



# Childhood exposure to non-persistent endocrine disrupting chemicals and multi-omic markers in a population-based child cohort

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## Background & Objectives

- The general population is exposed to a cocktail of chemical exposures
- Non-persistent endocrine disruptors (EDCs) are a class of chemicals that interfere with the endocrine system
- The early stages of life are particularly vulnerable to the effects of EDCs
- Multi-omic signatures might provide mechanistic insights into the effect of EDC exposure, in particular before the onset of clinical symptoms in children
- We aimed to identify multi-omic signatures associated with non-persistent EDCs using an integrative approach based on Partial Correlation Networks

## Methods

### Study design



HELIX child study [1,2]

Week 1 - pool of 15 urines

Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7	Day 8
M	NM	NM	NM	NM	NM	NM	NM
							Blood

N<sub>t=1</sub>=117

~6-month interval

N<sub>t=2</sub>=120

Day 1	Day 2	Day 3	Day 4	Day 5	Day 6	Day 7	Day 8
M	NM	NM	NM	NM	NM	NM	NM
							Blood

Week 2 - pool of 15 urines

### Exposure assessment

Organophosphate (OP) pesticides (P=7)  
Phenols (P=10)  
Phthalates (P=5)  
(P<sub>tot</sub>=22)

### Omics

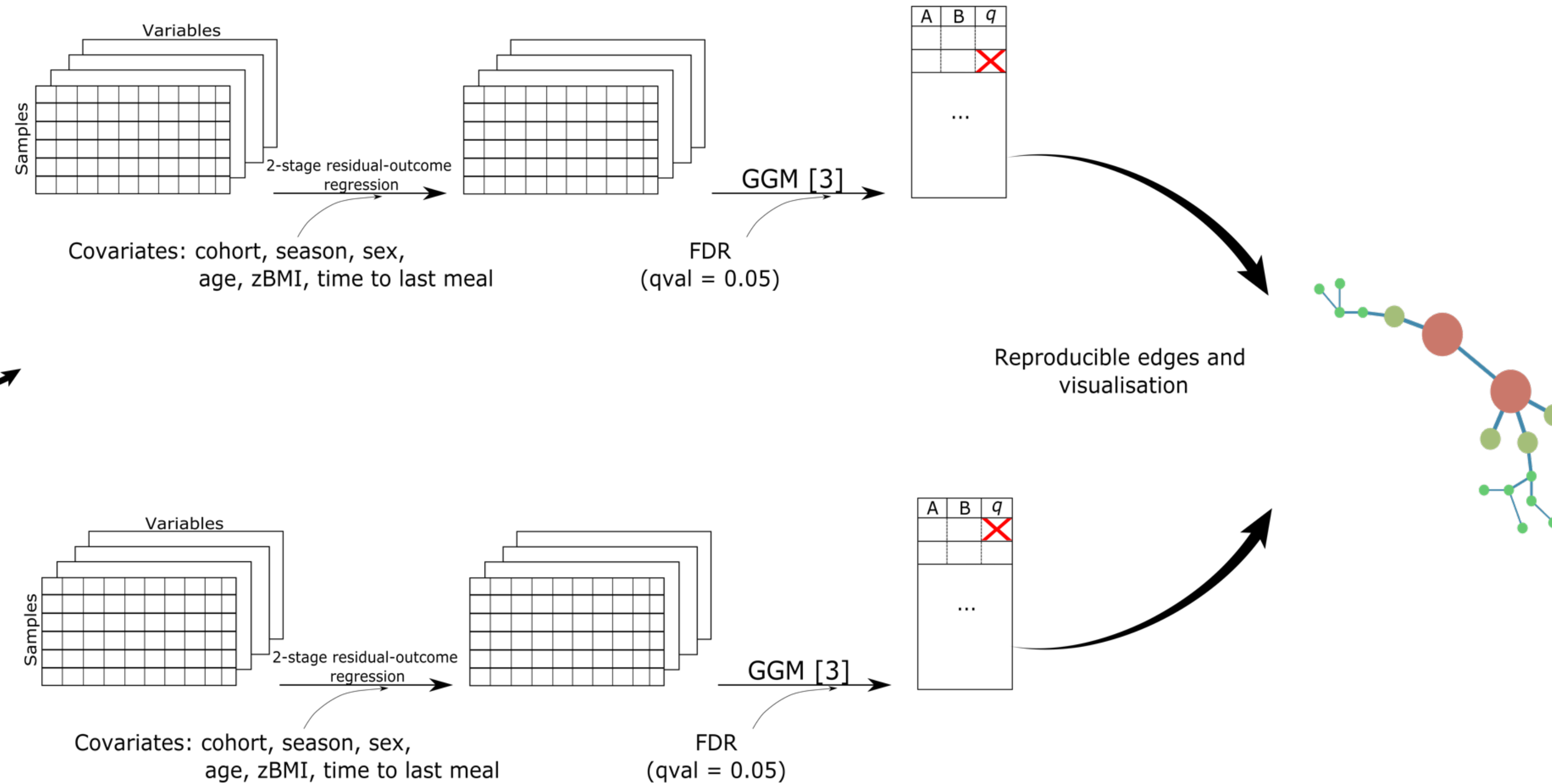
Serum metabolites (AbsoluteIDQ p180, P=177)  
Urine metabolites (<sup>1</sup>H NMR, P=44)  
Plasma proteins (Luminex assays, P=36)

