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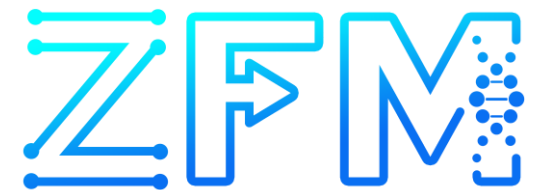
24-26 September 2025

**Nuclear and Medical Physics**

Faculty of Physics, Astronomy and Applied Computer Science,  
Jagiellonian University in Krakow, Poland

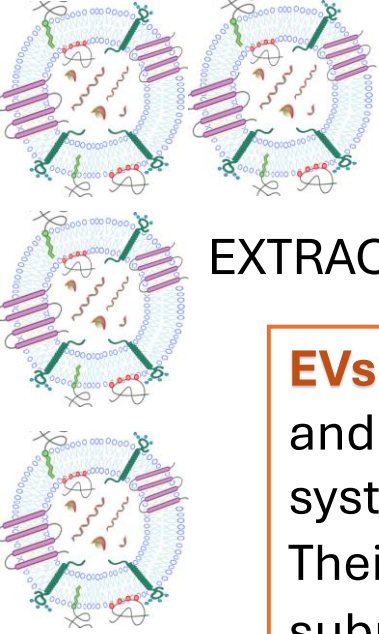
# Lipid Remodelling in Extracellular Vesicles from $\beta$ -Cells under Hyperglycemic Stress - Multimodal Mass Spectrometry Approach

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# Background & Rationale



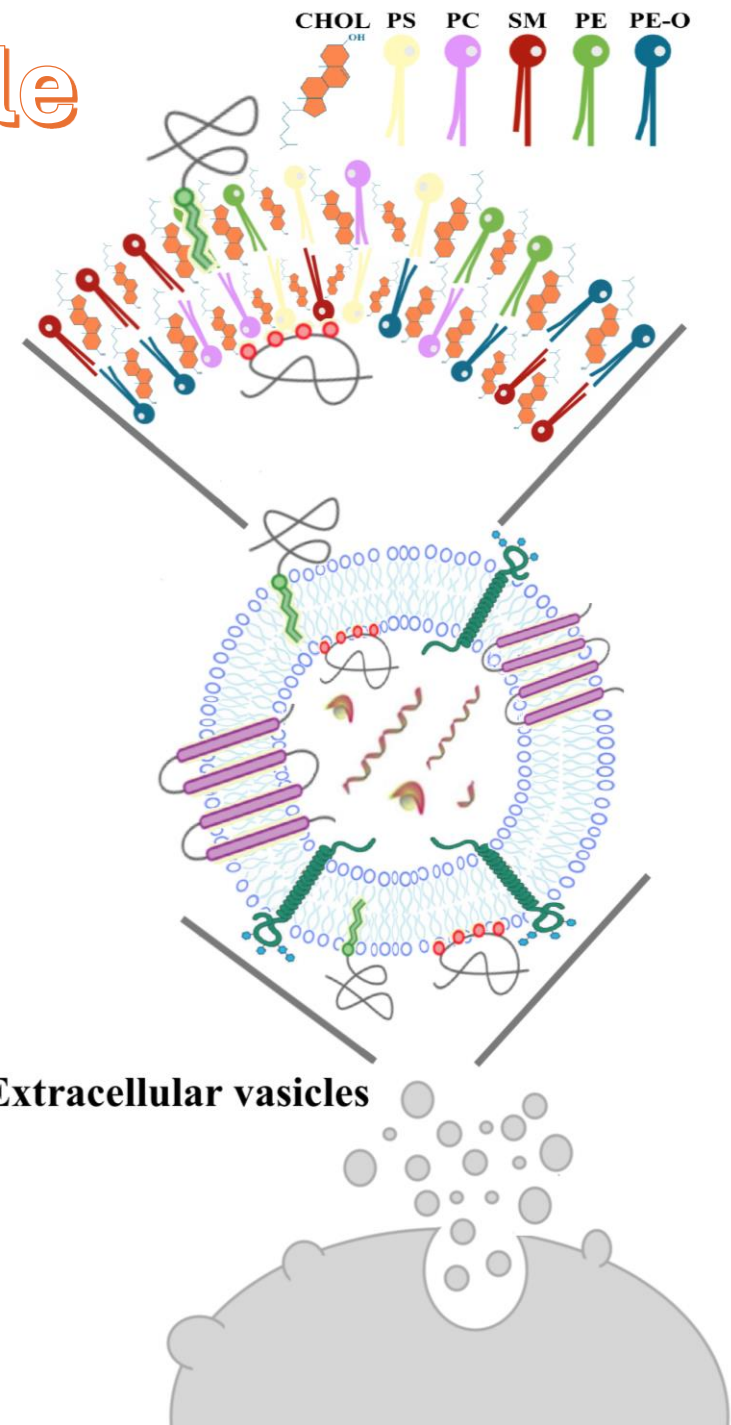
## EXTRACELLULAR VESICLES (EVs) AS BIOMARKERS

**EVs** are spherical structures surrounded by a lipid bilayer and contribute to a variety of functions in biological systems.

Their basic classification distinguishes three subpopulations based on their size and biogenesis:

- **exosomes** (50–150 nm) = **SMALL EVs**,
- **ectosomes** (100–1000 nm) = **LARGE EVs**
- **apoptotic bodies**.

