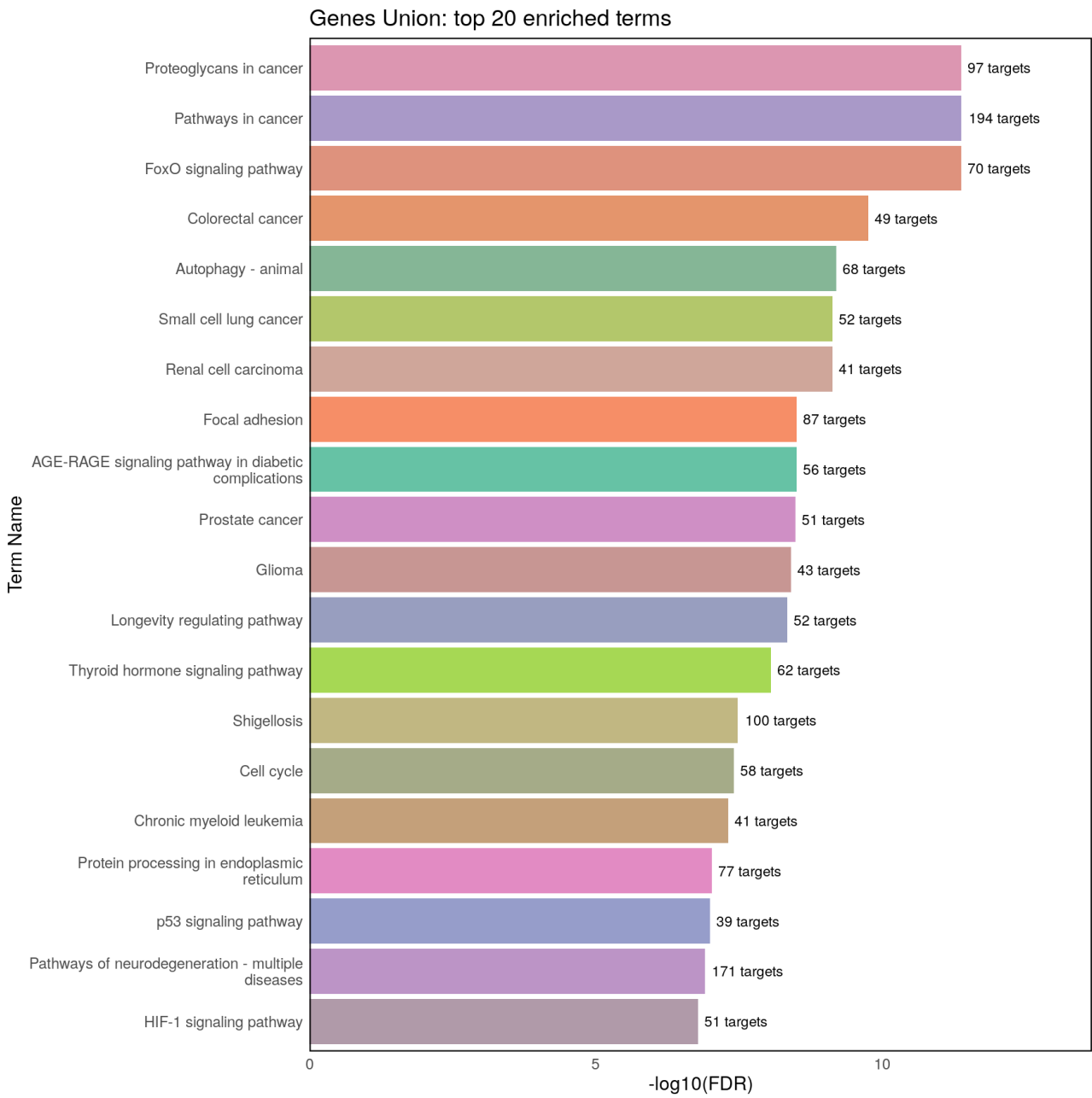


Supplementary File 4

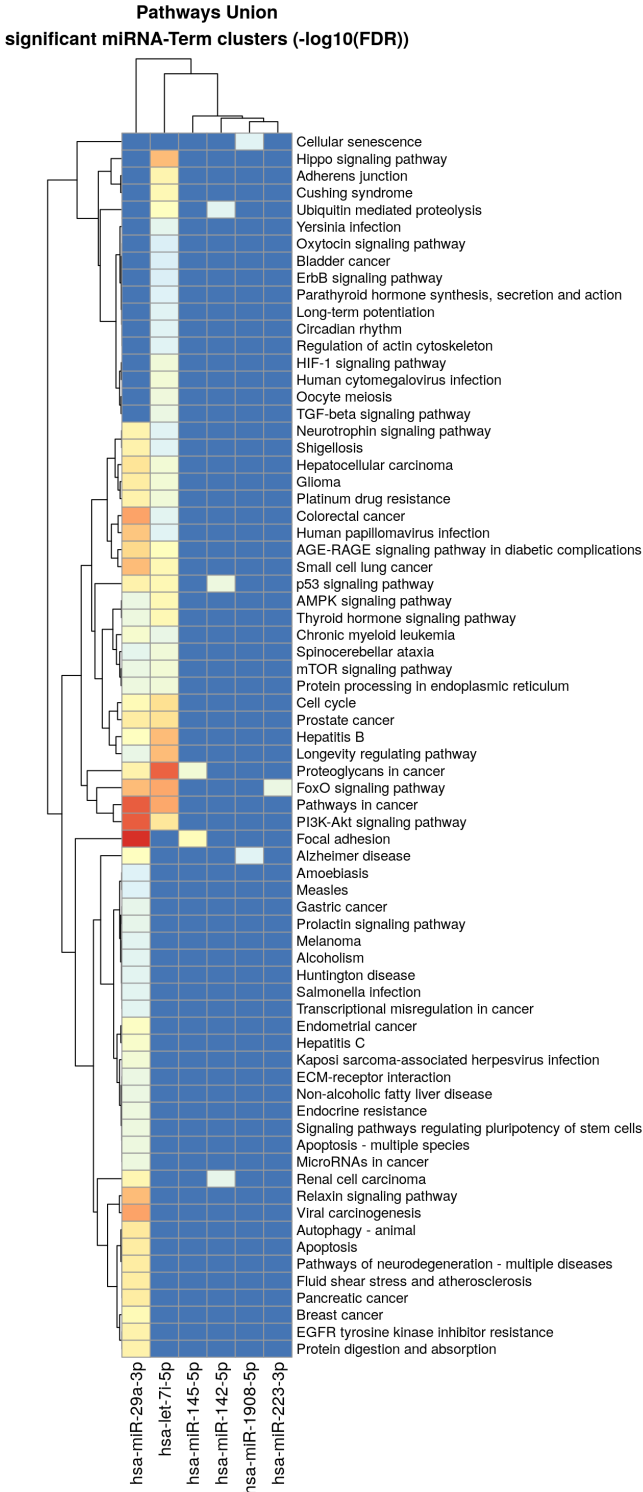
Functional enrichment analysis of pathways