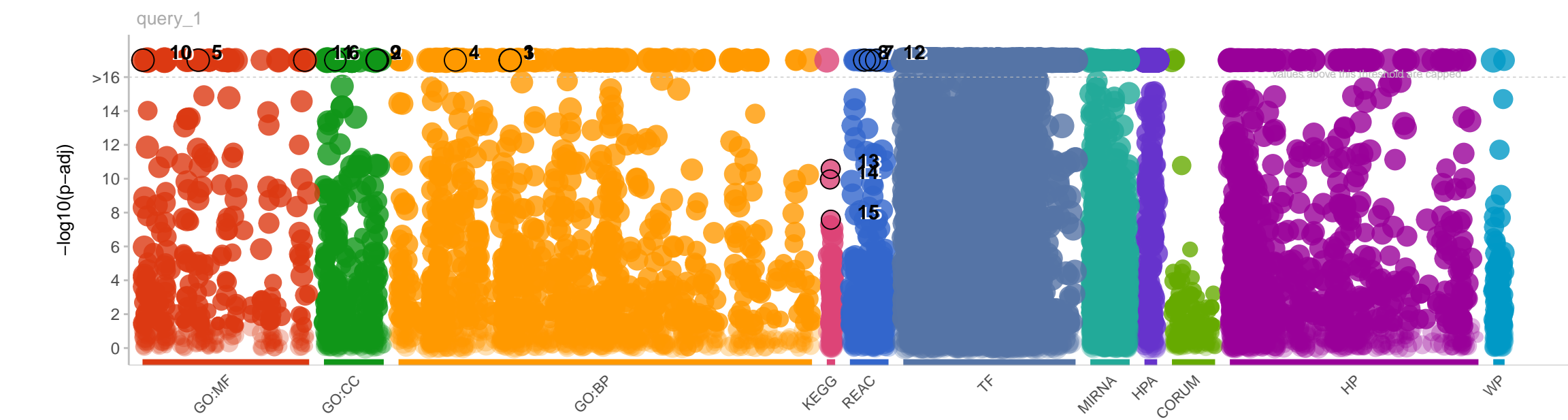