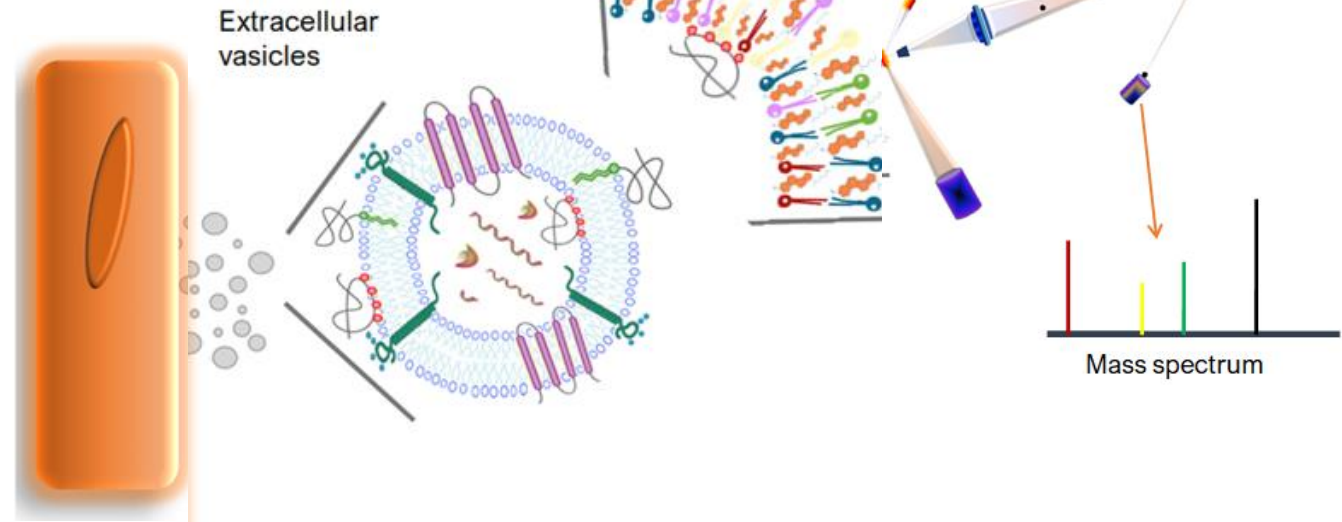
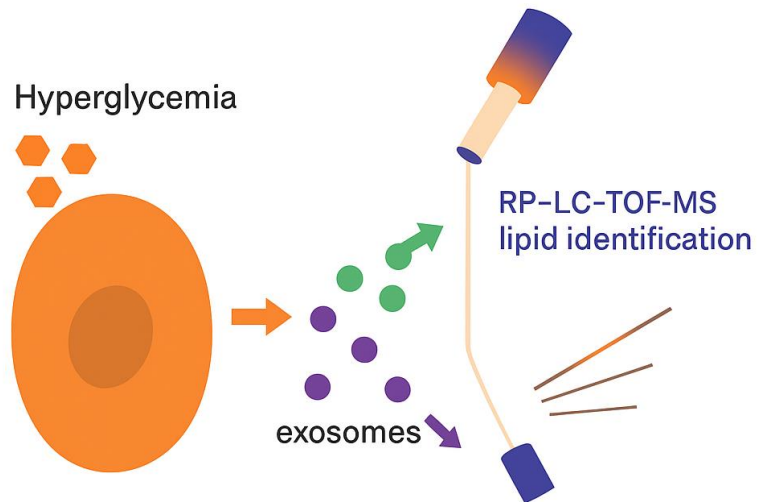
EV content varies according to the cell or organ of origin and the microenvironment at the time of their generation, determining in this manner their fate and biological activity.

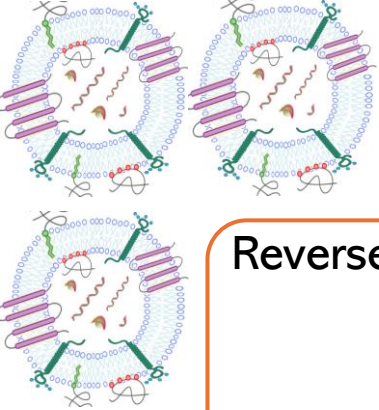




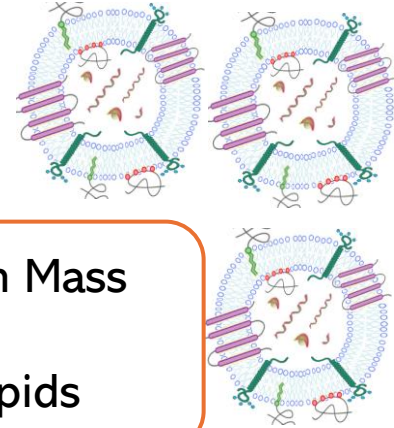
# Impact of hyperglycemia on $\beta$ -cell EV lipid composition

**To compare and discover molecular profile changes of EVs subpopulations derived from cells cultured in hyperglycemic conditions.**





# Analytical Techniques



Reversed-Phase Liquid Chromatography – Quadrupole  
Time-of-Flight Mass Spectrometry  
(RP-LC-Q-TOF-MS)  
high-resolution lipid identification