Blood samples  
Urine samples

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### Multi-Omics Exposures networks



## Results

- The time-specific networks (N<sub>edges</sub>=1,064, N<sub>edges</sub>=1,109) included associations of comparable strength ( $\rho=0.09$  (-0.09, 0.11) for both) and statistical significance ( $q=0.008$  (0.001, 0.025),  $q=0.01$  (0.001, 0.027)). The significant edges represented less than 3% of the possible connections
- The merged network consisted of N<sub>edges</sub>=229
- Graph merging led to the exclusion of the majority of exposure-omic connections (Figure 1). Notably, none of the protein-exposure associations were reproducible
- The merged network consisted of 32 connected components, 3 of which included mixed exposure-omic connections (Figure 2)

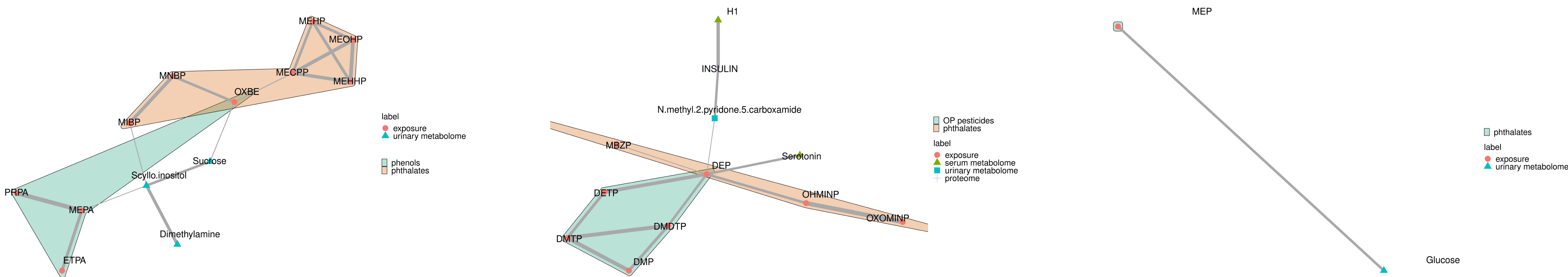


Figure 2. Clusters (i.e. connected components) of EDC exposure-omic associations.

