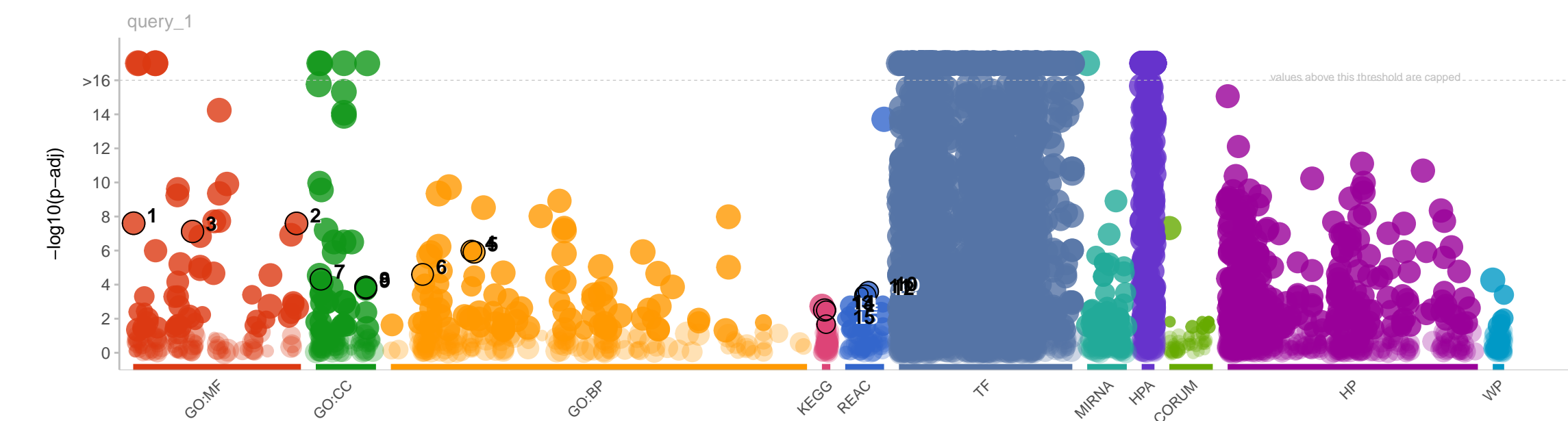
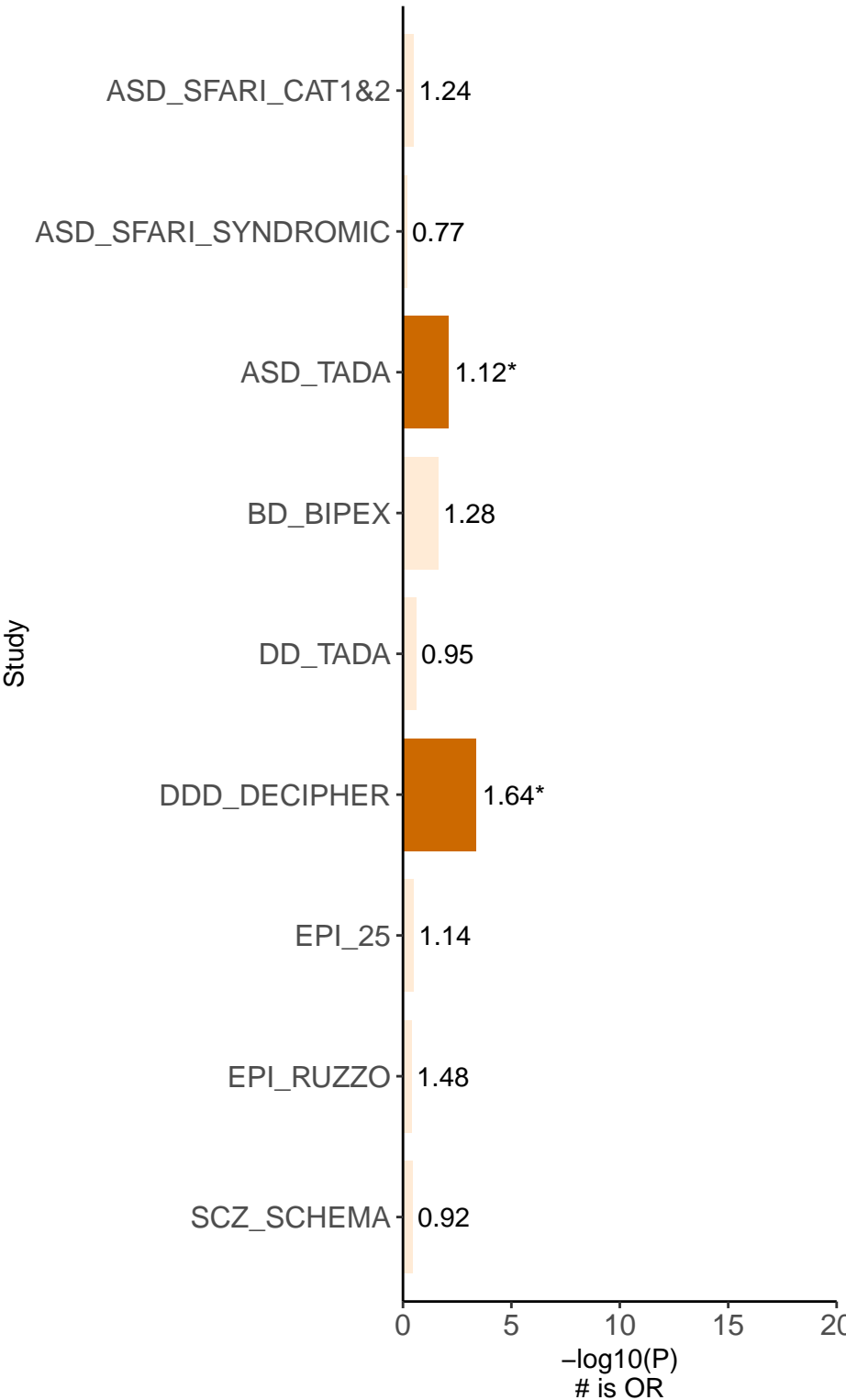
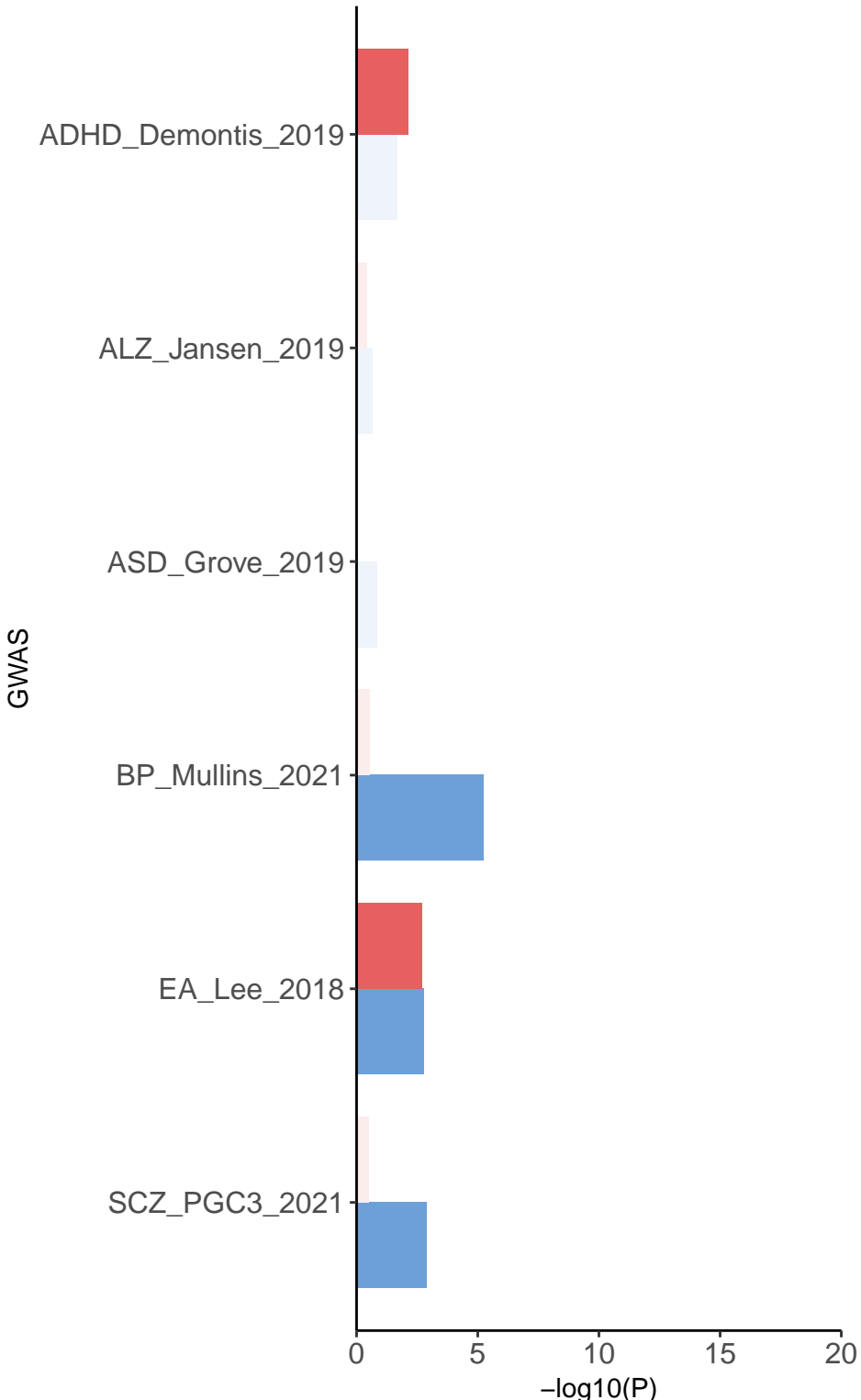
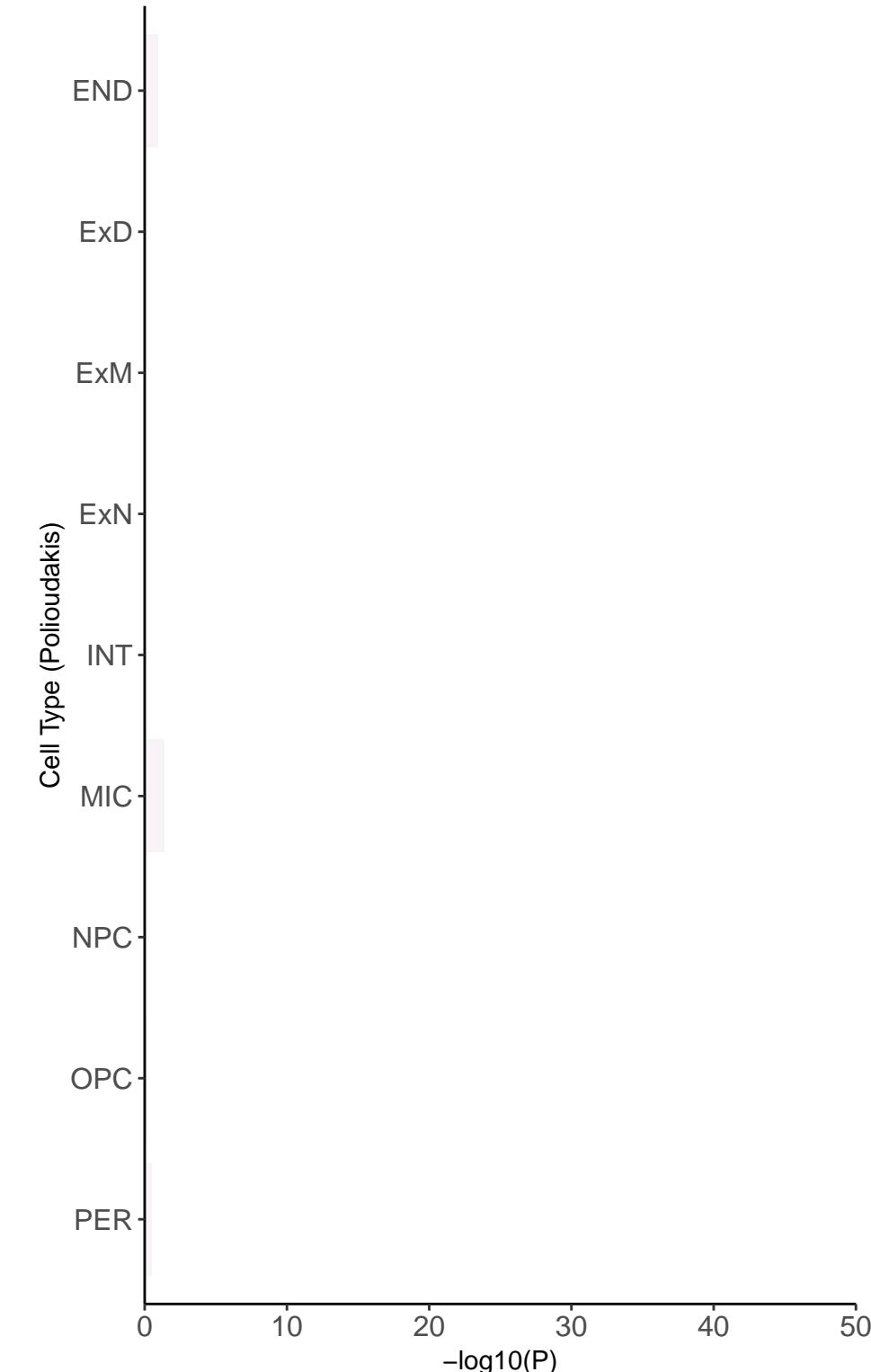
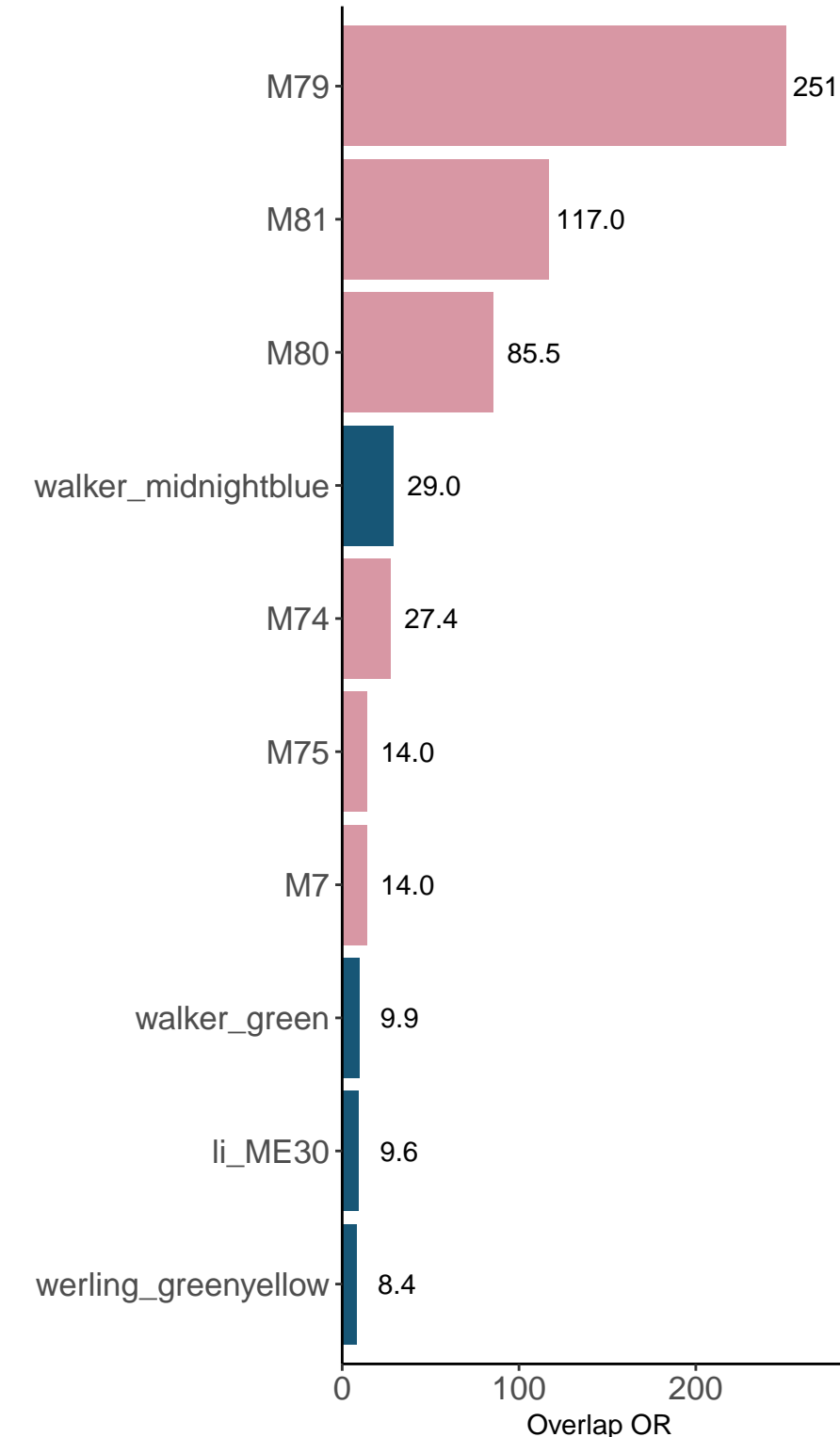