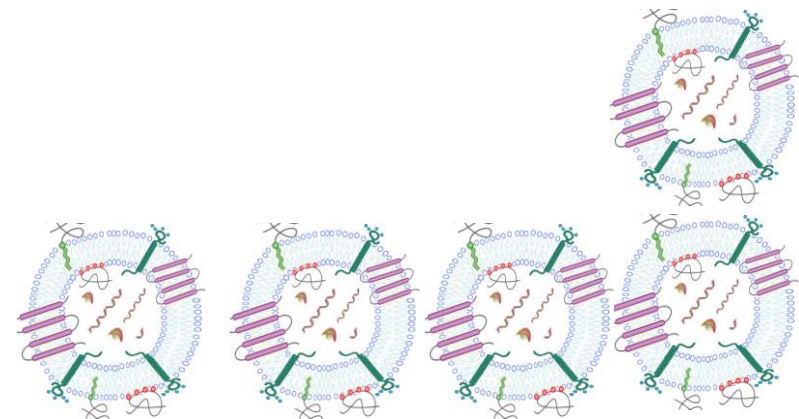
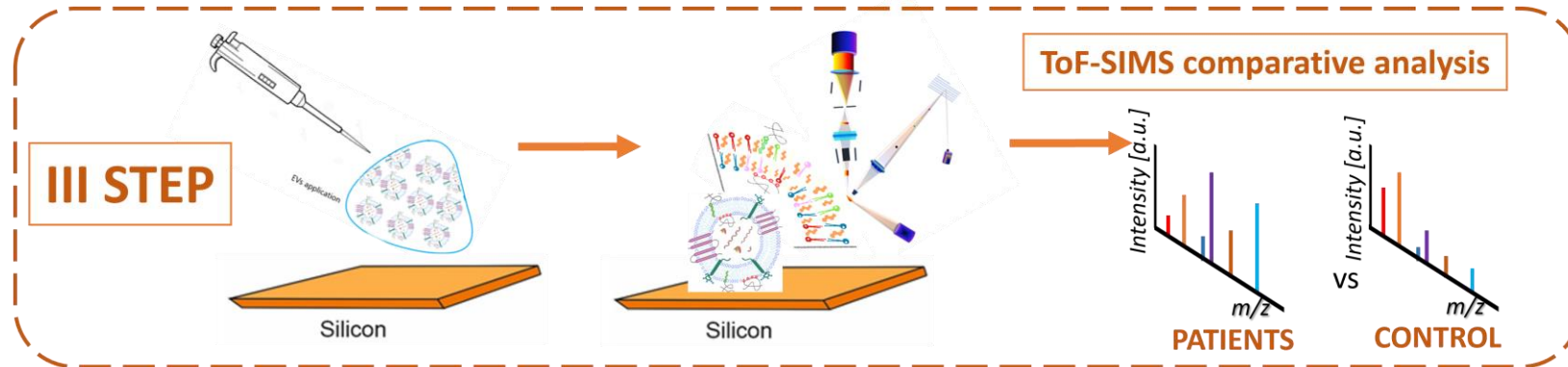
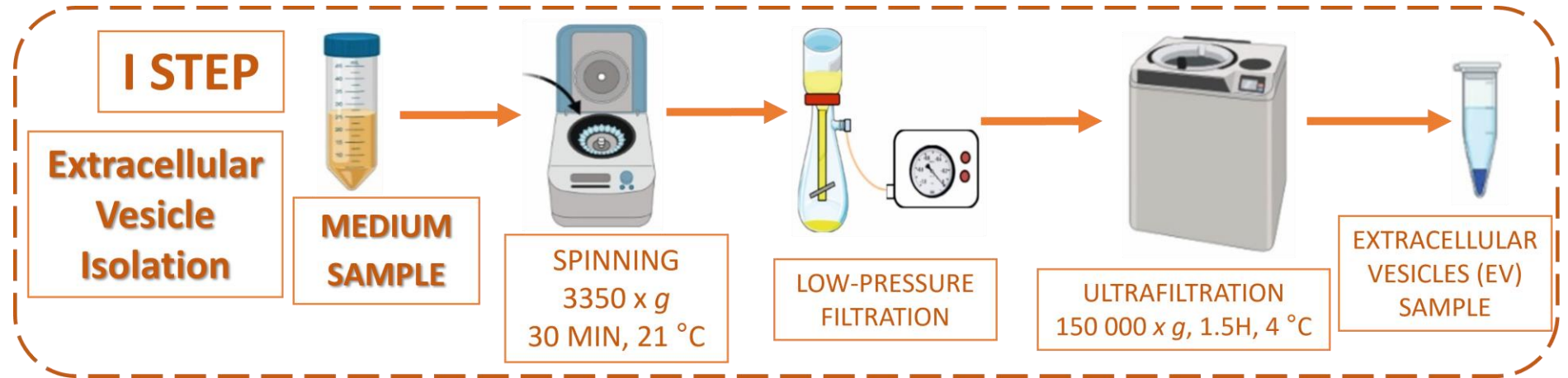
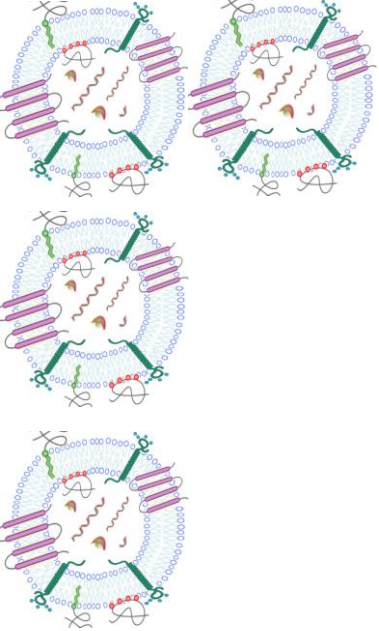
Time of Flight – Secondary Ion Mass  
Spectrometry  
direct surface mapping of lipids

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	RP-LC-Q-TOF-MS	ToF-SIMS
<b>Purpose</b>	High-resolution lipid ID	Surface mapping of lipids
<b>Advantages</b>	Accurate mass, broad lipid coverage, quantitative	Sub-micron imaging, minimal prep, label-free
<b>Limitations</b>	Requires extraction, no spatial info	Semi-quantitative, fewer intact lipids