involved in the development of SARD-ILD and in disease subgroups

Whole cohort

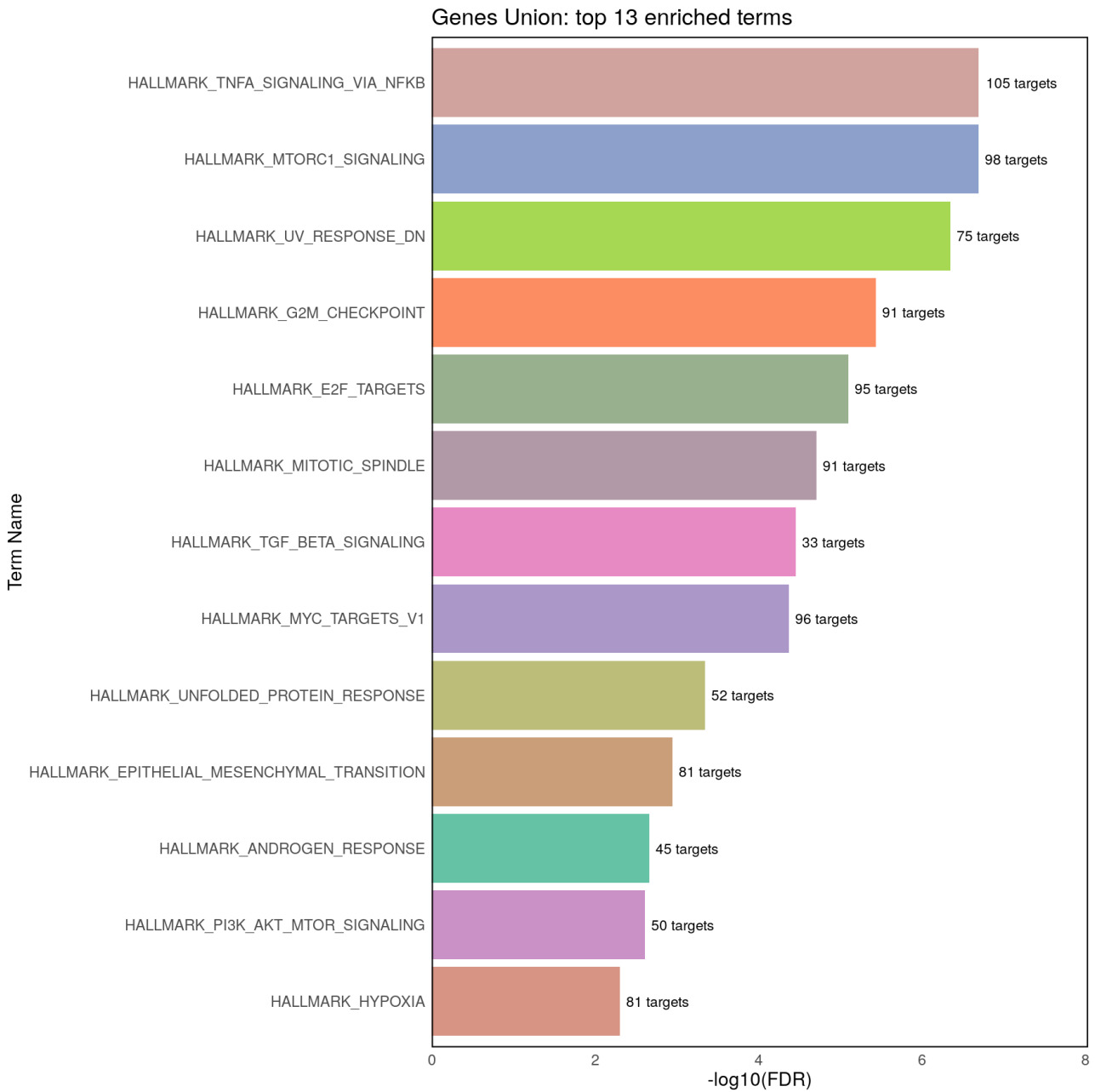
Genes union (KEGG, p 0.01)



