

## Microarray data

- Chronic exposure (E-MEXP-1828)
- Acute exposure (E-MEXP-1823)

Probes to genes  
(medians)

Keep more variable genes  
(DCGL R package)

Classification of genes in modules  
(WGCNA R package)

Based on modules from each condition

- Differential Co-expression Analysis  
(DCLR R package)

Enrichment analysis

- TFBS  
(Babelomics 4.3)