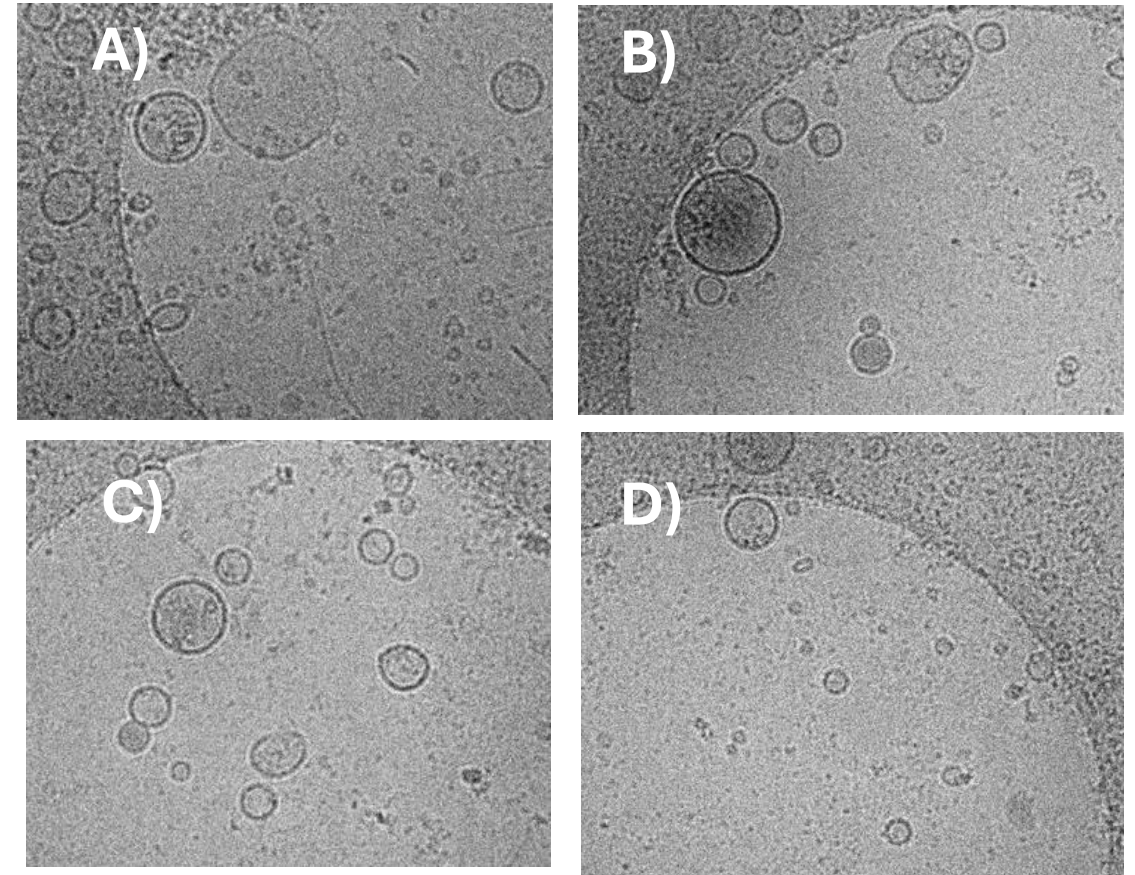
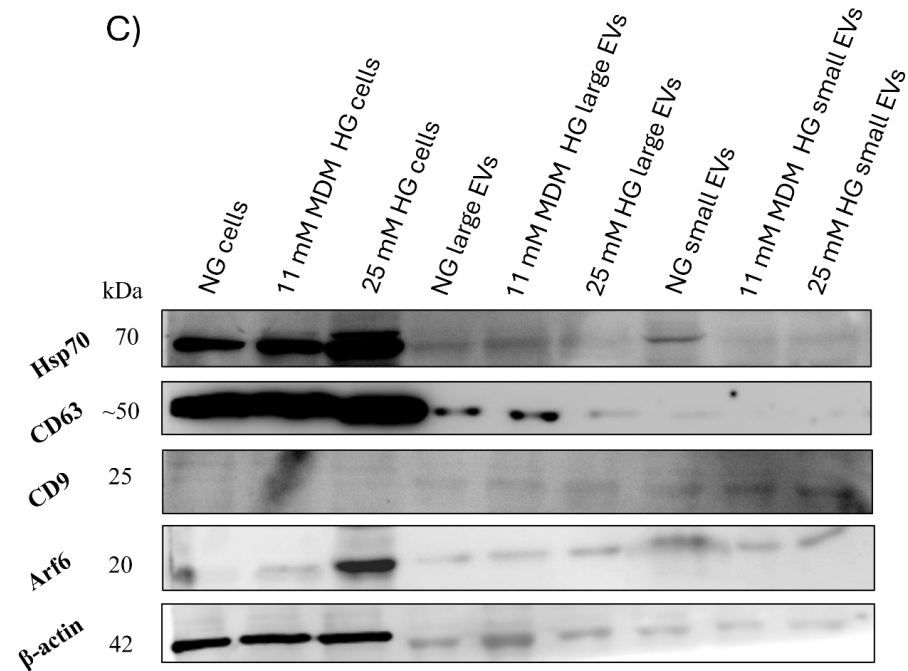
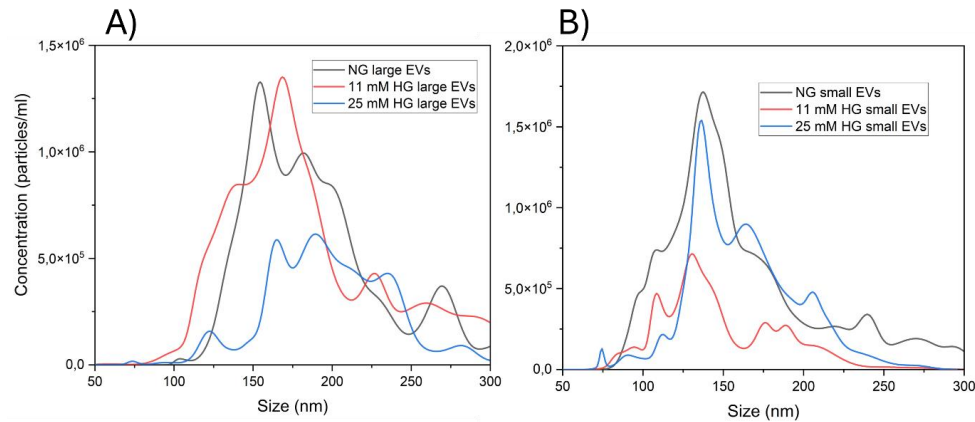
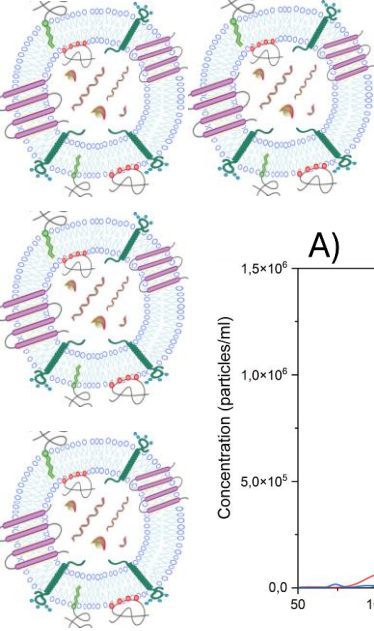
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# Experimental Design

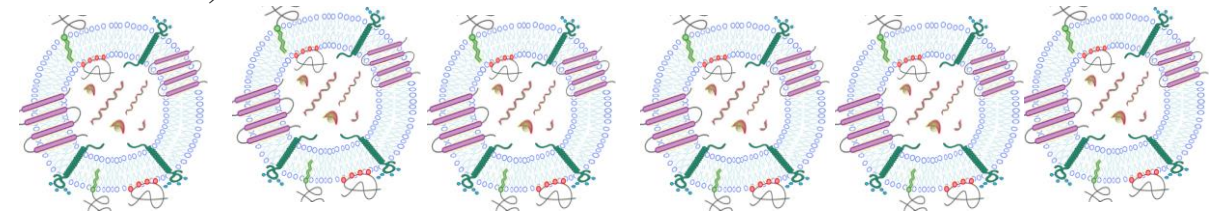




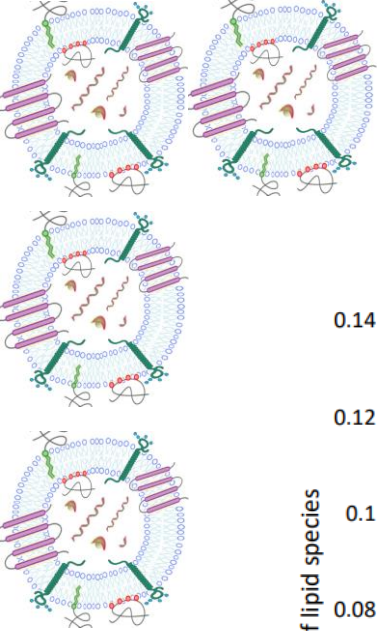
# RESULTS – Evs characterisation



*The cryo-TEM image of A) large HG-EVs, B) large NG-EVs, C) small HG-EVs, D) small NG-EVs.*