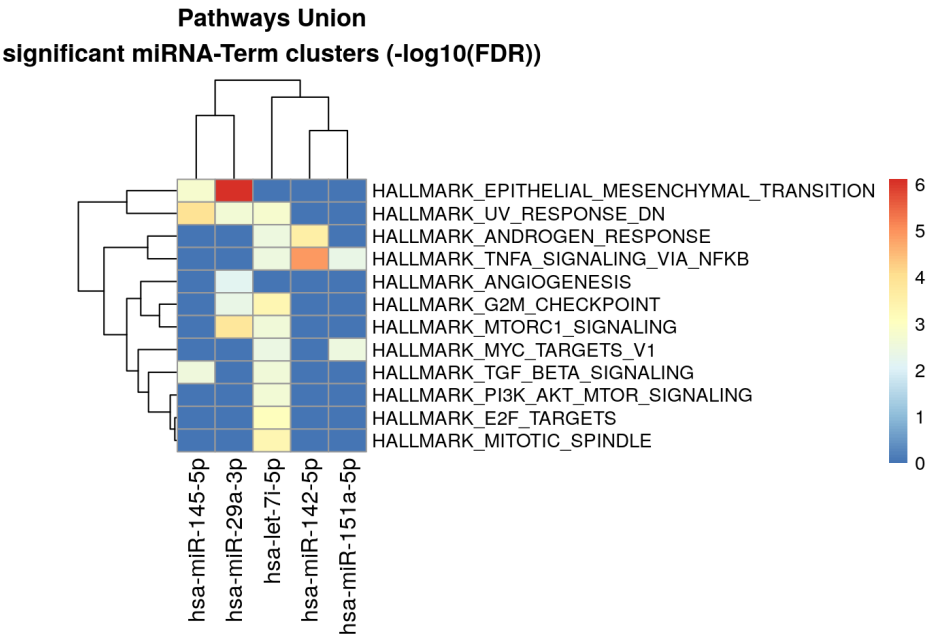
Whole cohort
Pathway union (KEGG, p 0.01)