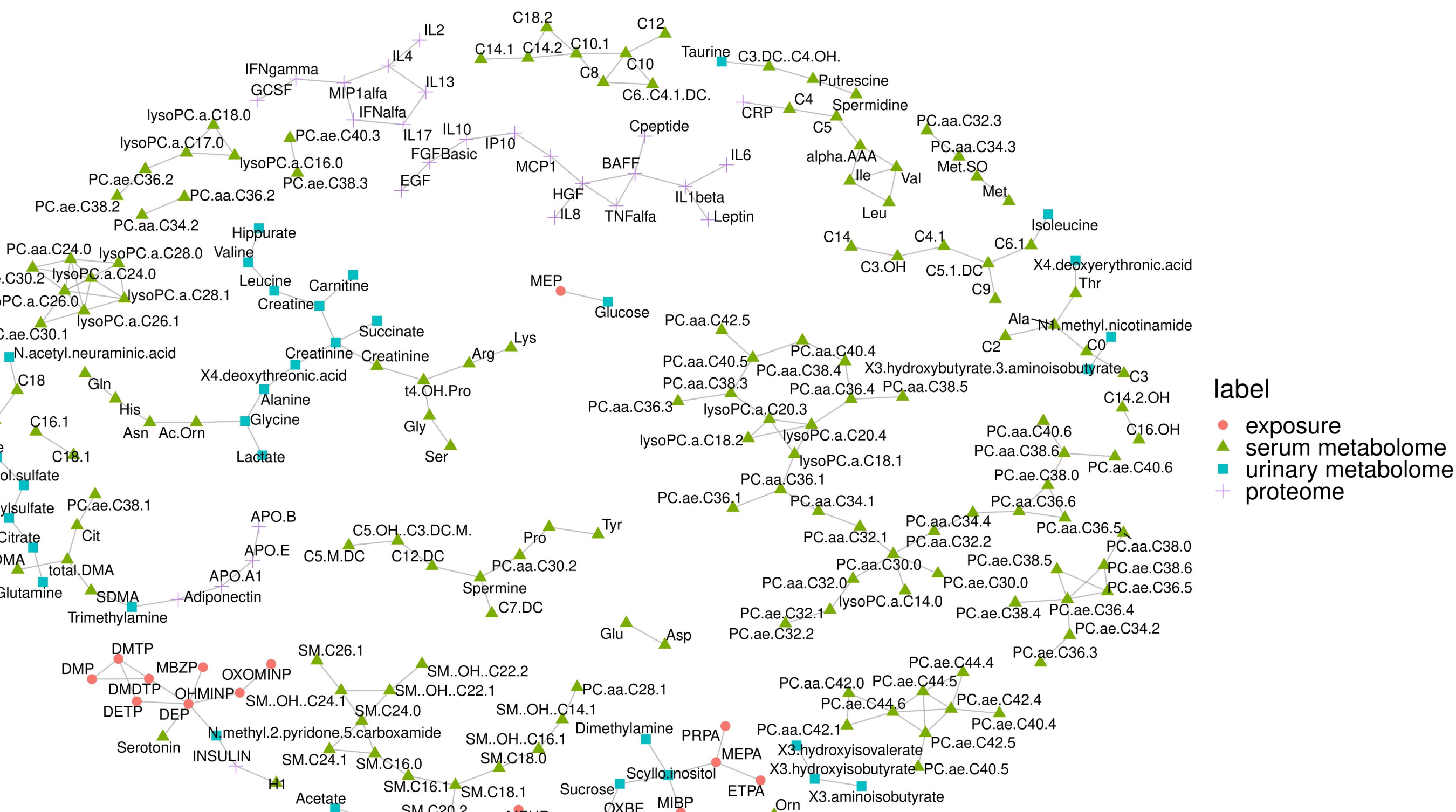


Figure 3. Merged network showing all the connected components.