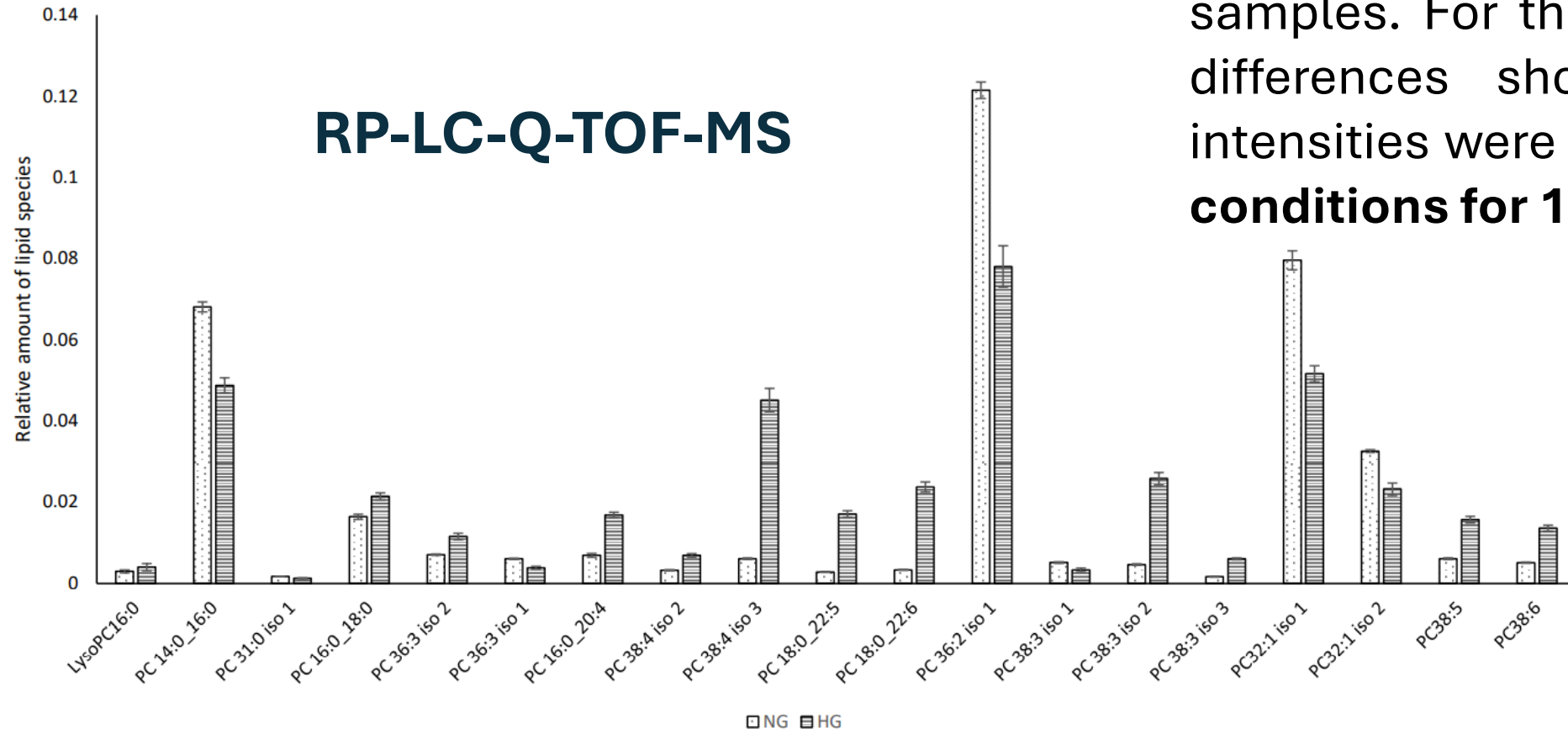


# ECTOSOMES



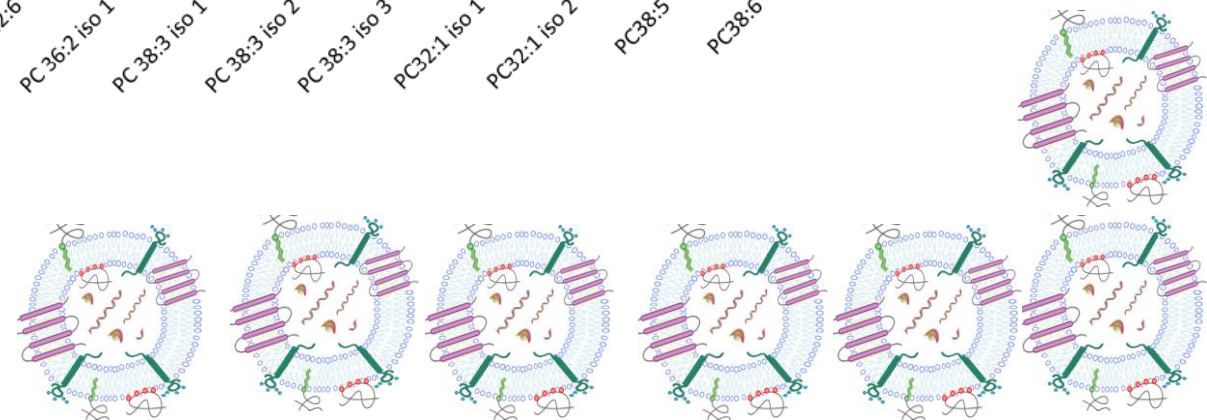
ectosomes -only significantly ( $p < 0.05$ ) changed PC

