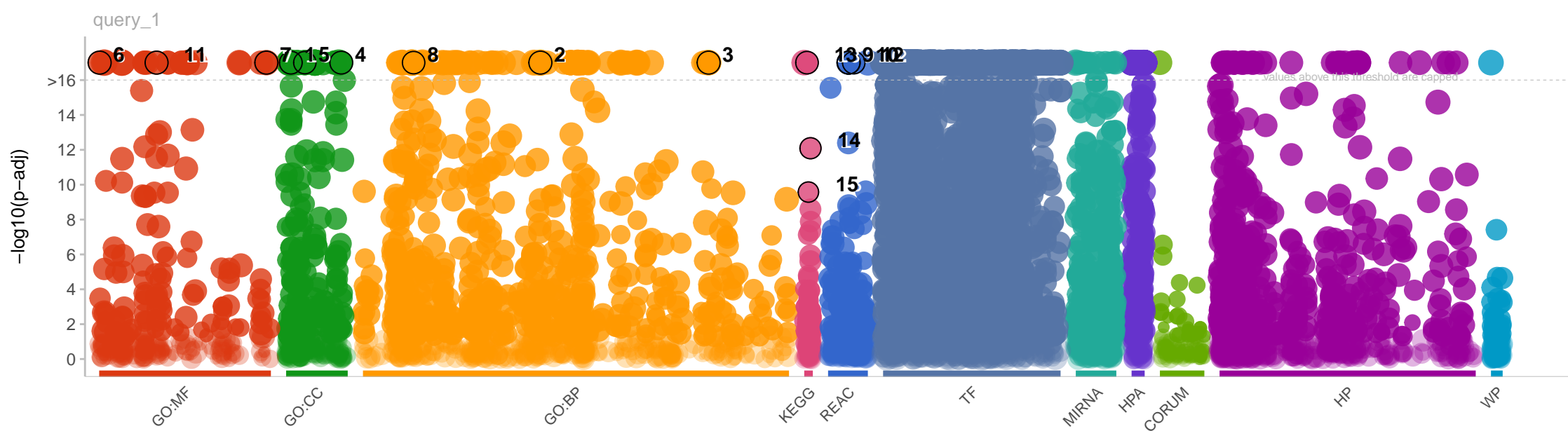
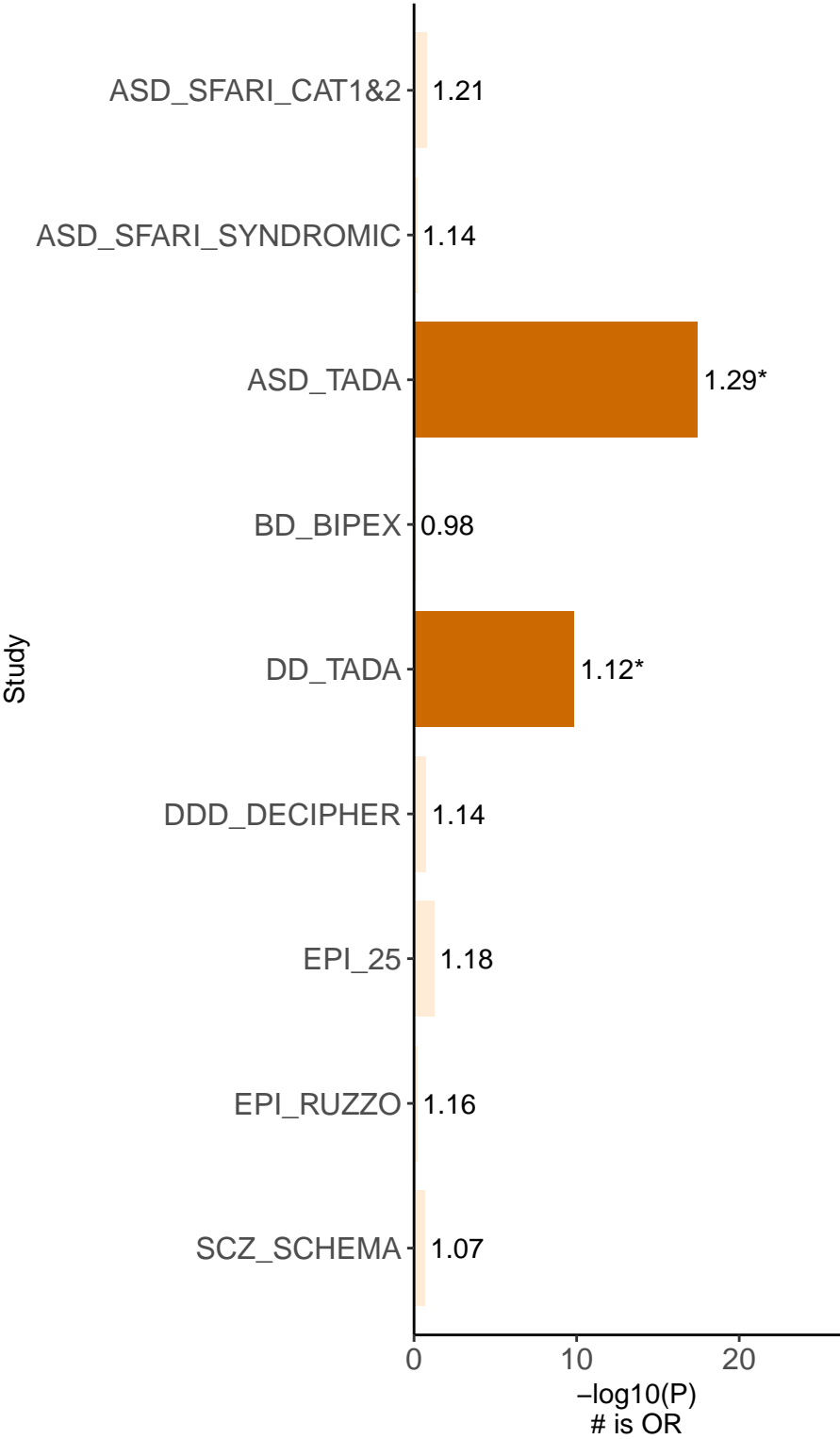
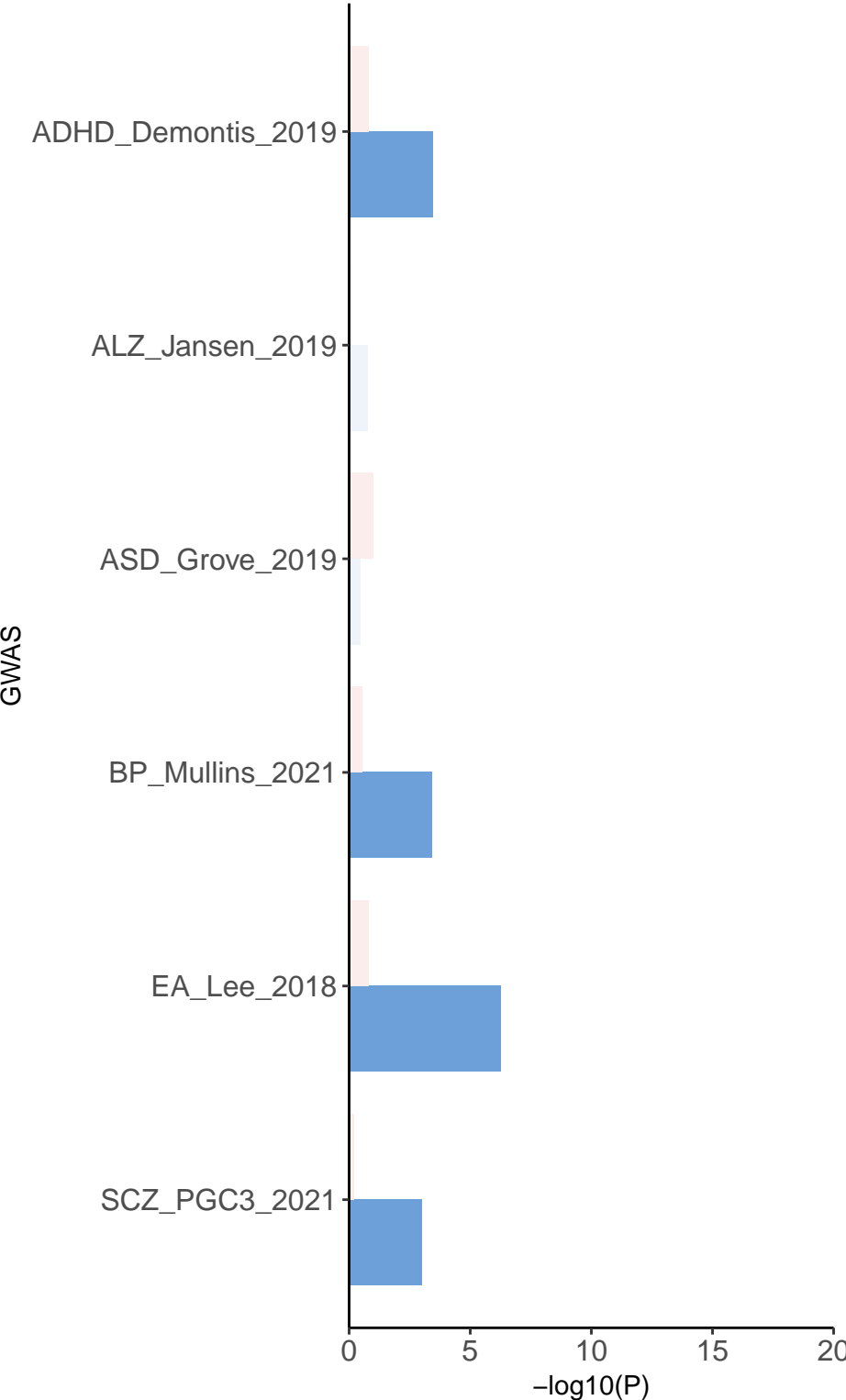
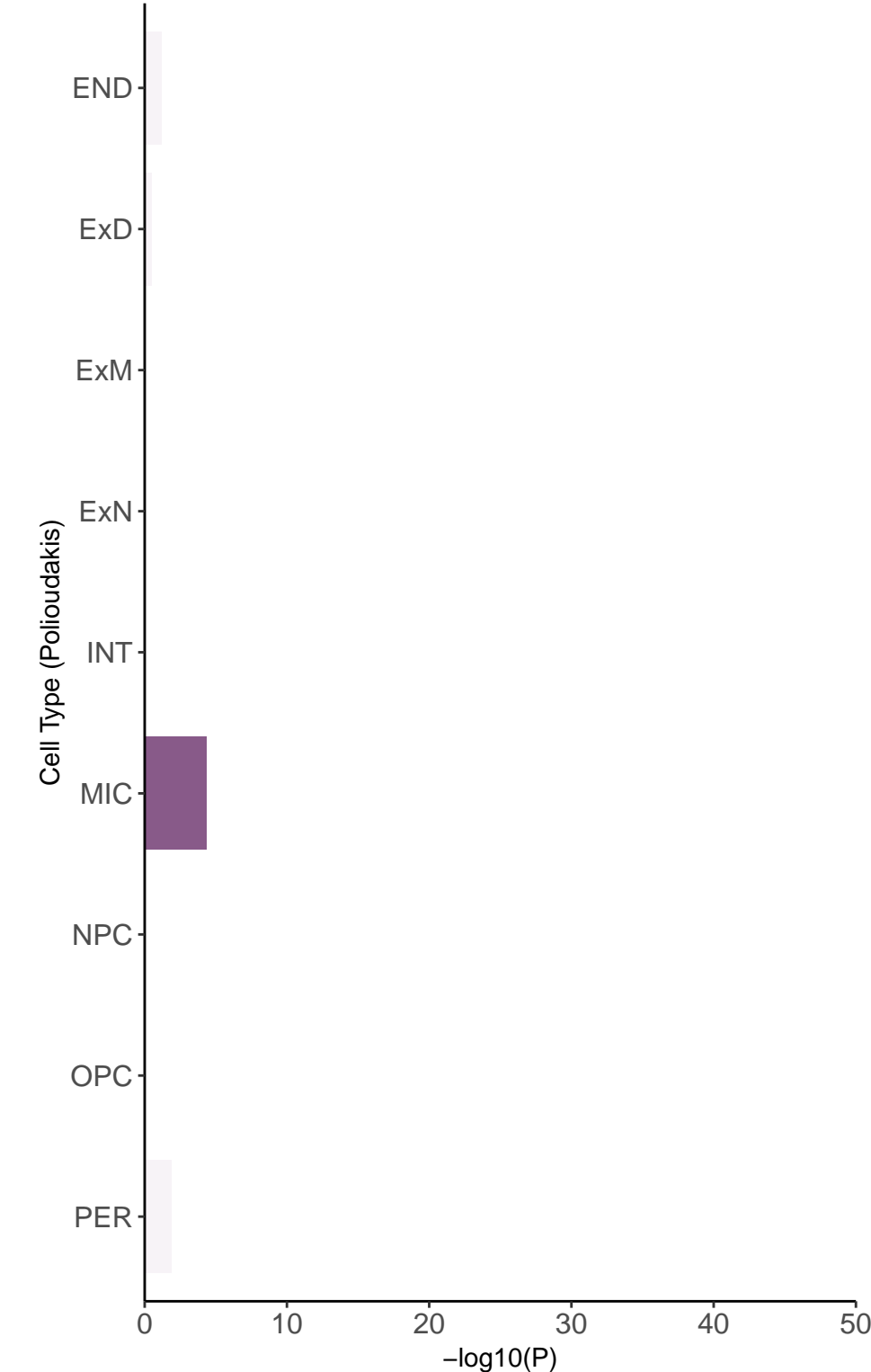
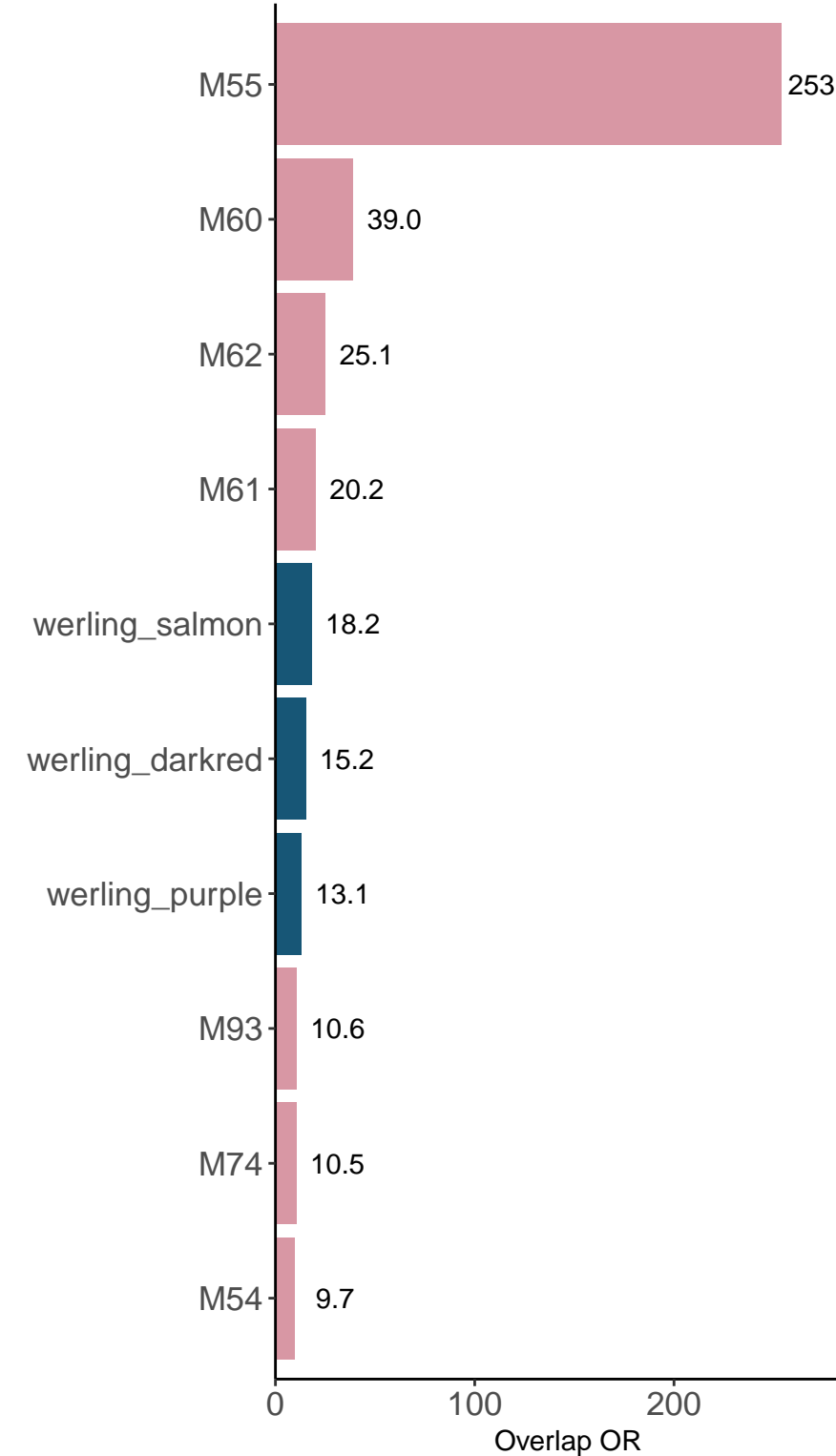
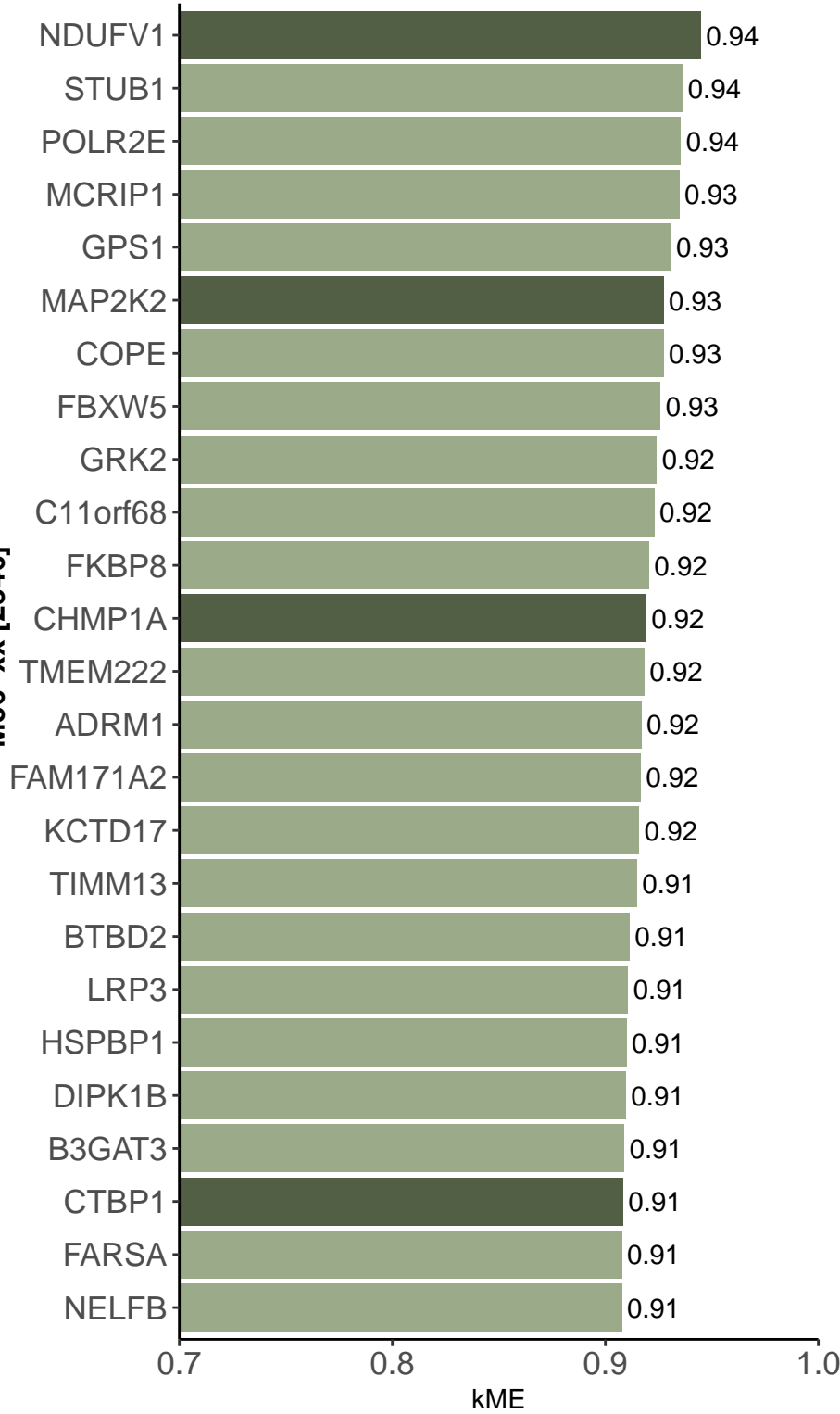


M56-xx [2346]



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:0005739	mitochondrion	1556	3.1e-43
2	GO:BP	GO:0044248	cellular catabolic process	1938	2.4e-40
3	GO:BP	GO:1901575	organic substance catabolic process	1896	2.6e-34
4	GO:CC	GO:1902494	catalytic complex	1628	1.4e-32
5	GO:CC	GO:0031975	envelope	1164	2.1e-31
6	GO:MF	GO:0000166	nucleotide binding	1973	6.1e-30
7	GO:MF	GO:1901265	nucleoside phosphate binding	1974	6.5e-30
8	GO:BP	GO:0009057	macromolecule catabolic process	1239	7.4e-29
9	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	2.0e-25
10	REAC	REAC:R-HSA-1430728	Metabolism	1832	6.1e-25
11	GO:MF	GO:0019899	enzyme binding	1888	1.2e-24
12	REAC	REAC:R-HSA-597592	Post-translational protein modification	1257	1.7e-22
13	KEGG	KEGG:01100	Metabolic pathways	1334	2.5e-18
14	KEGG	KEGG:05022	Pathways of neurodegeneration – multiple diseases	421	8.1e-13
15	KEGG	KEGG:04144	Endocytosis	238	2.7e-10

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