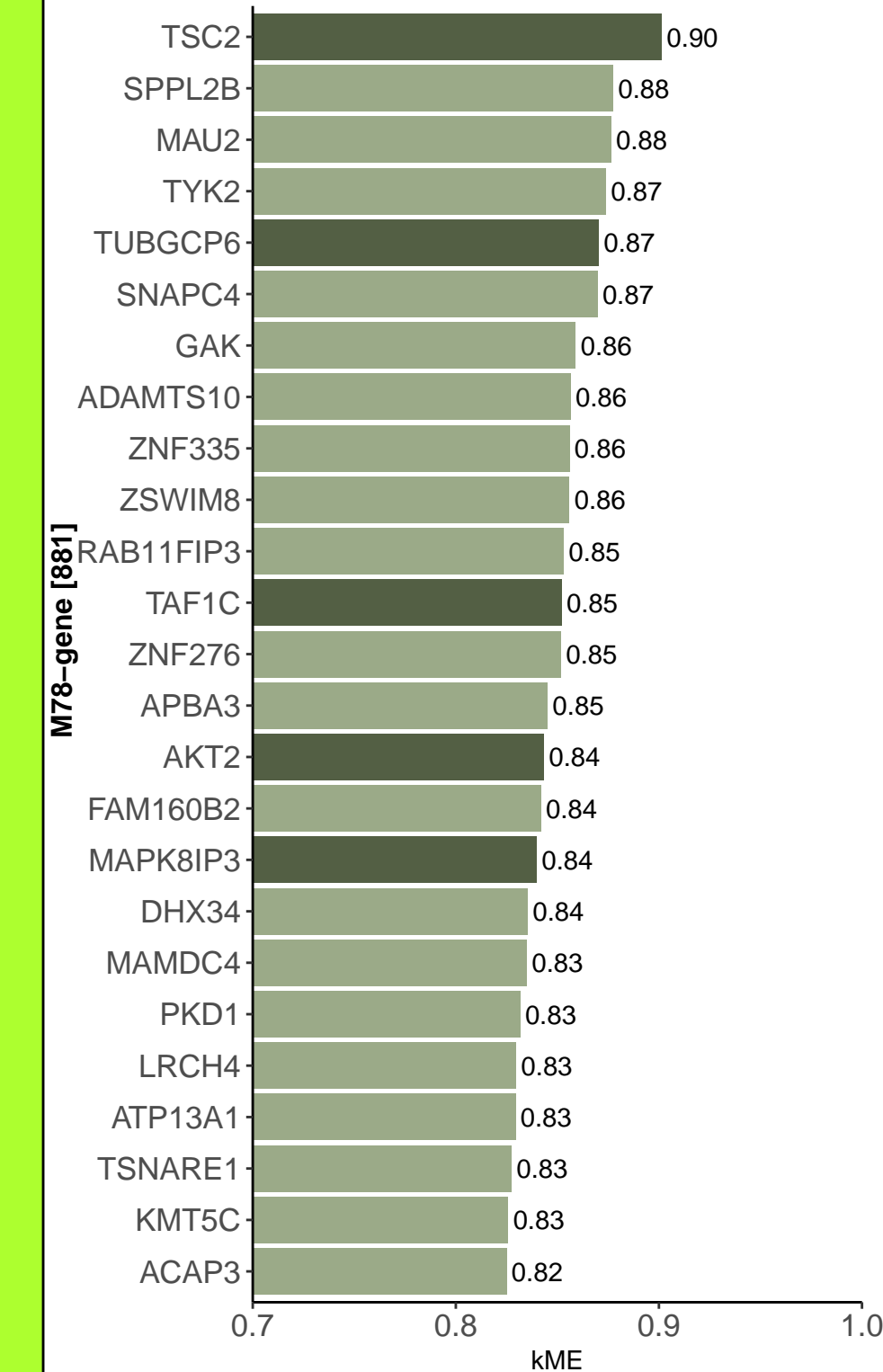


M78-gene [881]



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0000166	nucleotide binding	1973	2.5e-08
2	GO:MF	GO:1901265	nucleoside phosphate binding	1974	2.5e-08
3	GO:MF	GO:0030554	adenyl nucleotide binding	1523	7.5e-08
4	GO:BP	GO:0016570	histone modification	434	1.0e-06
5	GO:BP	GO:0018193	peptidyl-amino acid modification	1120	1.2e-06
6	GO:BP	GO:0006259	DNA metabolic process	973	2.5e-05
7	GO:CC	GO:0005773	vacuole	764	4.9e-05
8	GO:CC	GO:009080	supramolecular complex	1150	1.4e-04
9	GO:CC	GO:0099512	supramolecular fiber	812	1.6e-04
10	REAC	REAC:R-HSA-1852241	Organelle biogenesis and maintenance	270	2.6e-04
11	REAC	REAC:R-HSA-165181	Inhibition of TSC complex formation by PKB	3	3.4e-04
12	REAC	REAC:R-HSA-556833	Metabolism of lipids	648	4.4e-04
13	KEGG	KEGG:01100	Metabolic pathways	1334	3.1e-03
14	KEGG	KEGG:04142	Lysosome	122	3.3e-03
15	KEGG	KEGG:04211	Longevity regulating pathway	83	2.1e-02

[g:Profiler \(biit.cs.ut.ee/gprofiler\)](https://biit.cs.ut.ee/gprofiler)