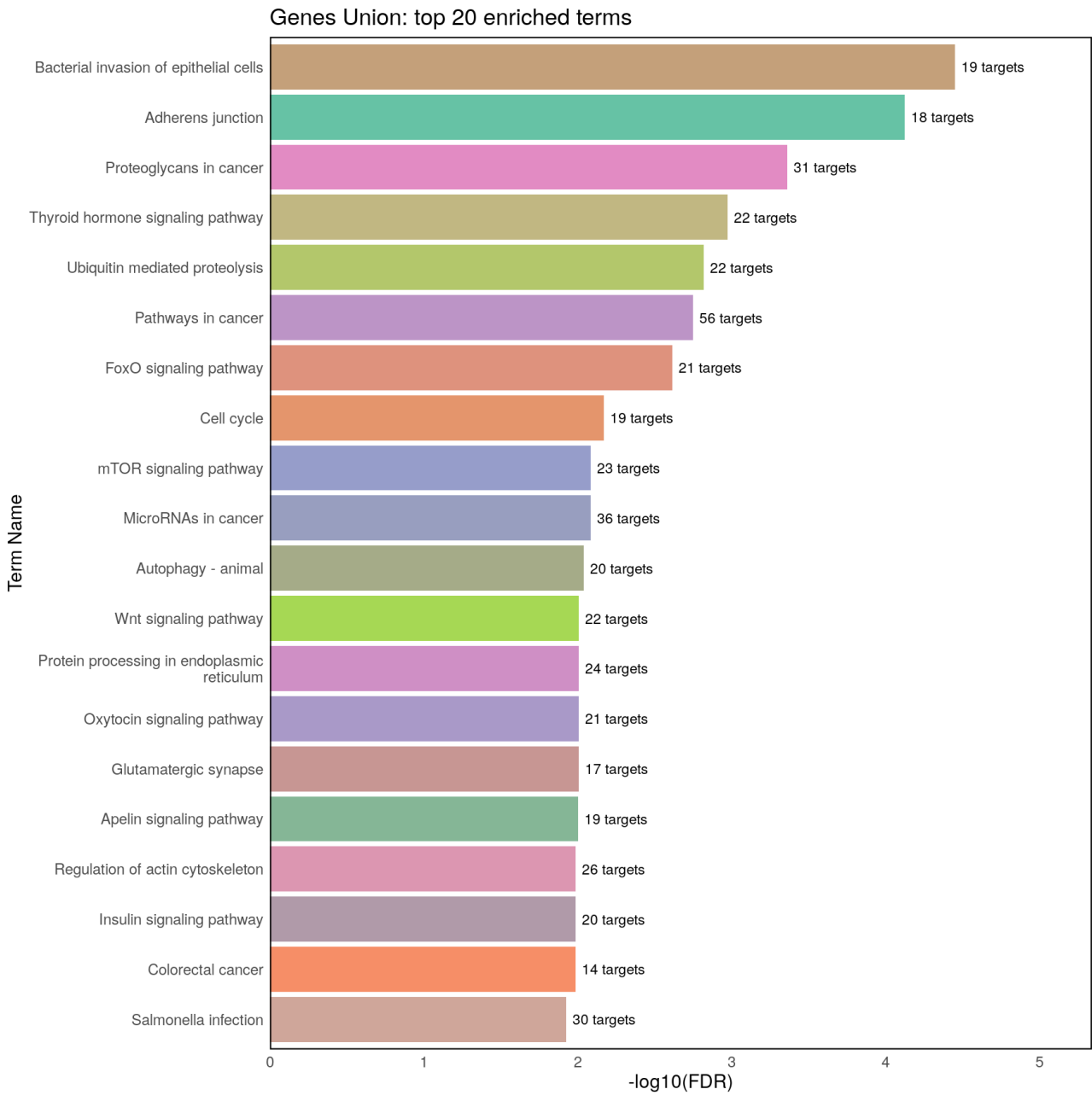
Whole cohort
Genes union (MSigDB p 0.01)



