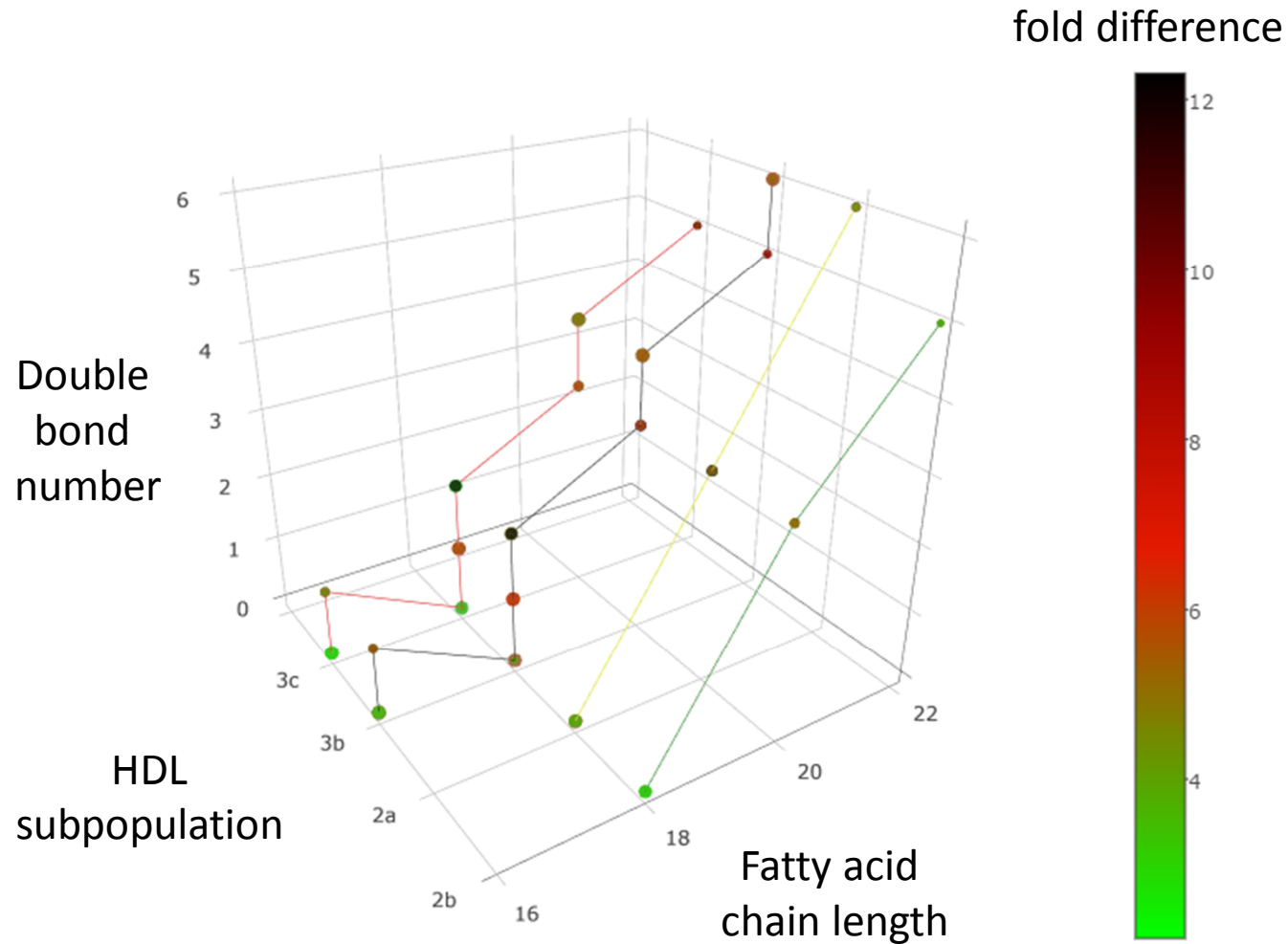
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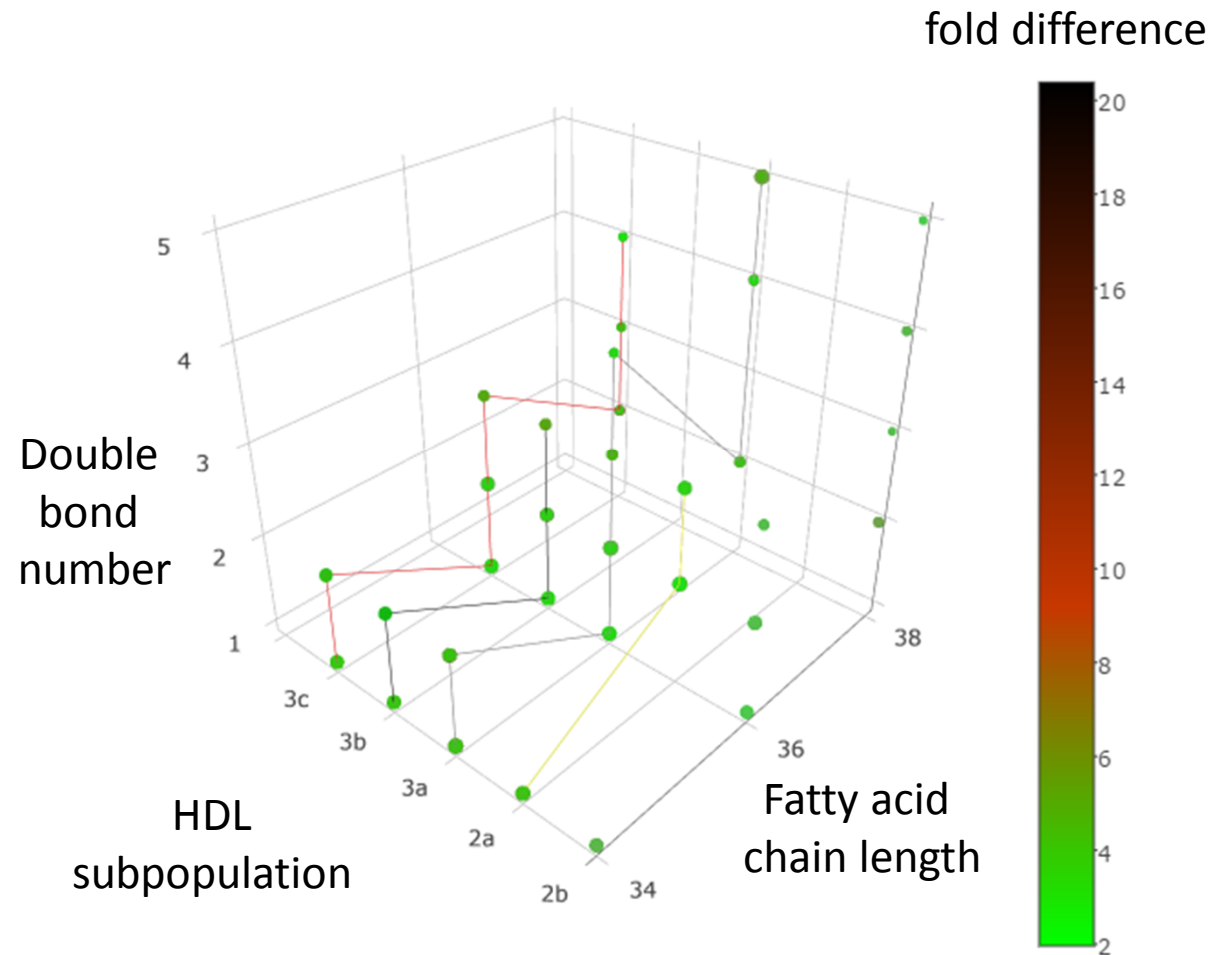
Structural analysis: PUFA-containing molecular species of PA (36:3, 36:4, 38:4, 38:5, 38:6) are markedly affected by apoA-I deficiency