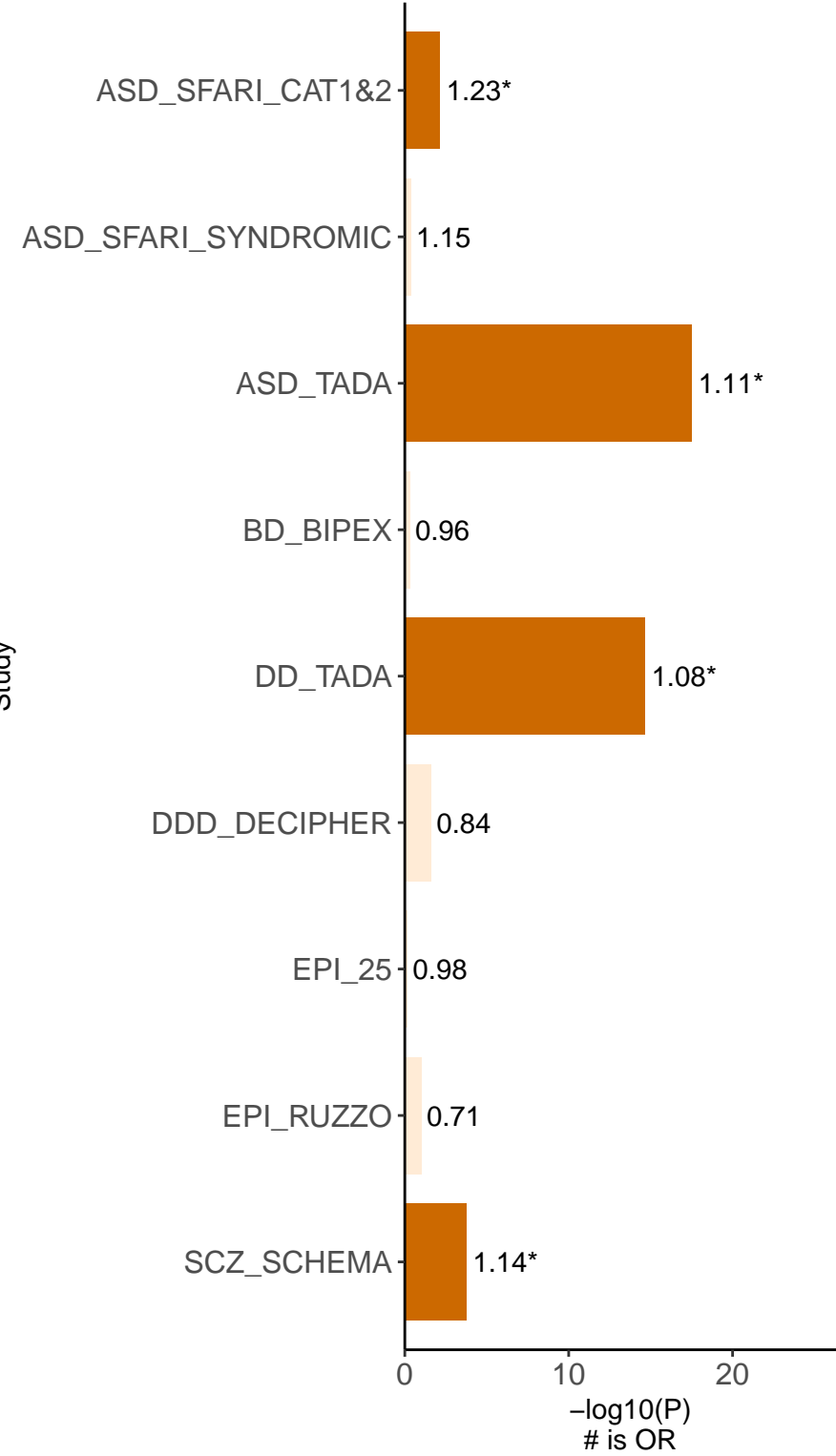
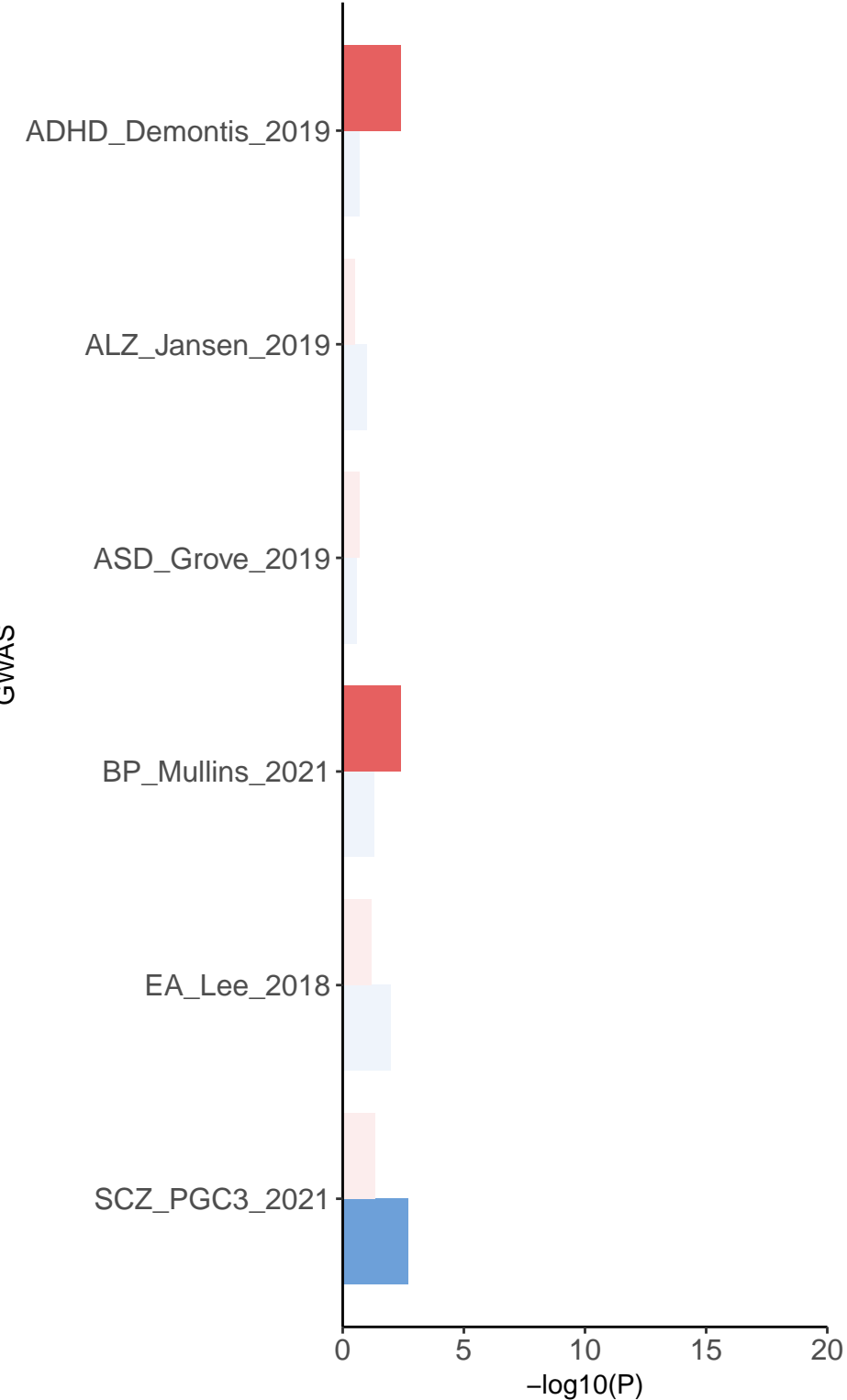