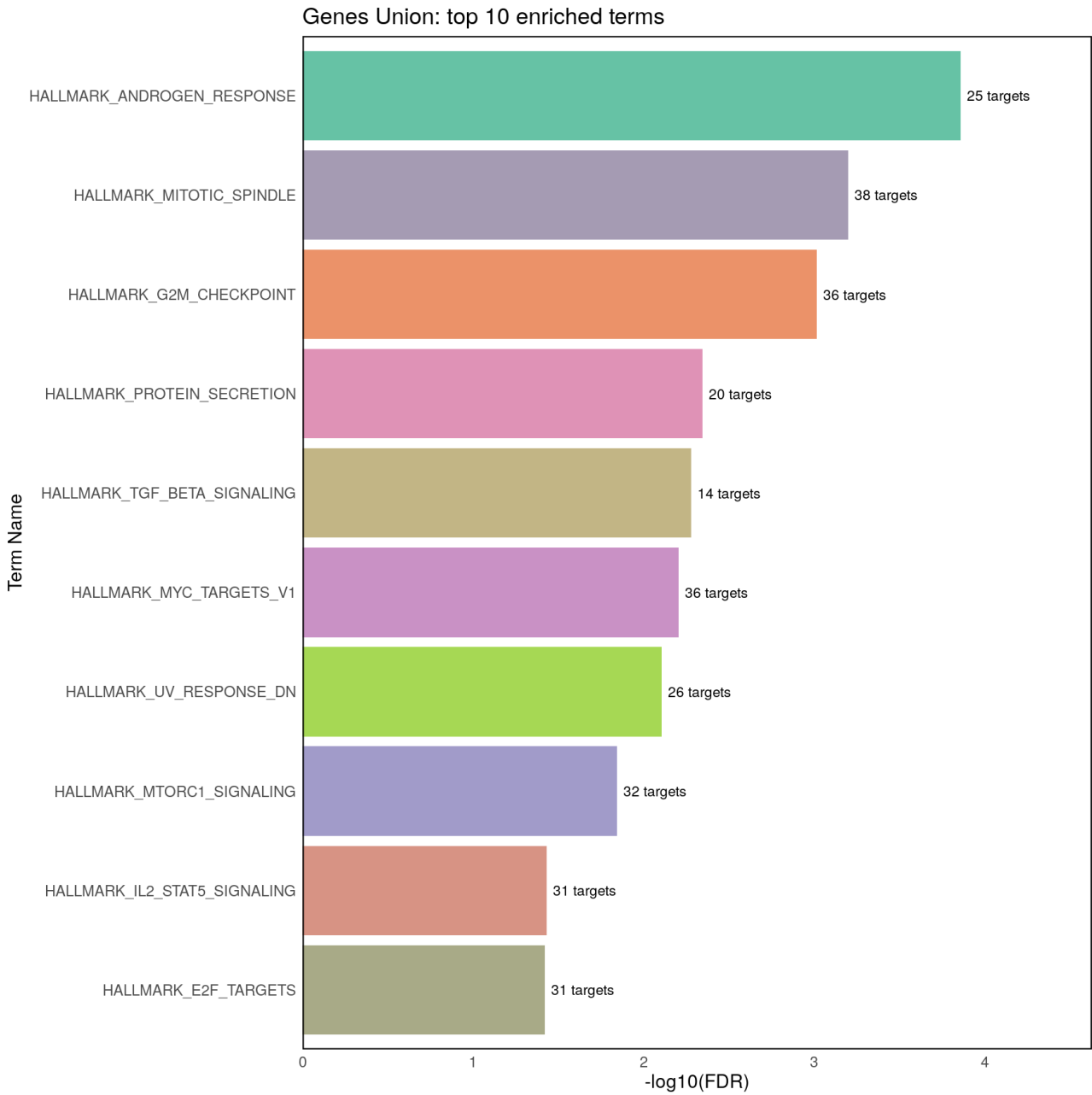
Whole cohort
Pathway union (MSigDB p 0.01)



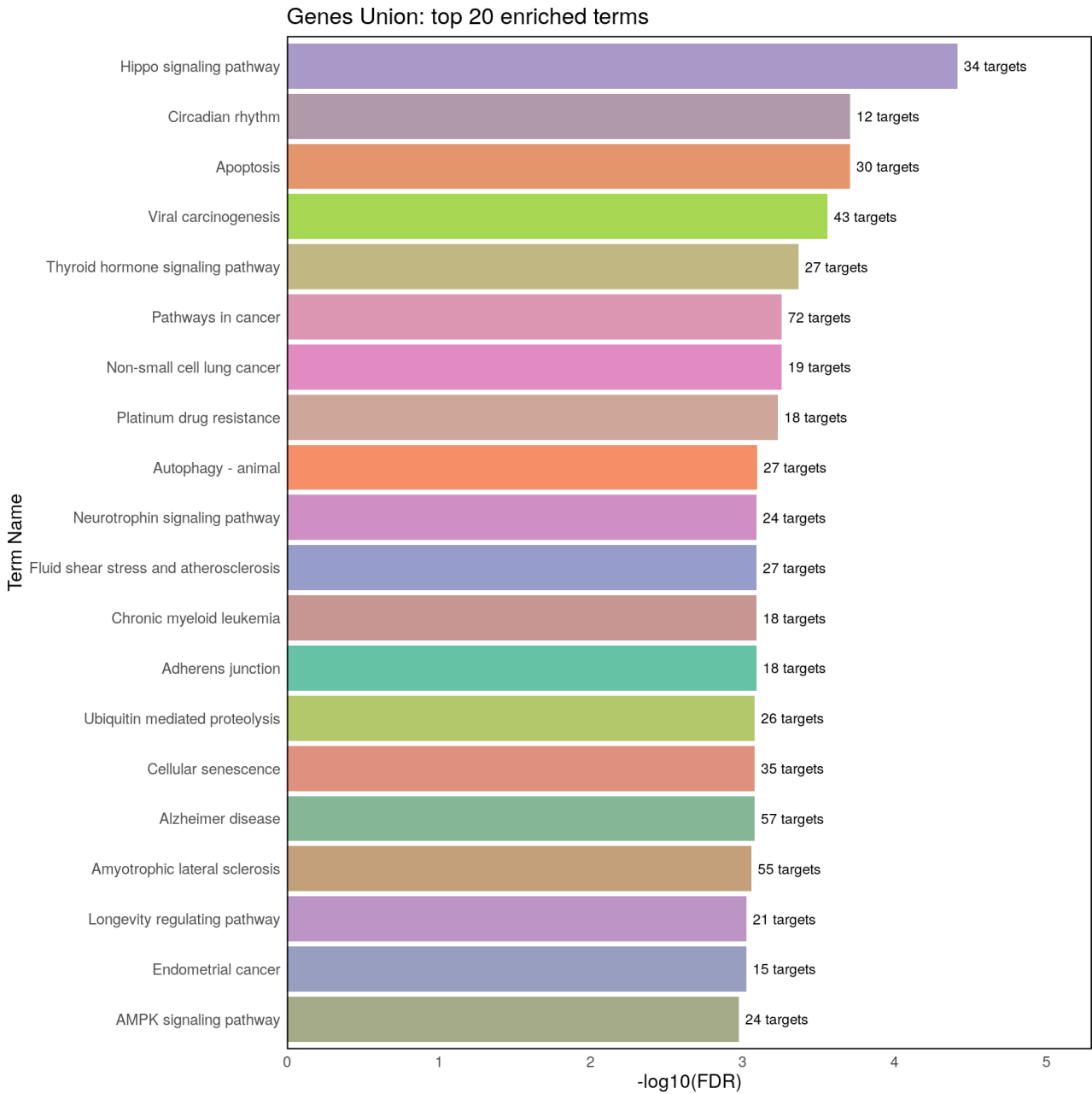
Early disease
Genes Union (KEGG, p 0.01)



