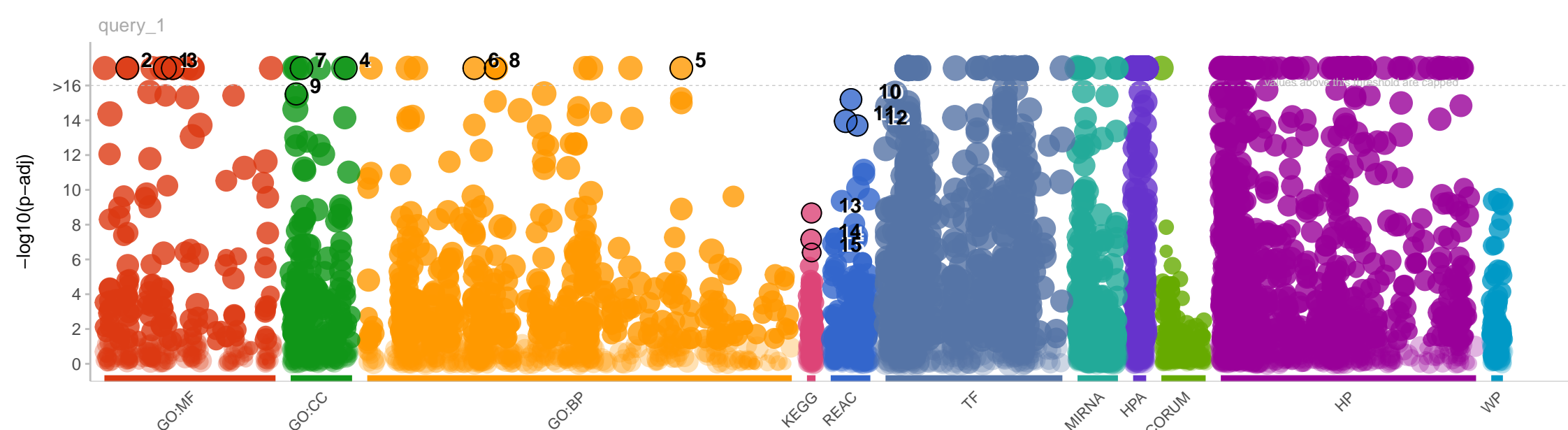
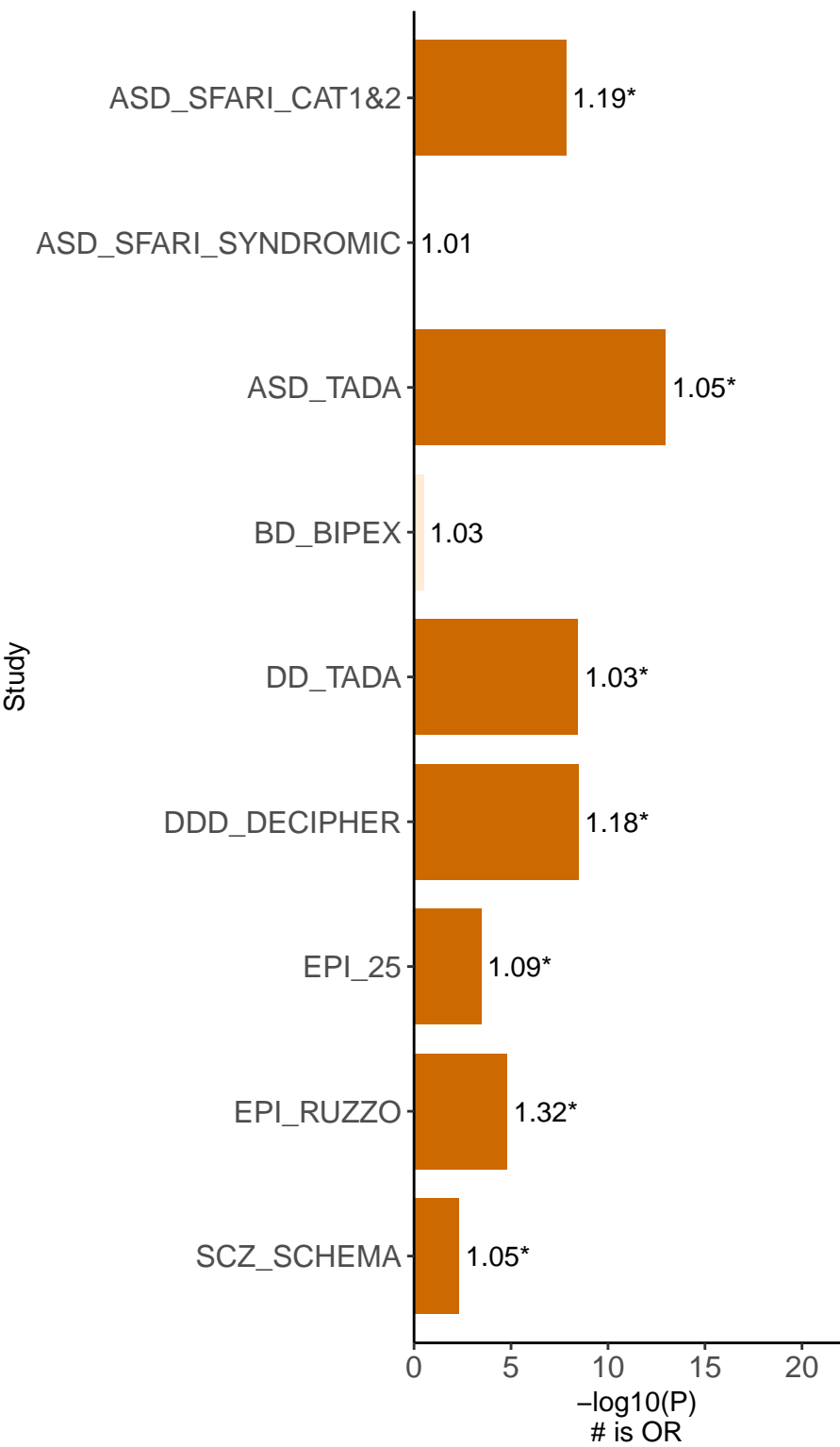
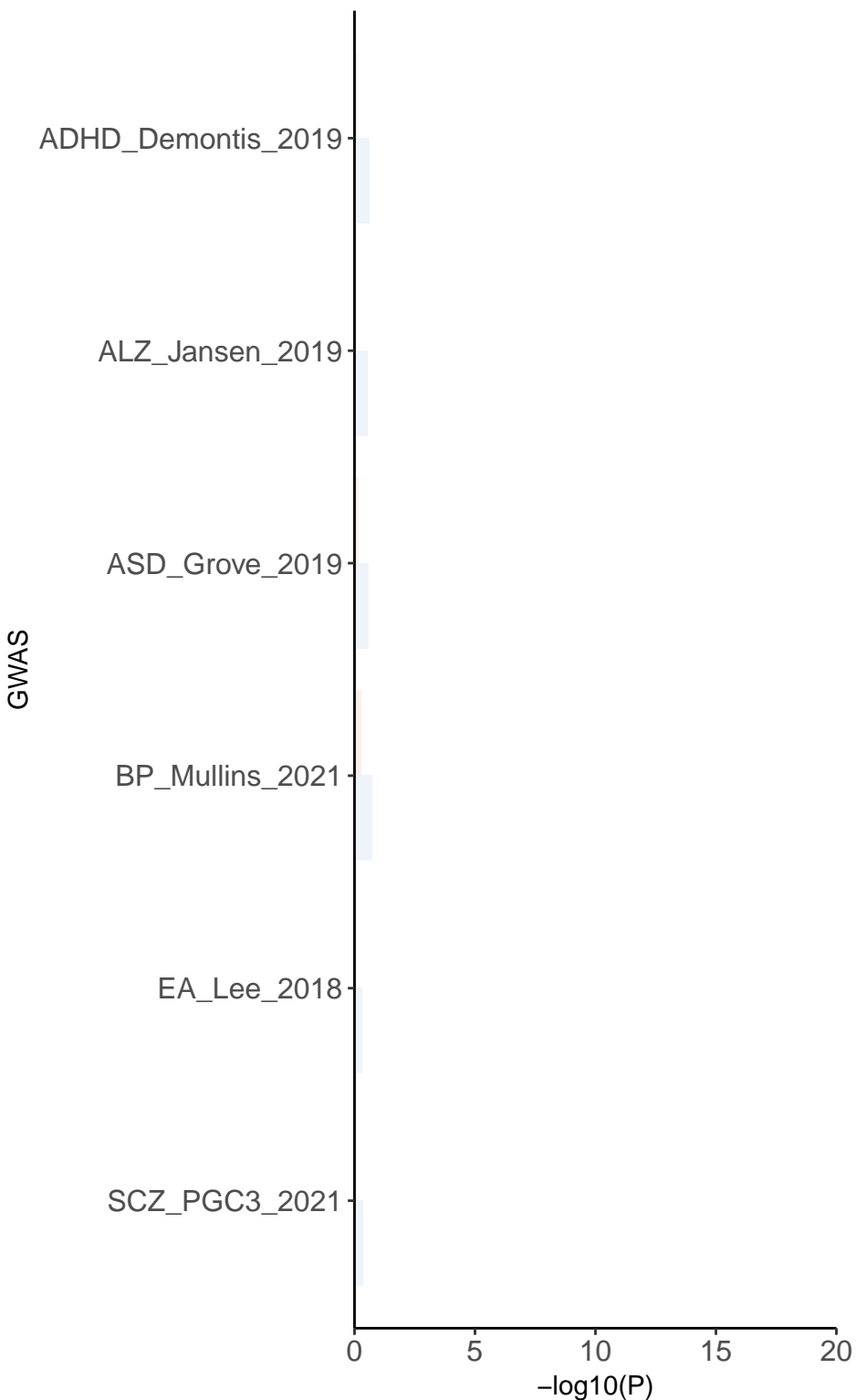
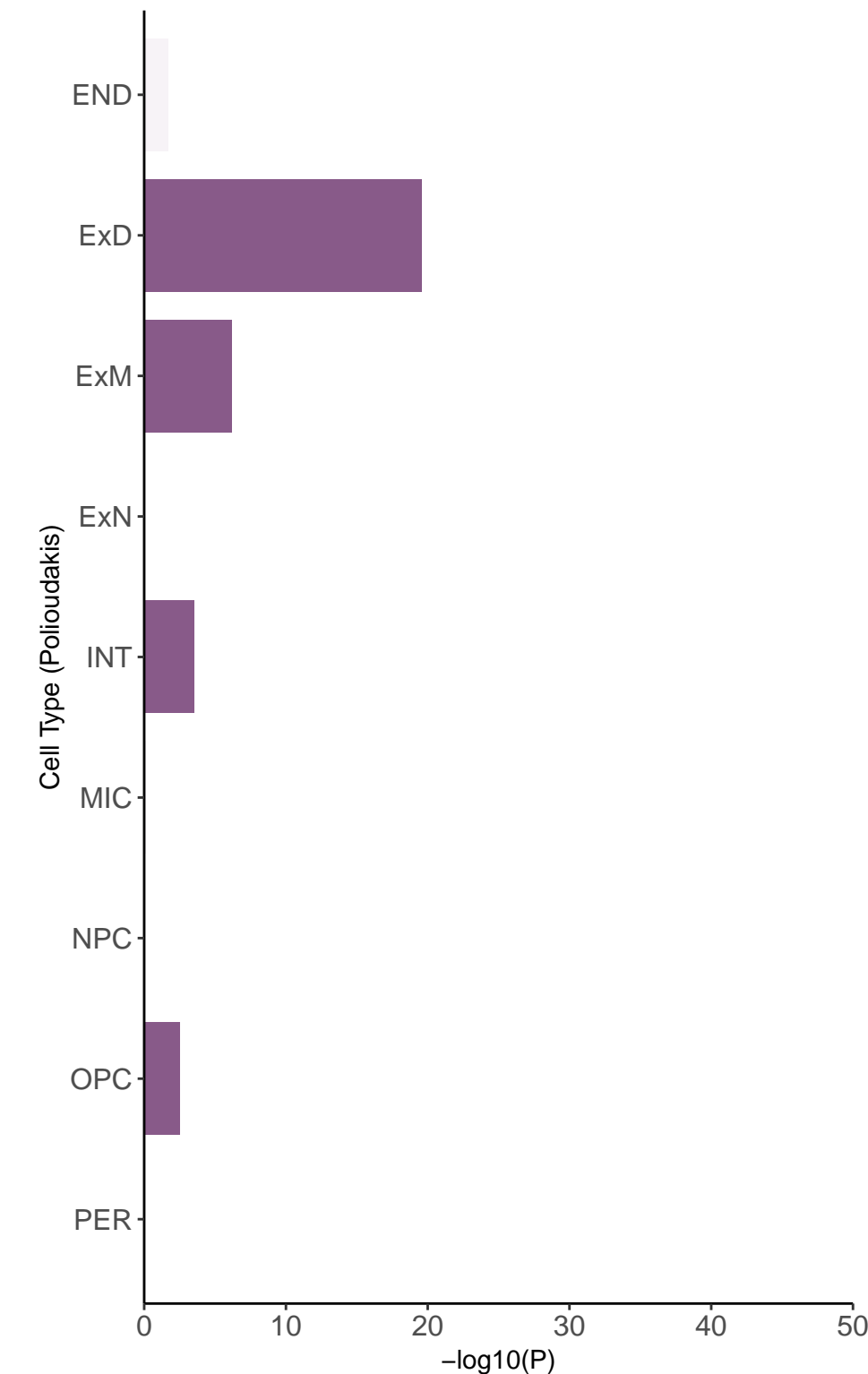
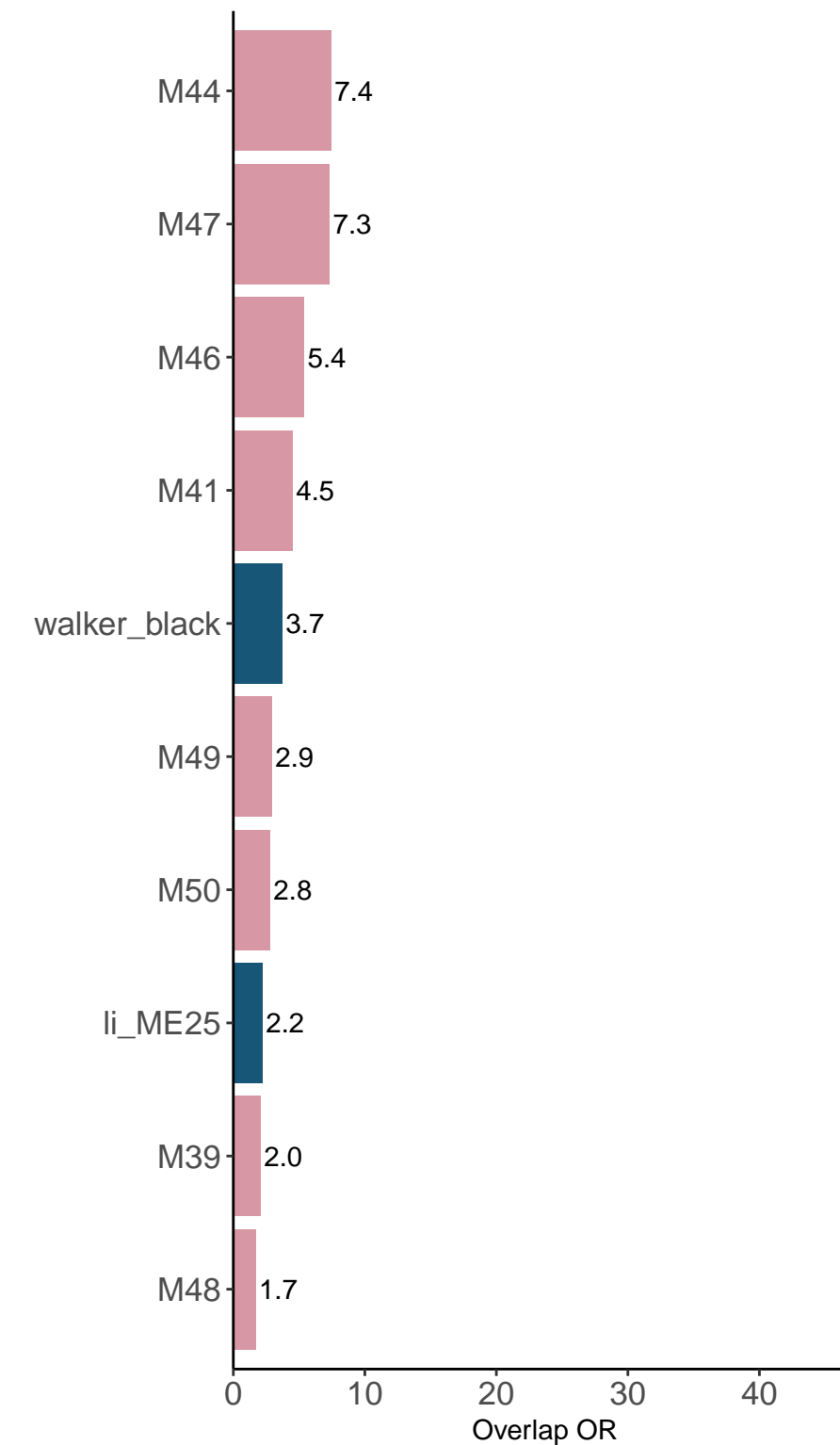
