

Ensemble of Gene Set Enrichment Analyses (EGSEA) - Report

Analysis Parameters

Total number of genes: 17343

Total number of samples: 8

Number of contrasts: 2

Base GSEA methods: camera,safe,gage,padog,plage,zscore,gsva,ssgsea,globaltest,ora

P-value combine method: fisher

Sorting statistic: avg.rank

Fold changes calculated: Yes

Gen IDs - Symbols mapping used: Yes

Organism: Homo sapiens

Analysis Results

X24IL13 - X24

- c2 Curated Gene Sets ([Stats Table](#), [Heatmaps](#), [Summary Plots](#), [Download Stats](#))
- KEGG Pathways ([Stats Table](#), [Heatmaps](#), [Pathways](#), [Summary Plots](#), [Download Stats](#))

X24IL13Ant - X24IL13

- c2 Curated Gene Sets ([Stats Table](#), [Heatmaps](#), [Summary Plots](#), [Download Stats](#))
- KEGG Pathways ([Stats Table](#), [Heatmaps](#), [Pathways](#), [Summary Plots](#), [Download Stats](#))

Comparison Analysis

- c2 Curated Gene Sets ([Stats Table](#), [Heatmaps](#), [Summary Plots](#), [Download Stats](#))
- KEGG Pathways ([Stats Table](#), [Heatmaps](#), [Pathways](#), [Summary Plots](#), [Download Stats](#))