## RP-LC-Q-TOF-MS

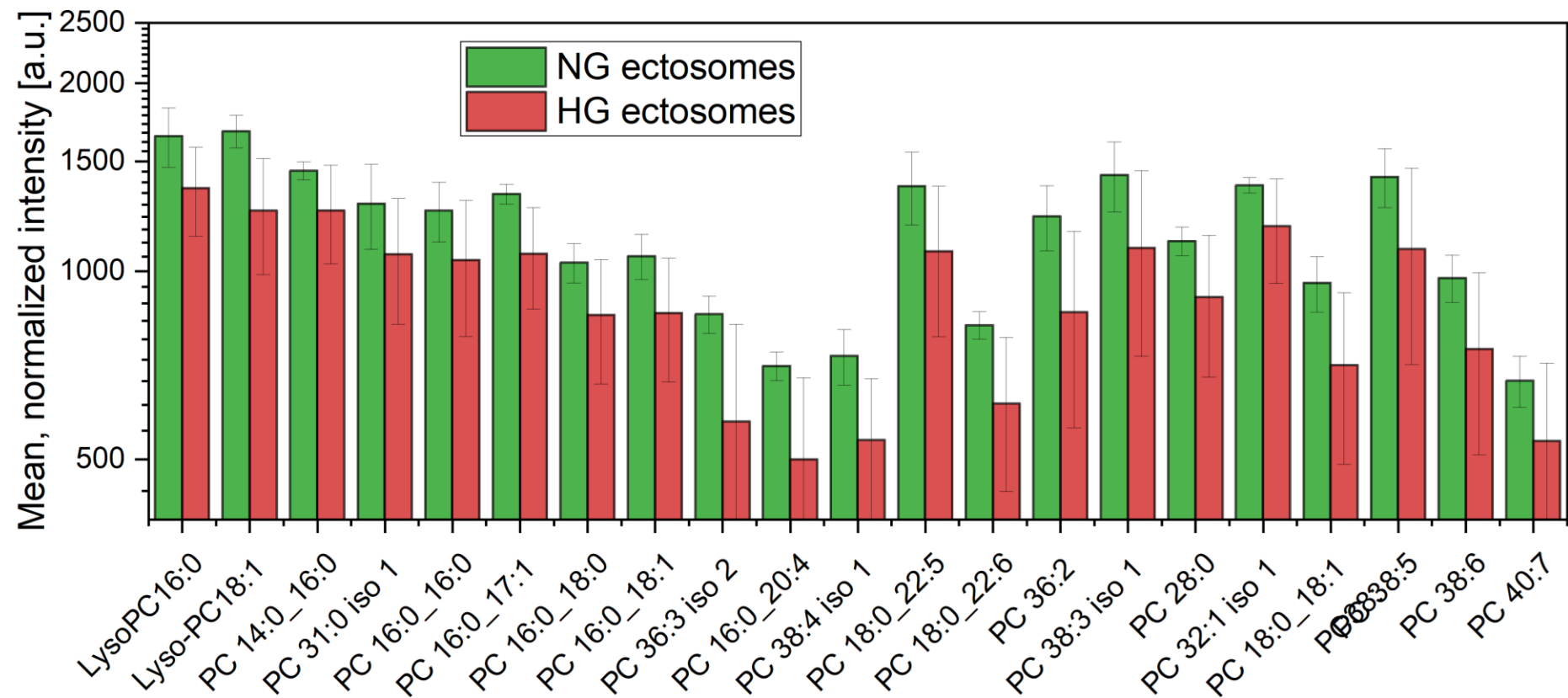
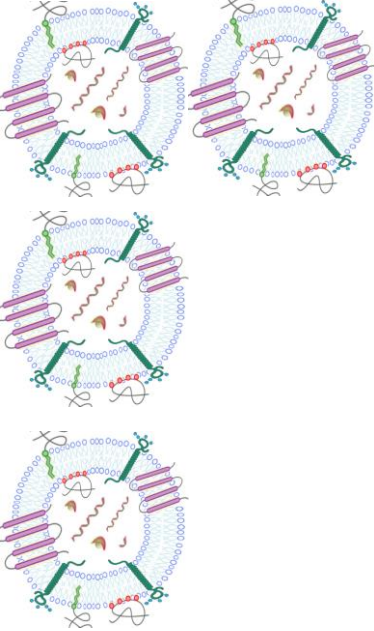


Of the 28 analysed PCs, 19 were statistically different between the NG and HG samples. For the significant differences shown, higher intensities were noted in **HG** conditions for 12 lipids.

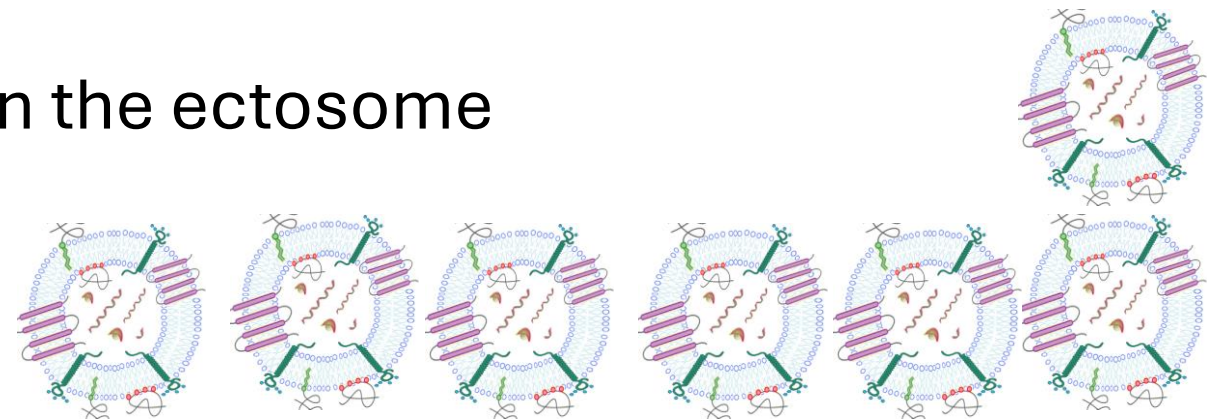
Comparison of PCs in NG and HG conditions. The error bars indicate SD corresponding to 3 independent biological replicates.



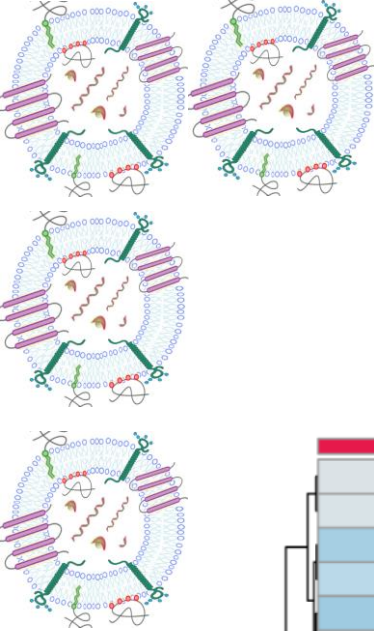




There are no statistical changes in the ectosome population.