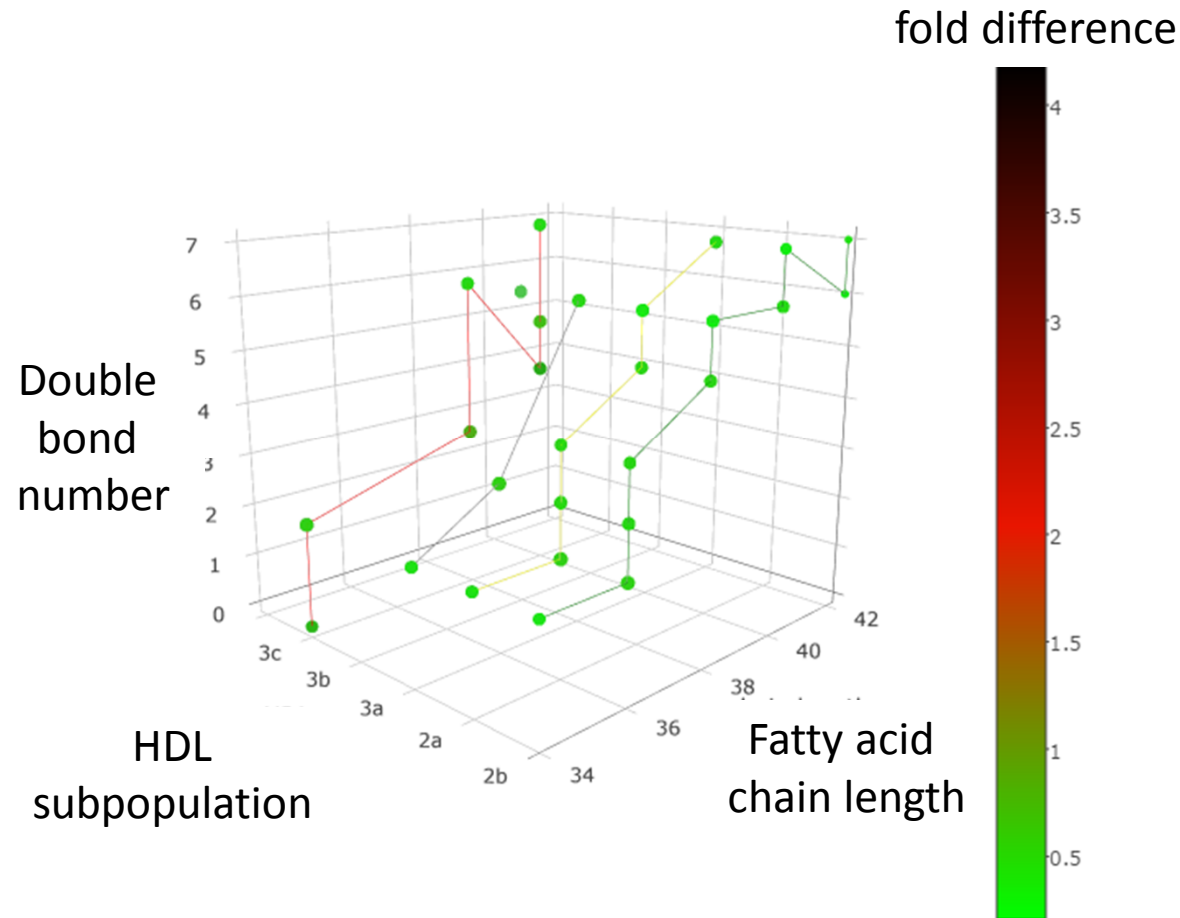
Structural analysis: PUFA-containing molecular species of LPC (18:2) are markedly affected by apoA-I deficiency



