

Phenotypic and metabolomic effects of atmospheric particulate matter from three different Catalonia locations on physiologically relevant human lung cell cultures

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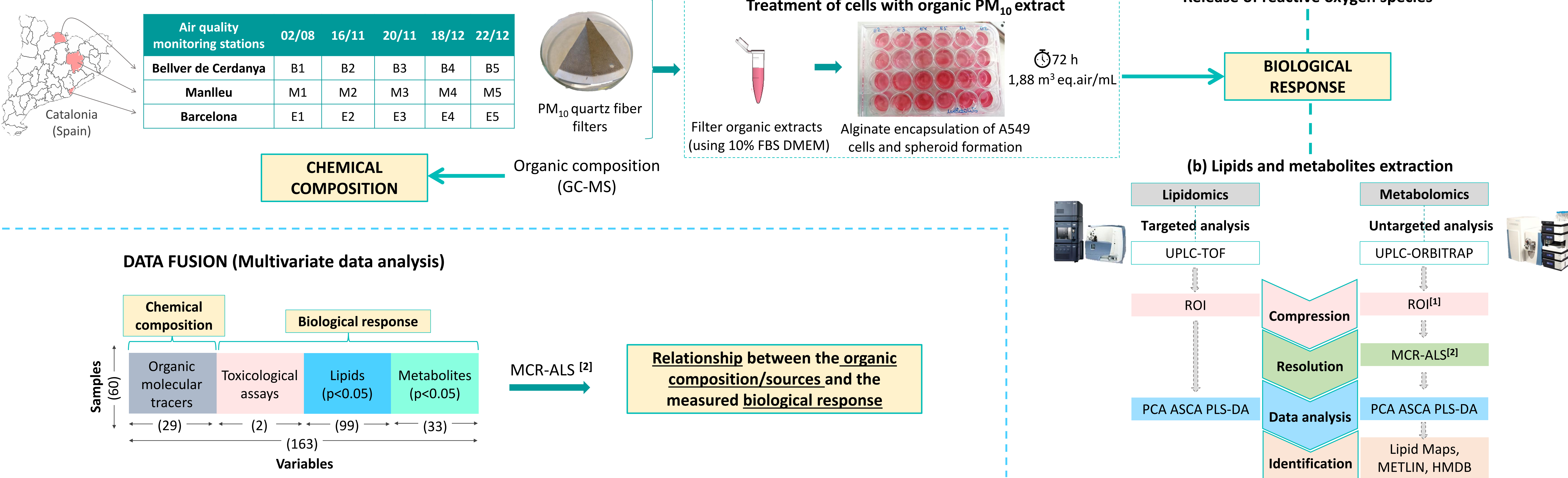
1 BACKGROUND

Airborne particulate matter (PM) pollution has raised great concern since its exposure has been associated to adverse respiratory outcomes. However, the underlying biological mechanisms that induce lung injury remain mostly unknown. In this work we propose the use of 3D bronchial cell cultures to study the biological effects of chemicals present in the organic soluble fraction of PM₁₀ by means of the combination of toxicological assays and metabolomic and lipidomic profiling. The application of multivariate data analysis methods are useful to establish the relationship between pollutants and their biological response.

