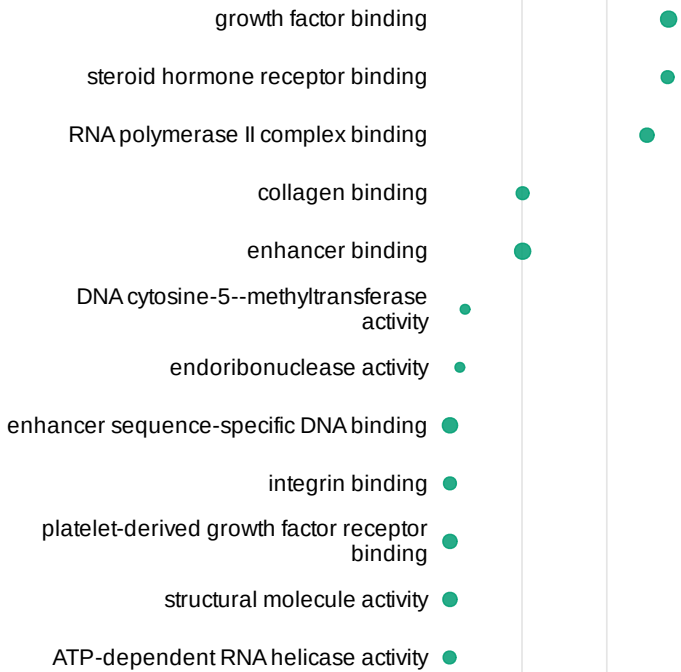
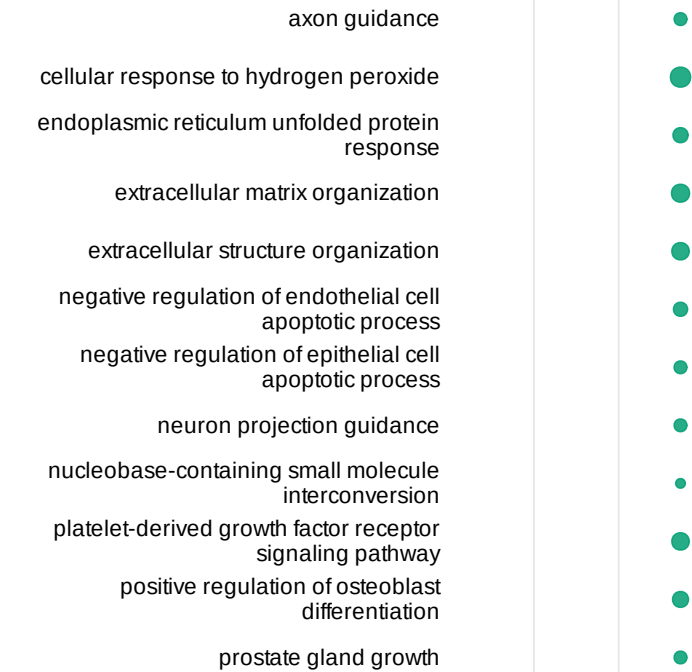


miRPathDB Enriched: GENERO\_1\_vs\_0

GO Biological  
process  
(miRPathDB)

GO Molecular  
function  
(miRPathDB)

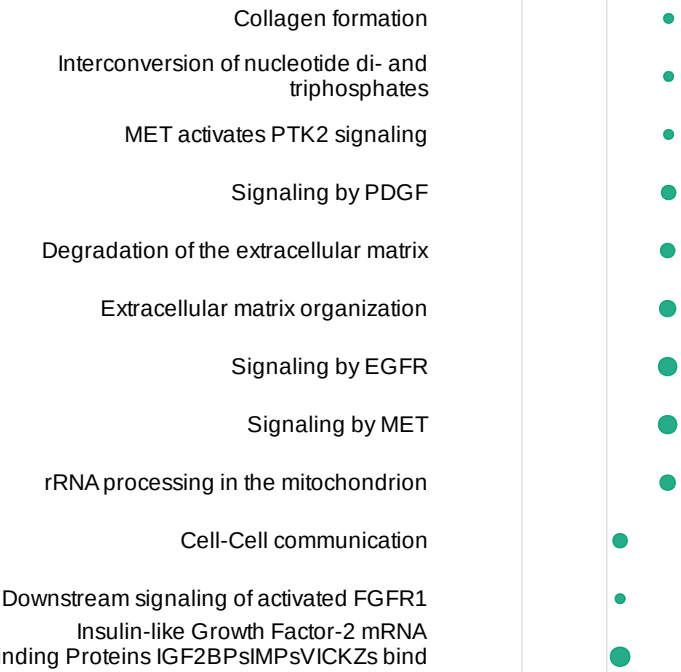
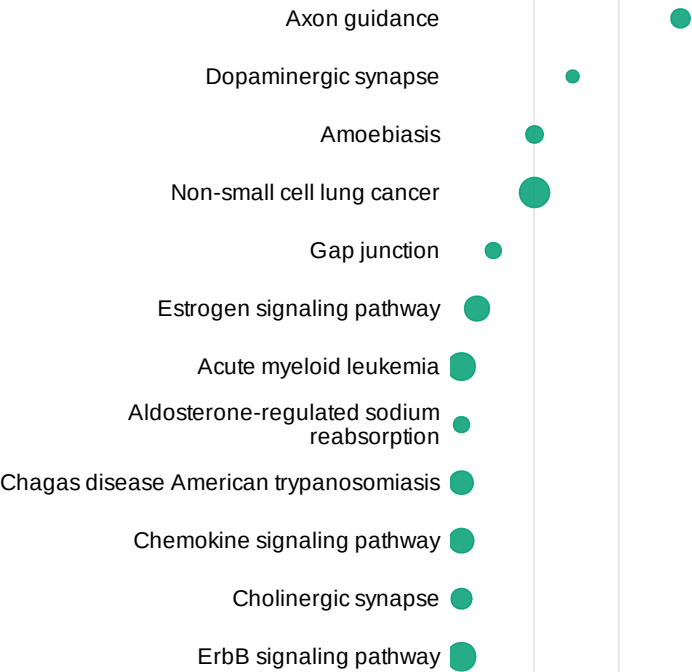


Observed

- 10
- 20
- 30
- 40

KEGG  
(miRPathDB)

Reactome  
(miRPathDB)



-log10(P-adjusted)