10 common differentially expressed biological processes

Cluster analysis (3 clusters) enriched GO-terms and chromosomal regions in each cluster

CCA (21 chromosomal regions)

Cluster analysis (12 clusters) enriched GO-terms and chromosomal regions in each cluster

Cluster analysis (16 clusters) enriched GO-terms and chromosomal regions in each cluster

Cluster analysis (3 clusters) enriched GO-terms and chromosomal regions in each cluster

GO analysis

number of up (grey) and down (black) regulated GO-terms

0 1h 6h 24h 48h 7 days

0 1h 6h 24h 48h

10 common differentially expressed biological processes