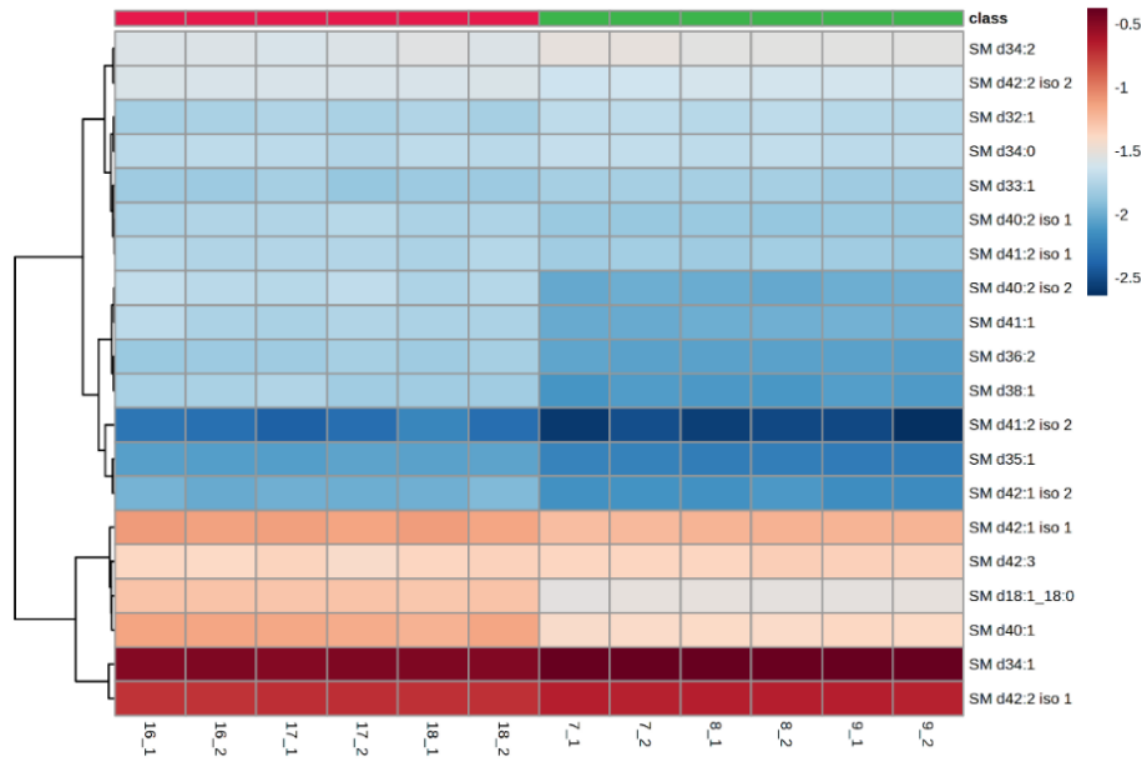




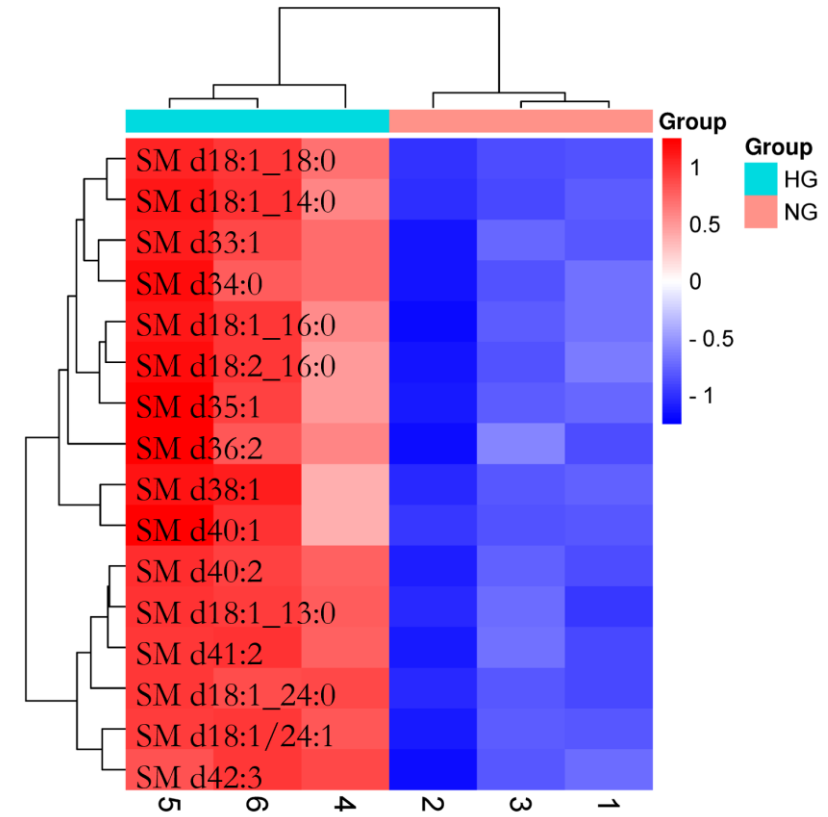


# Heatmaps presenting the sphingomyelins (SM) lipid profile in NG and HG conditions

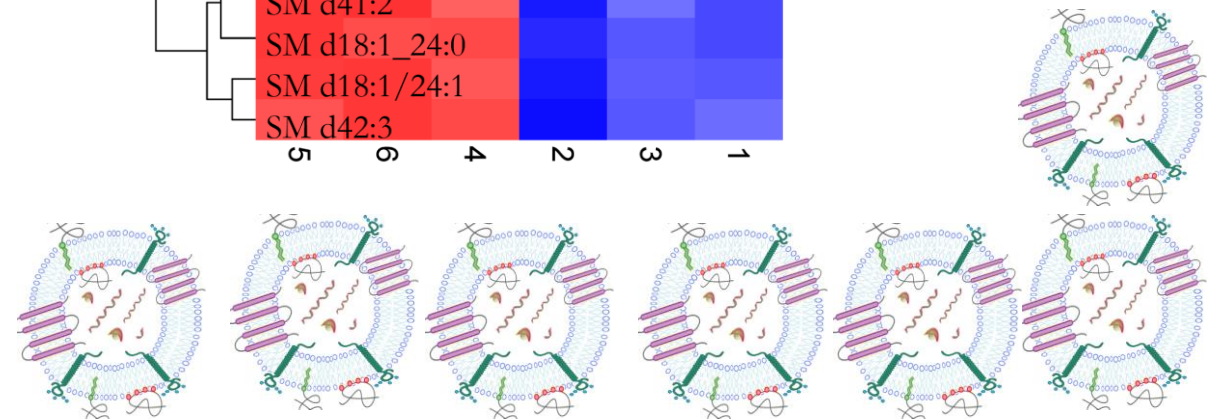
RP-LC-Q-TOF-MS



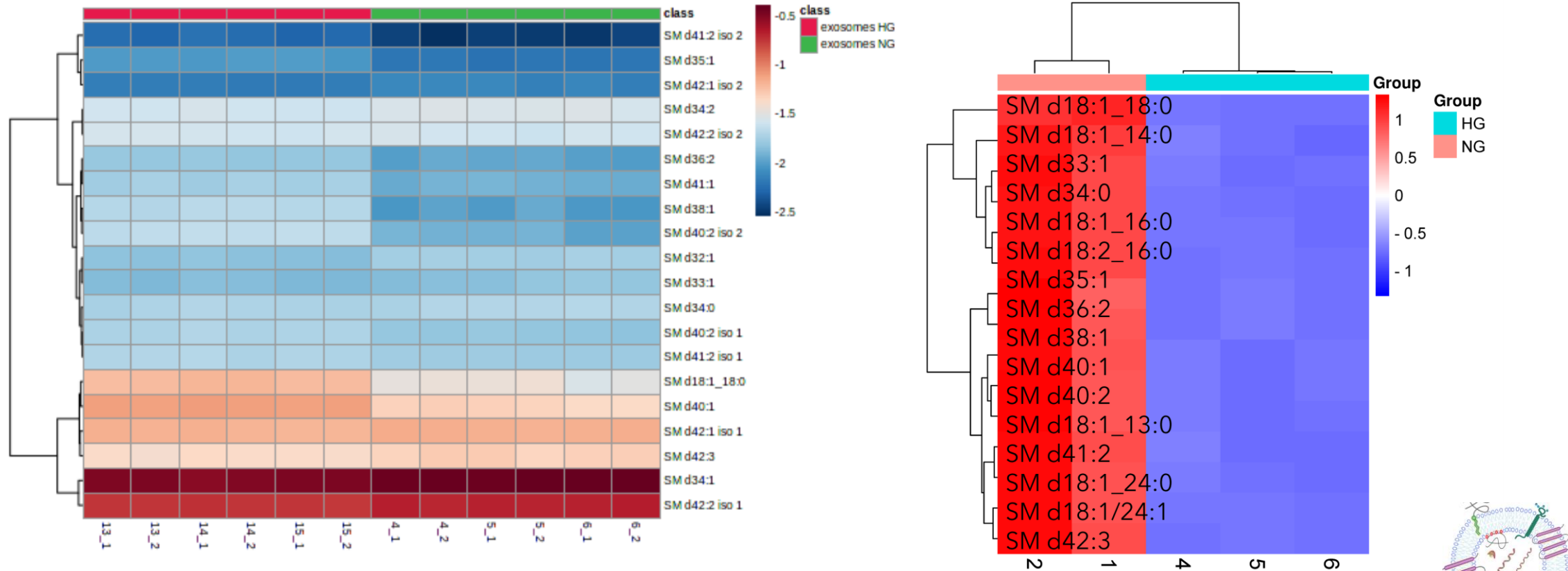
ToF-SIMS



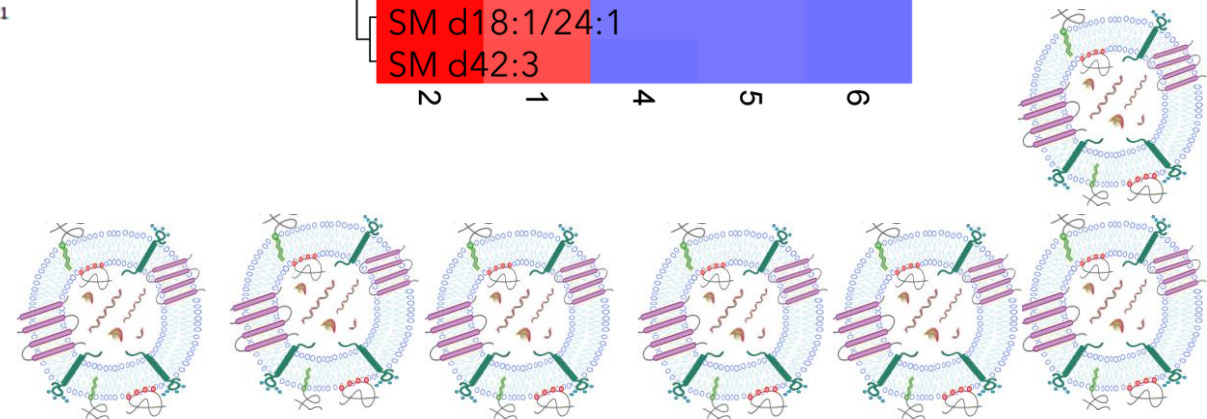
ectosomes



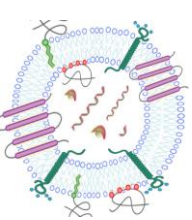
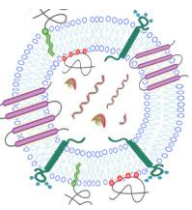
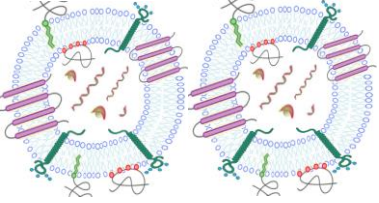
# Heatmaps presenting the sphingomyelins (SM) lipid profile in NG and HG conditions



**exosomes**



# CONCLUSION



**Exosomes** – RP-LC-Q-TOF-MS revealed significant increases in **8/20 sphingomyelins** and **19/28 glycerophospholipids** under hyperglycemia (HG). ToF-SIMS showed the same trend for surface SM but no major structural changes.