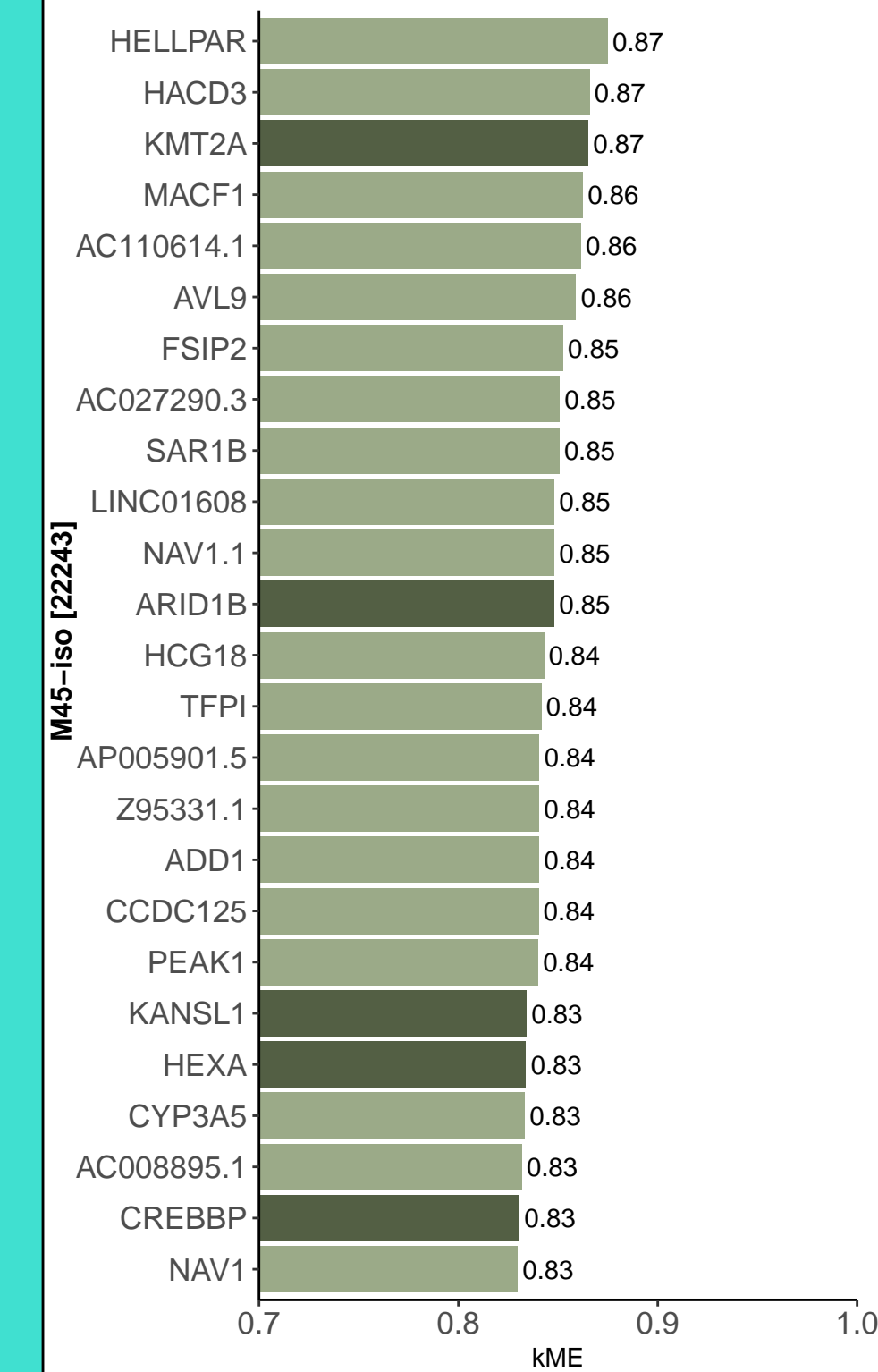


M45-iso [22243]



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0030554	adenyl nucleotide binding	1523	2.9e-26
2	GO:MF	GO:0005524	ATP binding	1369	2.9e-26
3	GO:MF	GO:0032559	adenyl ribonucleotide binding	1428	6.7e-26
4	GO:CC	GO:1902494	catalytic complex	1628	2.6e-23
5	GO:BP	GO:0120036	plasma membrane bounded cell projection organization	1399	3.5e-23
6	GO:BP	GO:0030030	cell projection organization	1436	4.2e-22
7	GO:CC	GO:0015630	microtubule cytoskeleton	1271	2.0e-19
8	GO:BP	GO:0032989	cellular component morphogenesis	713	5.6e-17
9	GO:CC	GO:0005815	microtubule organizing center	787	3.1e-16
10	REAC	REAC:R-HSA-199991	Membrane Trafficking	578	6.4e-16
11	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	1.1e-14
12	REAC	REAC:R-HSA-9012999	RHO GTPase cycle	419	2.0e-14
13	KEGG	KEGG:04360	Axon guidance	174	2.1e-09
14	KEGG	KEGG:04144	Endocytosis	238	7.1e-08
15	KEGG	KEGG:04520	Adherens junction	70	4.0e-07

g:Profiler (biit.cs.ut.ee/gprofiler)