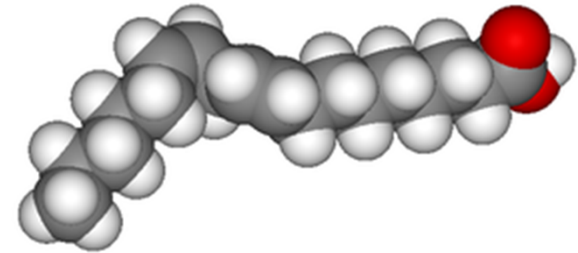
Structural analysis: PUFA-containing molecular species of **PG** (**36:2** and **36:3**) are markedly affected by apoA-I deficiency



Structural analysis: PUFA-containing molecular species of PE (34:2, 38:6) are markedly affected by apoA-I deficiency



Conclusions



- Low-to-moderately abundant, polyunsaturated fatty acid-containing phospholipid species of HDL, primarily PA(36:3), PA(36:4), PA(38:4), PA(38:5), PA(38:6), PG(36:2), PG(36:3), LPC(18:2), PE(34:2) and PE(38:6), are strongly affected by genetic apoA-I deficiency
- Elevated content of PA and LPC may reflect enhanced hydrolytic processing of HDL lipids
- Increased PA levels may result in activation of pro-inflammatory signalling pathways and may serve as biomarkers of impaired HDL function

Thank you for your attention!