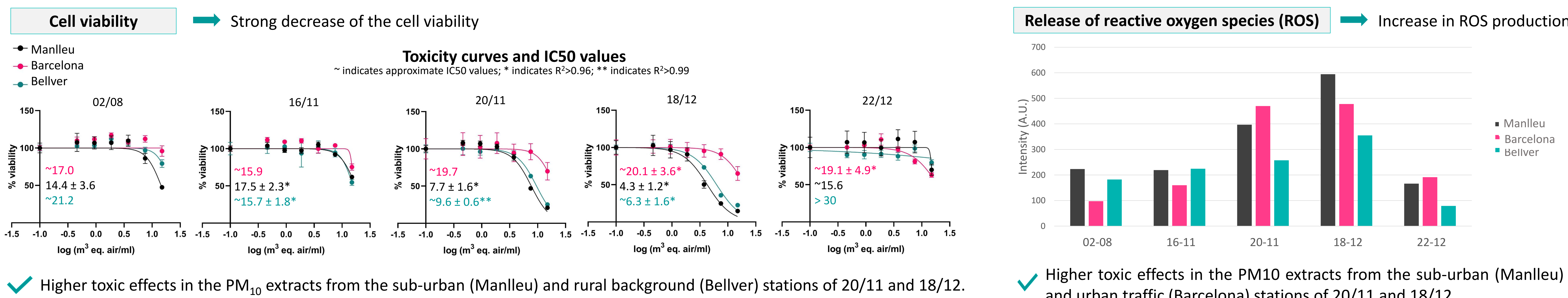
2 OBJECTIVE

→ Establishment of potential relationships between the organic composition/sources of PM₁₀ pollutions and the measured biological response by toxicologic studies (cell viability and ROS), lipidomics and metabolomics.

3 METHODOLOGY

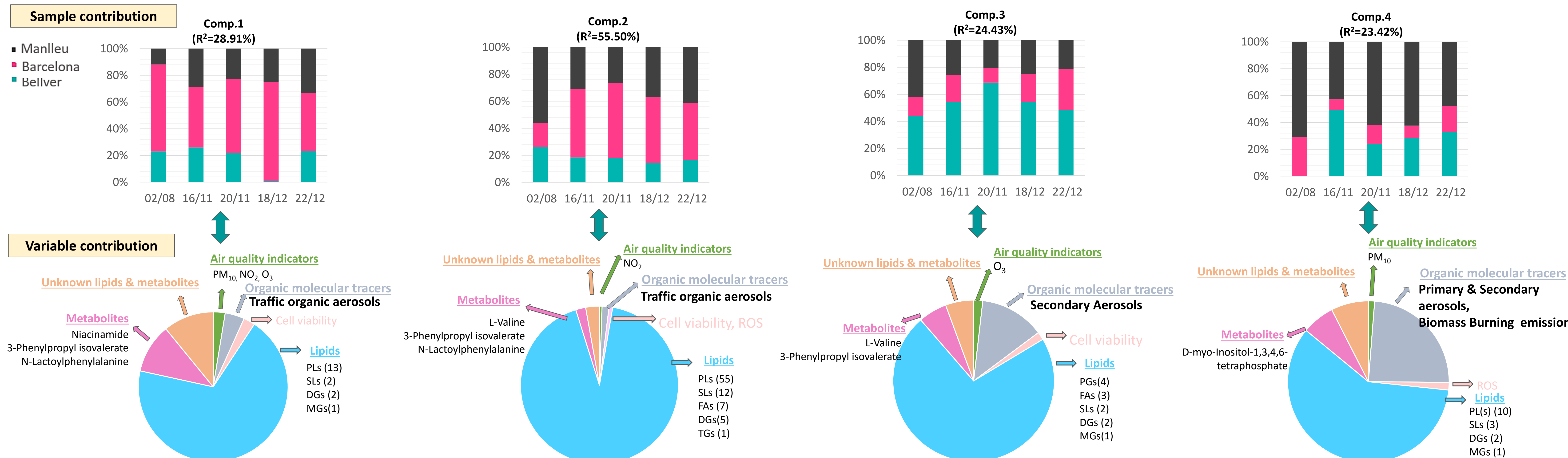


4 RESULTS



Data fusion analysis

→ 4 MCR-ALS resolved components, R²=91.85%



5 CONCLUSIONS

→ Phenotypic, metabolomic and lipidomic characterization of the PM₁₀ samples effects on A549 lung cells has been performed. Significant differences were found between locations and sampling days.

→ The joint analysis of chemical composition and omics data fusion enabled the identification of the most relevant relationships between PM₁₀ pollution sources and biological responses.

6 REFERENCES

- [1] Pérez-Cova, M., Bedia, C., Stoll, D. R., Tauler, R., & Jaumot, J. (2021). MSroi: a pre-processing tool for mass spectrometry-based studies. *Chemometrics and Intelligent Laboratory Systems*, 104333.
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