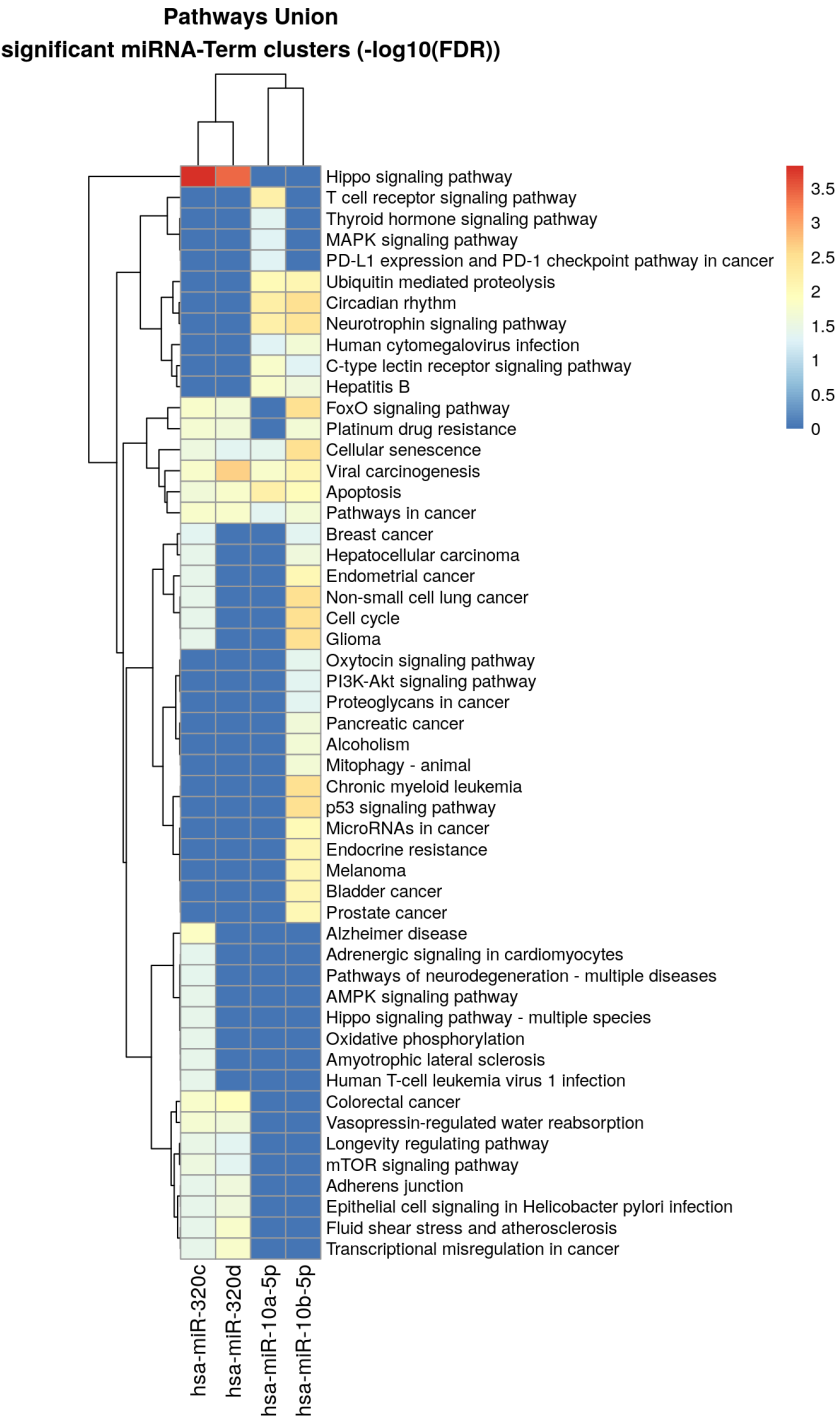
Early disease
Genes union (MSigDB, p 0.01)