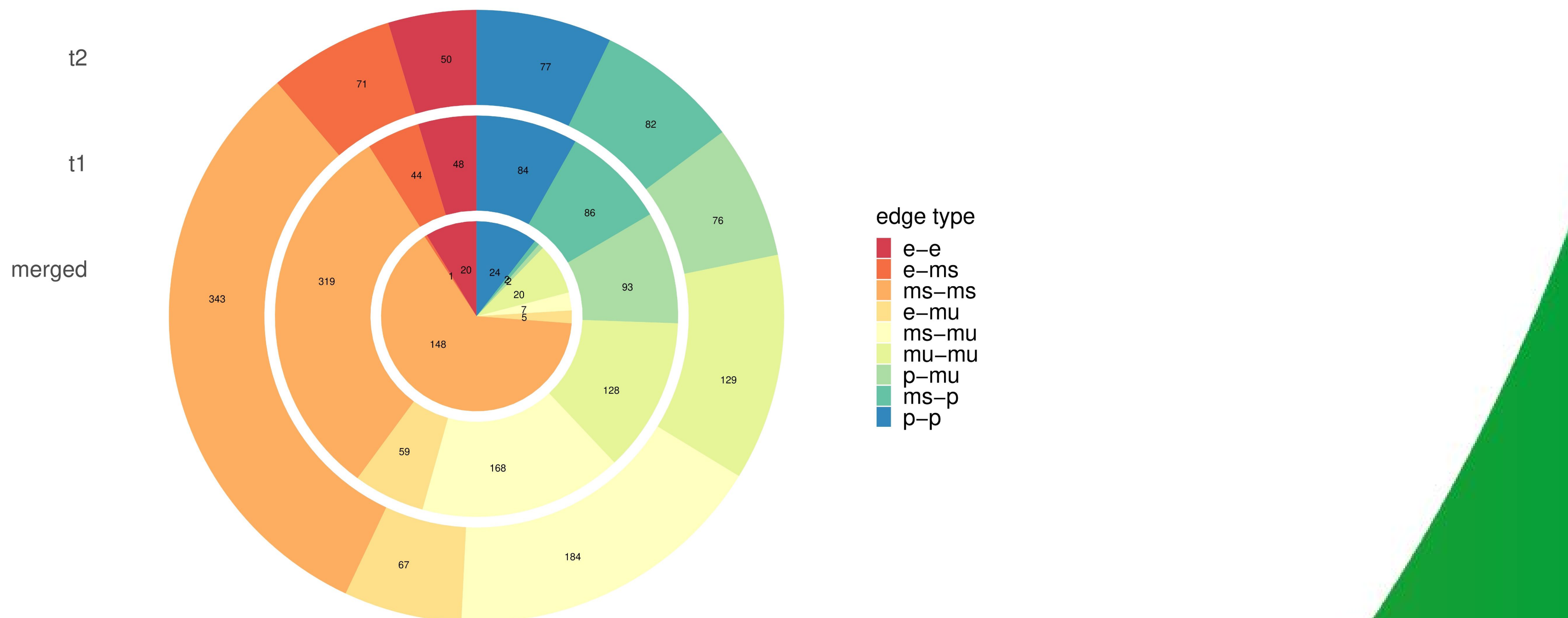


Figure 1. Multilevel pie chart showing the proportion of edge types for the time-specific and merged networks. e = exposure, ms = serum metabolite, mu = urinary metabolite, p = protein.

## Conclusions

- We integrated Multi-Omic and exposure data from a child cohort using an integrative approach, and we identified associations reproducible across time points
- The association between DEP and Serotonin ( $\rho=0.09$  for both time points) was reproducible. Exposure to Organophosphate pesticides has been linked to a variety of brain disorders [4], potentially through the serotonergic system
- In future work we plan to include methylation data

## References

[1] Vrijheid M, et al. "The human early-life exposome (HELIX): project rationale and design." (2014). [2] Casas M, et al. "Variability of urinary concentrations of non-persistent chemicals in pregnant women and school-aged children." (2018). [3] Schafer J, Strimmer K. "A Shrinkage Approach to Large-Scale Covariance Matrix Estimation and Implications for Functional Genomics." (2005). [4] Sarrouilhe D, et al. "Is the Exposome Involved in Brain Disorders through the Serotonergic System?" (2021)

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