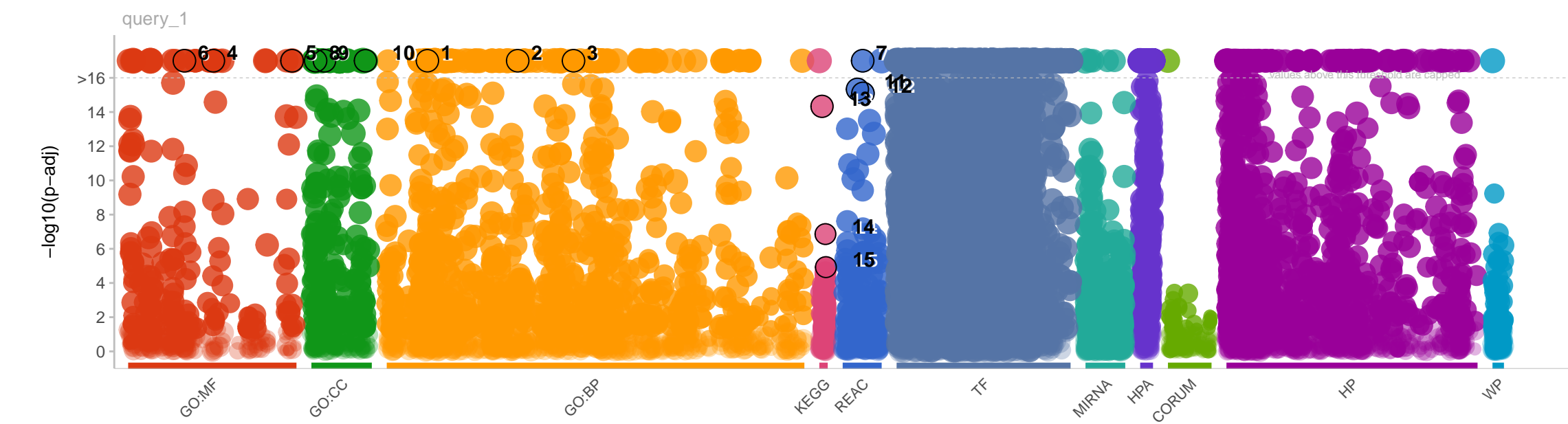
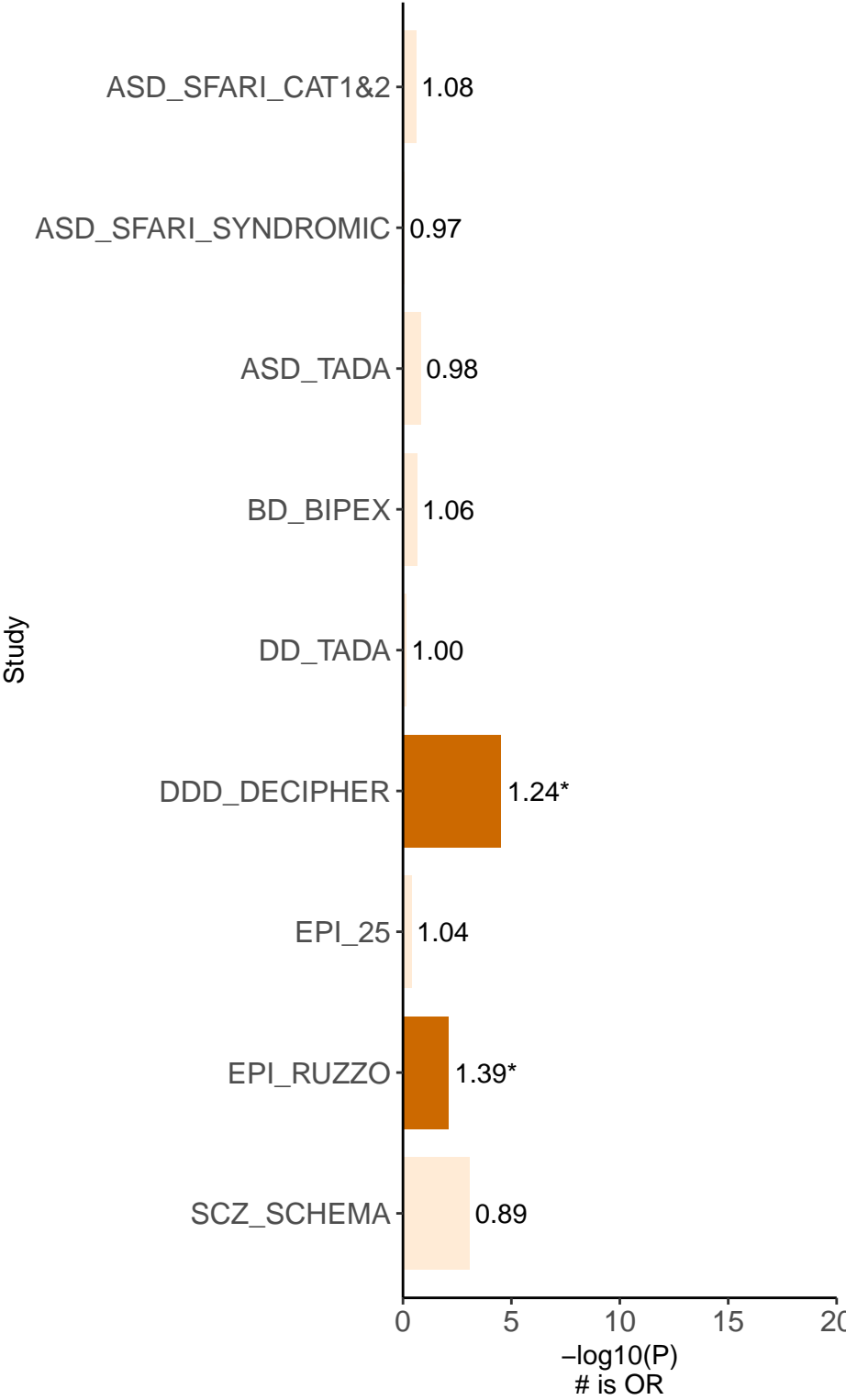
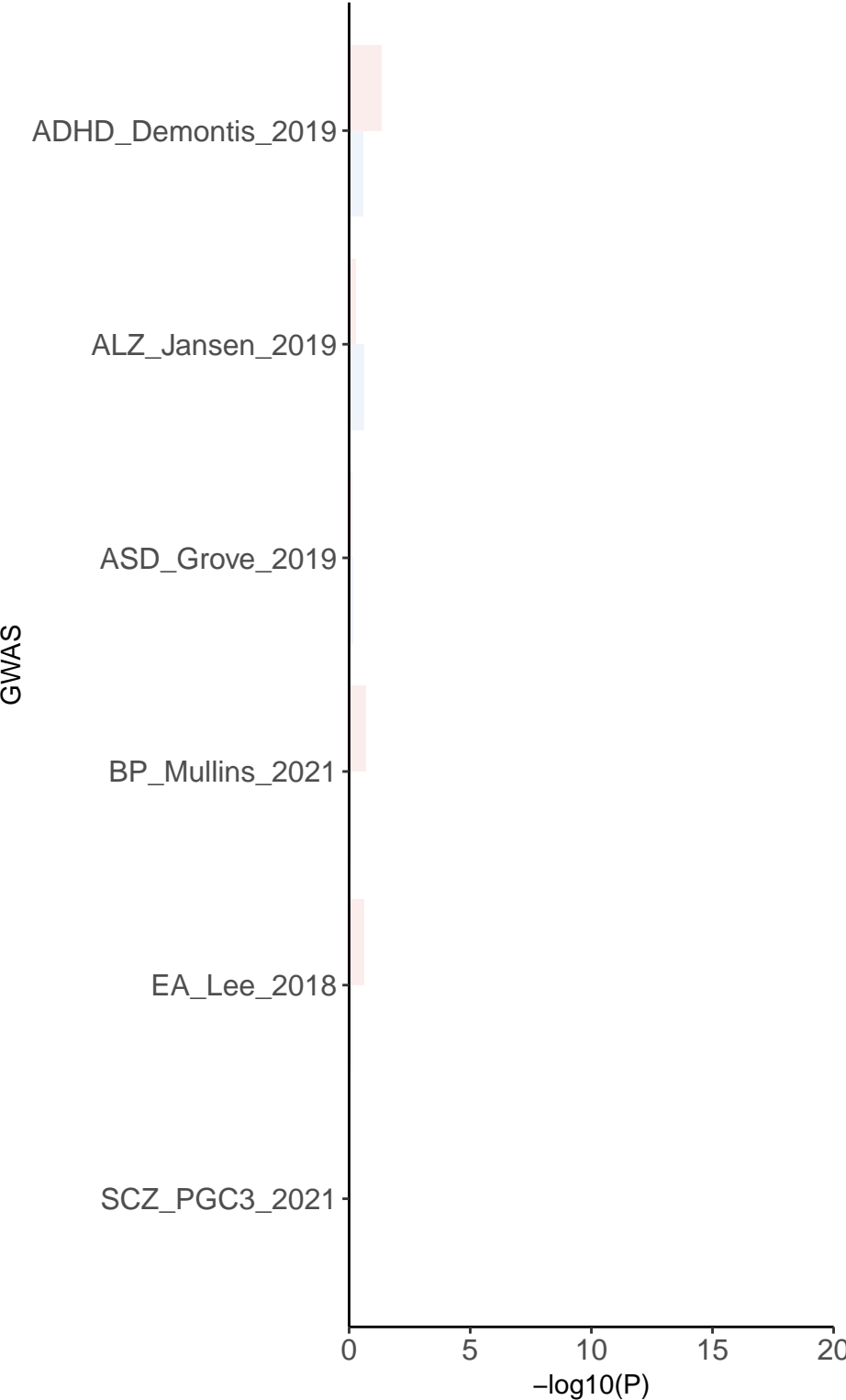
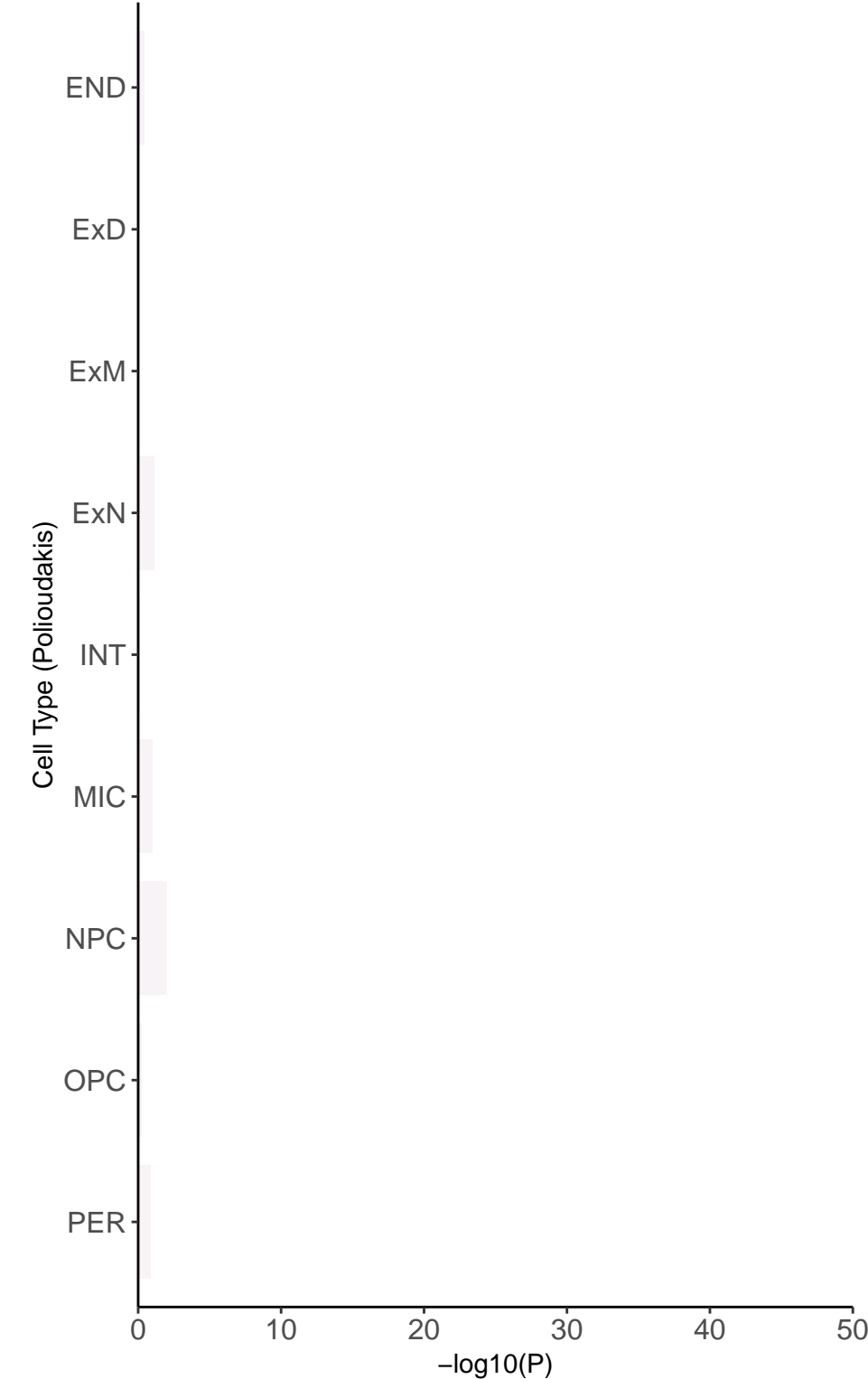
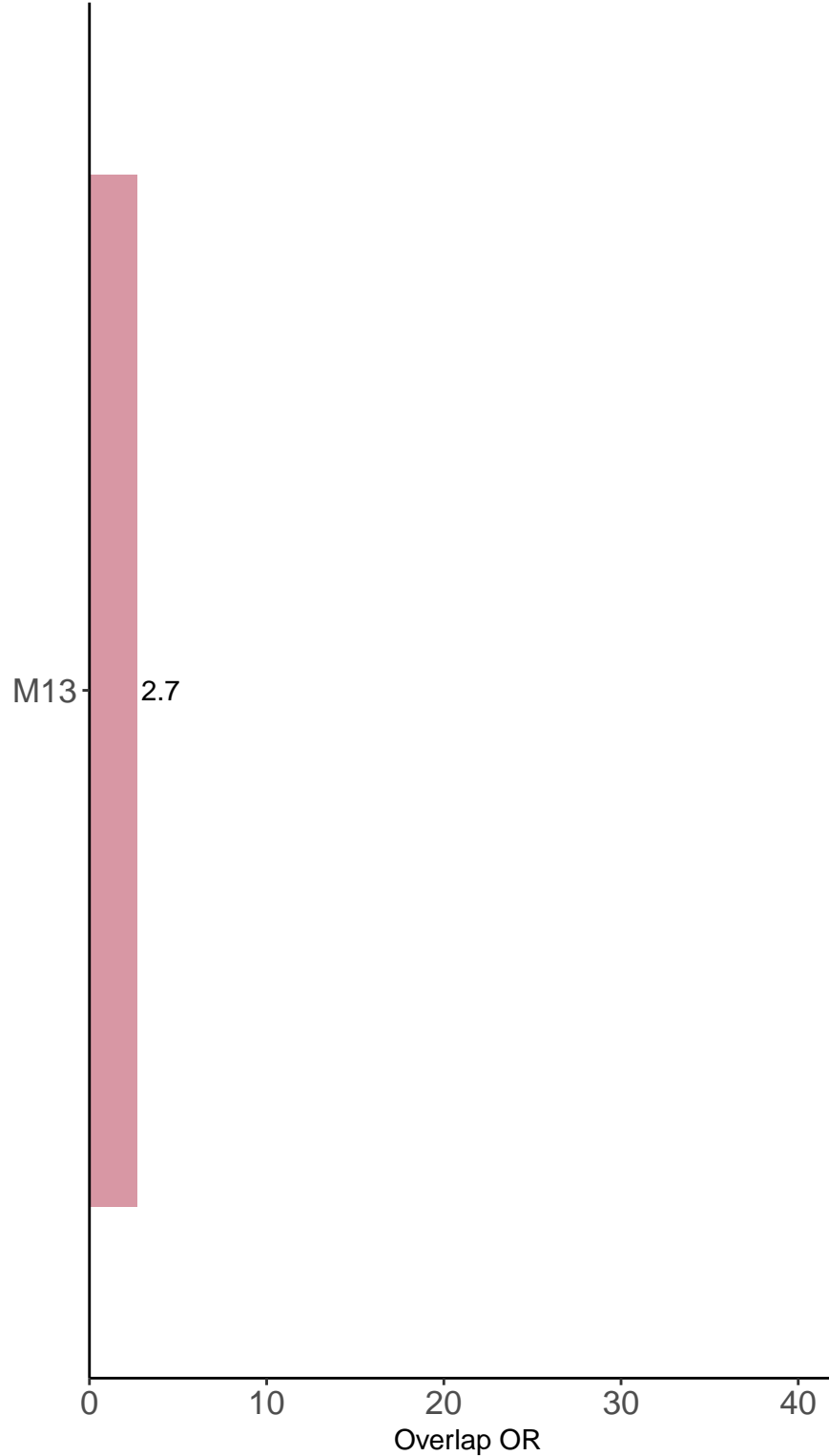
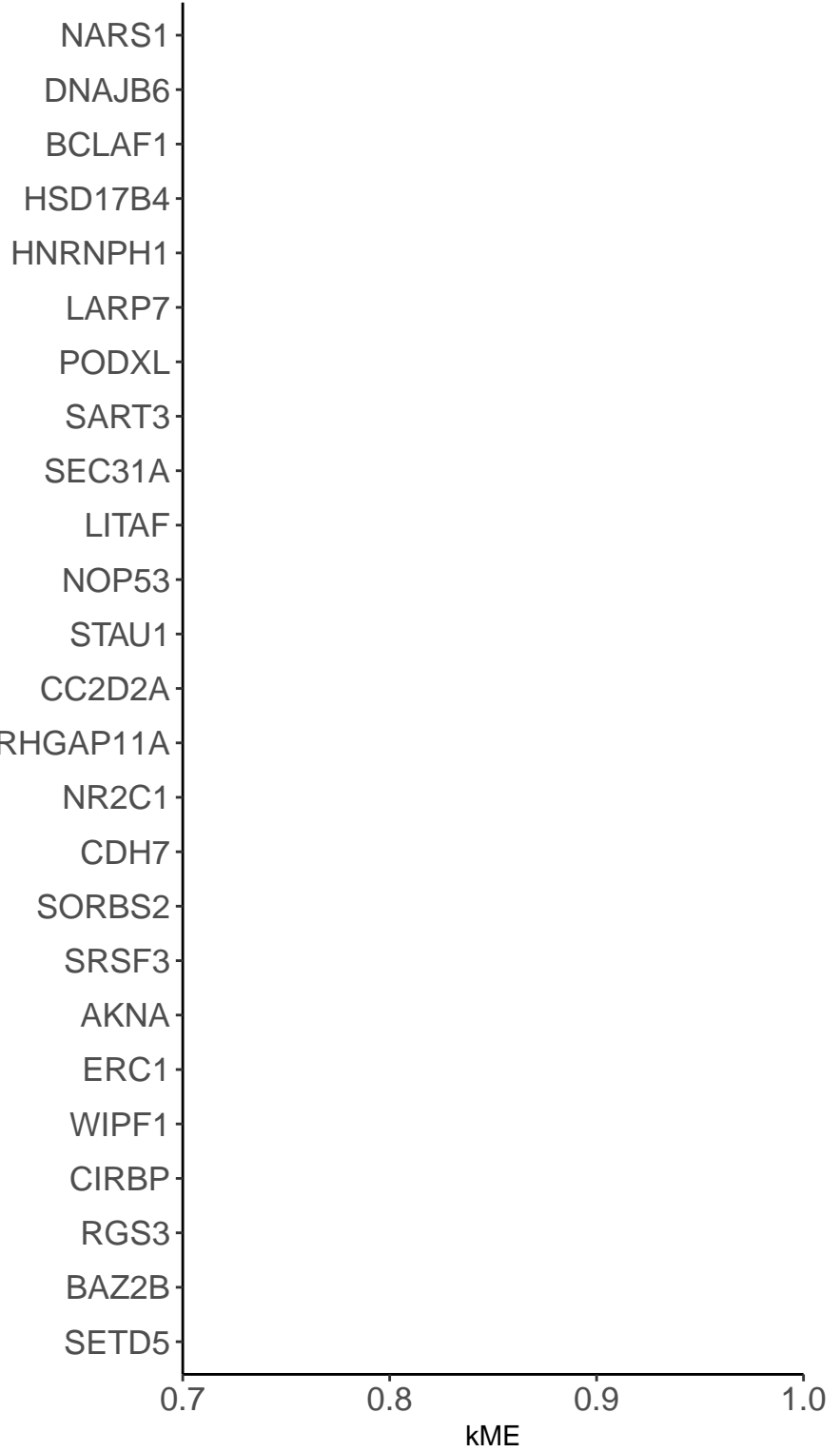


M10-Iso [4686]



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0007049	cell cycle	1655	1.6e-37
2	GO:BP	GO:0033554	cellular response to stress	1790	9.9e-34
3	GO:BP	GO:0045935	positive regulation of nucleobase-containing compound metabolic process	1798	7.8e-29
4	GO:MF	GO:0042802	identical protein binding	1838	1.5e-27
5	GO:MF	GO:1901265	nucleoside phosphate binding	1974	3.6e-27
6	GO:MF	GO:0019899	enzyme binding	1888	4.7e-27
7	REAC	REAC:R-HSA-1430728	Metabolism	1832	2.1e-25
8	GO:CC	GO:0005694	chromosome	1638	6.5e-25
9	GO:CC	GO:0030054	cell junction	1916	1.1e-21
10	GO:CC	GO:1902494	catalytic complex	1628	2.8e-19
11	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	4.7e-16
12	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	8.1e-16
13	KEGG	KEGG:01100	Metabolic pathways	1334	4.7e-15
14	KEGG	KEGG:05022	Pathways of neurodegeneration – multiple diseases	421	1.4e-07
15	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	1.2e-05

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