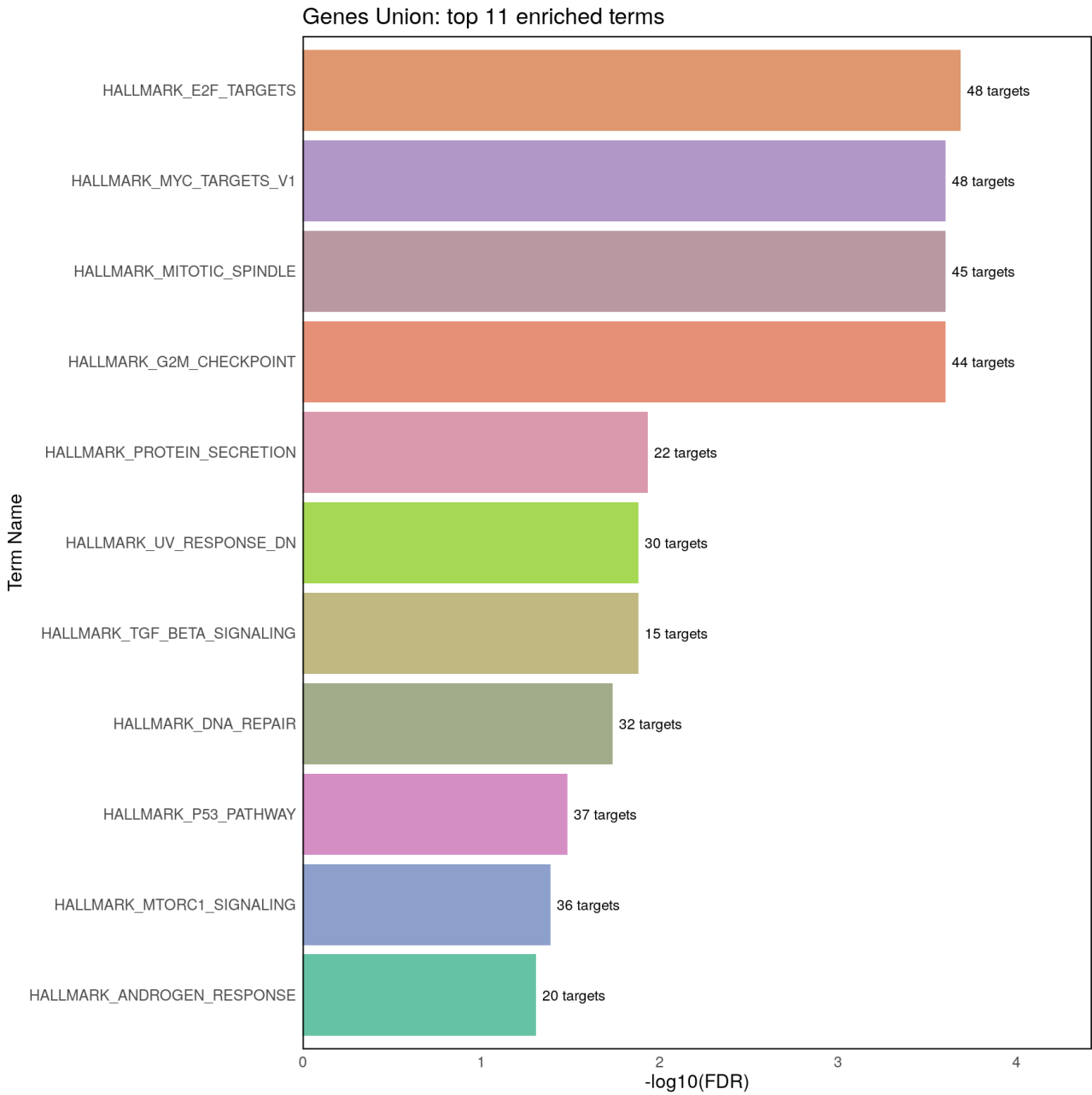
Vascular phenotype
Genes union (KEGG, p 0.01)