**Ectosomes** – LC-MS detected changes in **10/20 sphingomyelins** and **19/28 phosphatidylcholines** (12 higher in HG). ToF-SIMS again showed **no significant surface differences**.

## Overall Insight

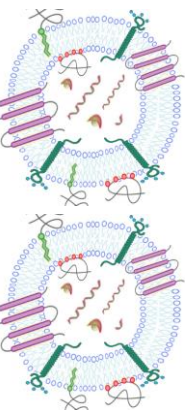
**Hyperglycemia enriches the internal lipid pool of EVs (especially SM and PC) while leaving the outer membrane largely unchanged.**

## Take Home Message

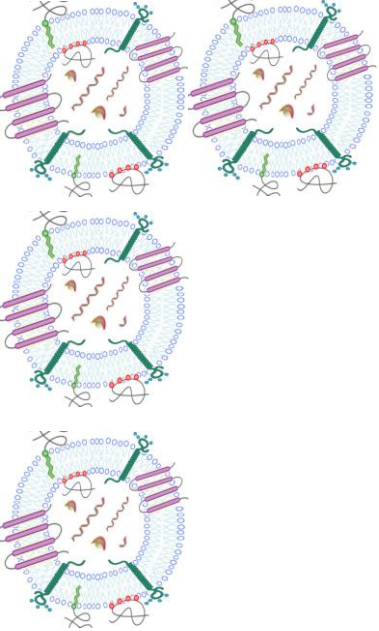
**Multimodal analysis** is essential:

- **LC-MS/MS** provides detailed molecular profiles.
- **ToF-SIMS** maps surface composition.

Together they reveal **bulk lipid remodelling without surface disruption**, highlighting EV lipids as potential **biomarkers of metabolic stress**.







# THANK YOU FOR YOUR KIND ATTENTION



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