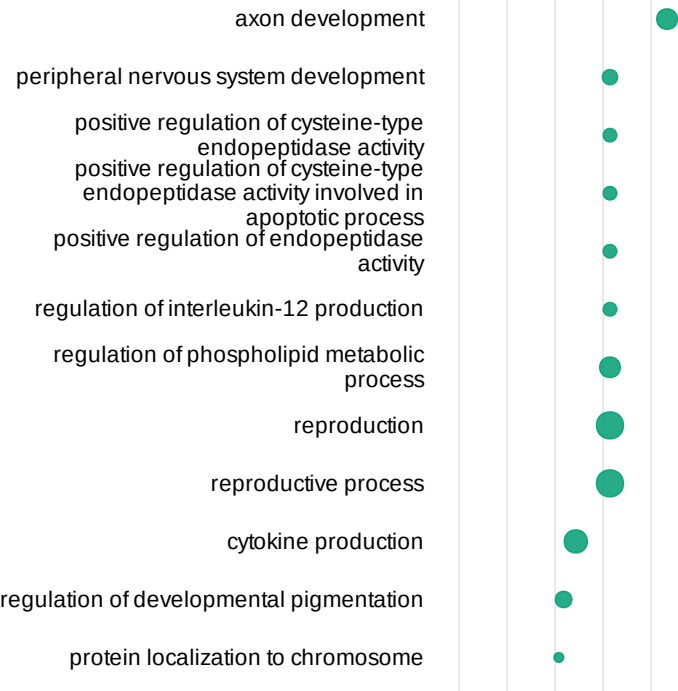


miRPathDB Enriched: P53_TP53_1_vs_0

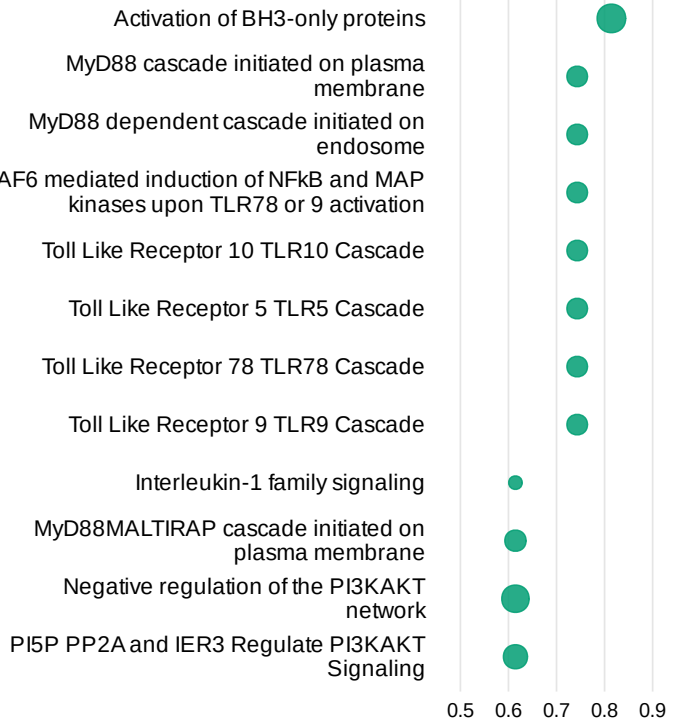
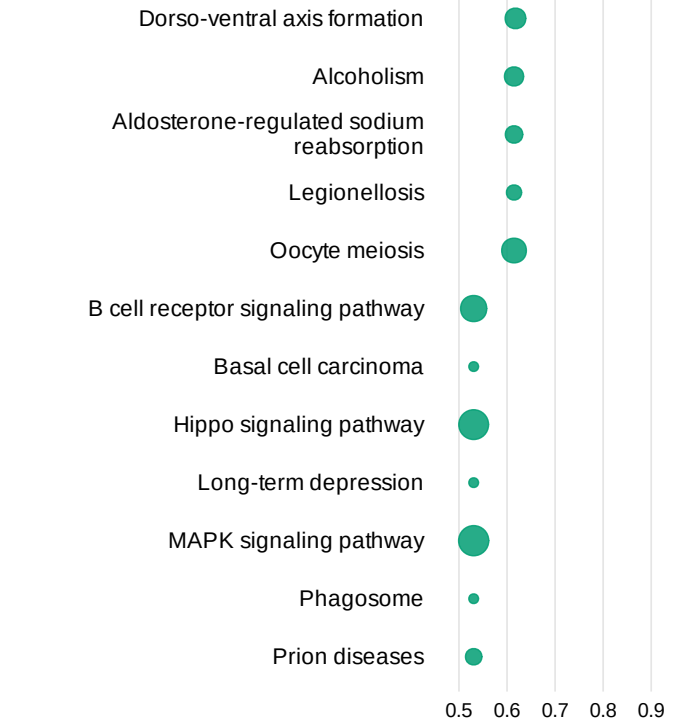
GO Biological
process
(miRPathDB)

GO Molecular
function
(miRPathDB)



KEGG
(miRPathDB)

Reactome
(miRPathDB)



Observed
10
20
30

-log10(P-adjusted)