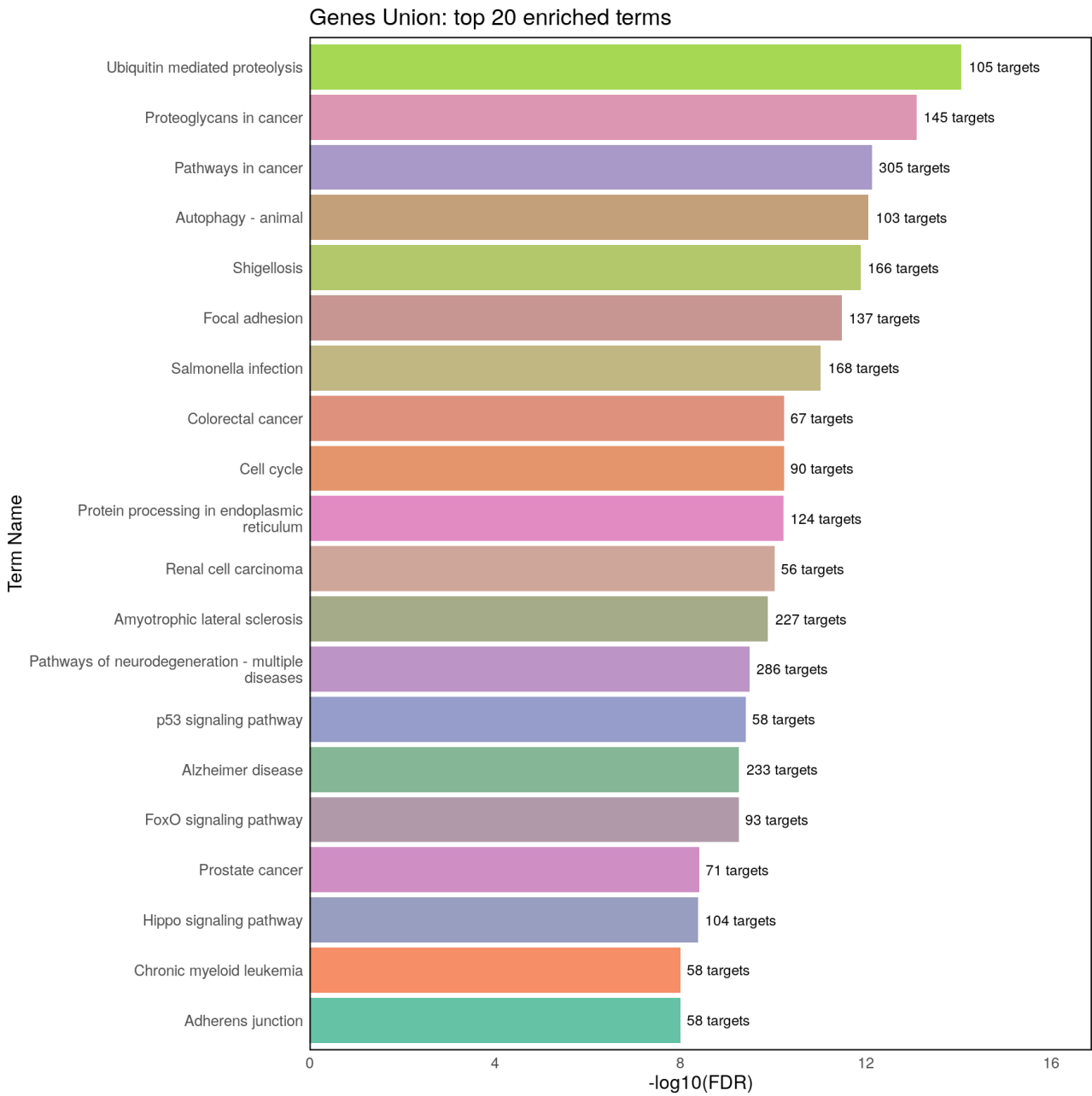
Vascular phenotype
Pathway union (KEGG, p 0.01)



Vascular phenotype
Genes union (MSigDB, p 0.01)



Progressing cases
Genes union (KEGG, p 0.01)



Progressing cases

(Pathways union, KEGG p 0.01)



Progressing cases
Genes union (MSigDB, p 0.01)