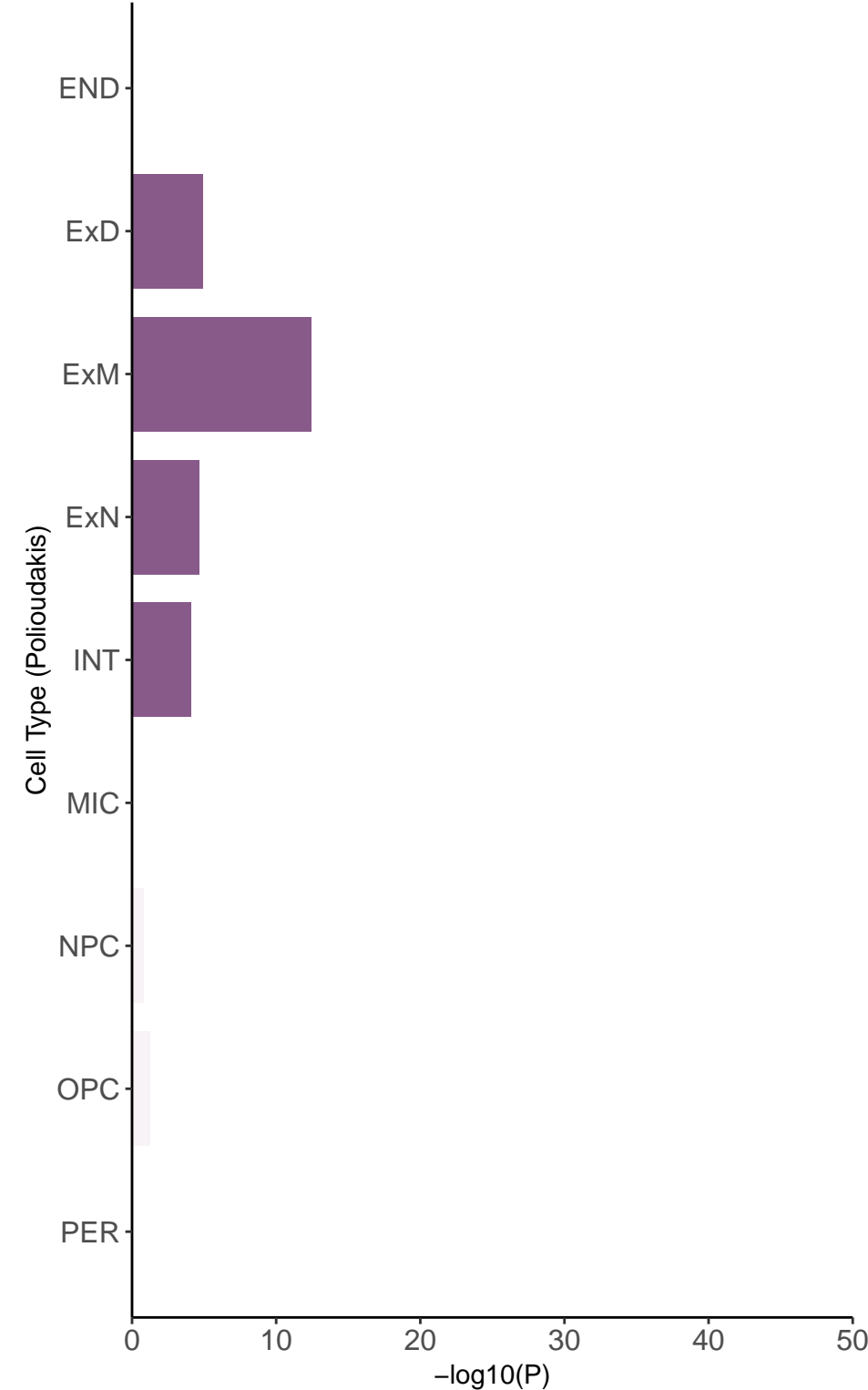
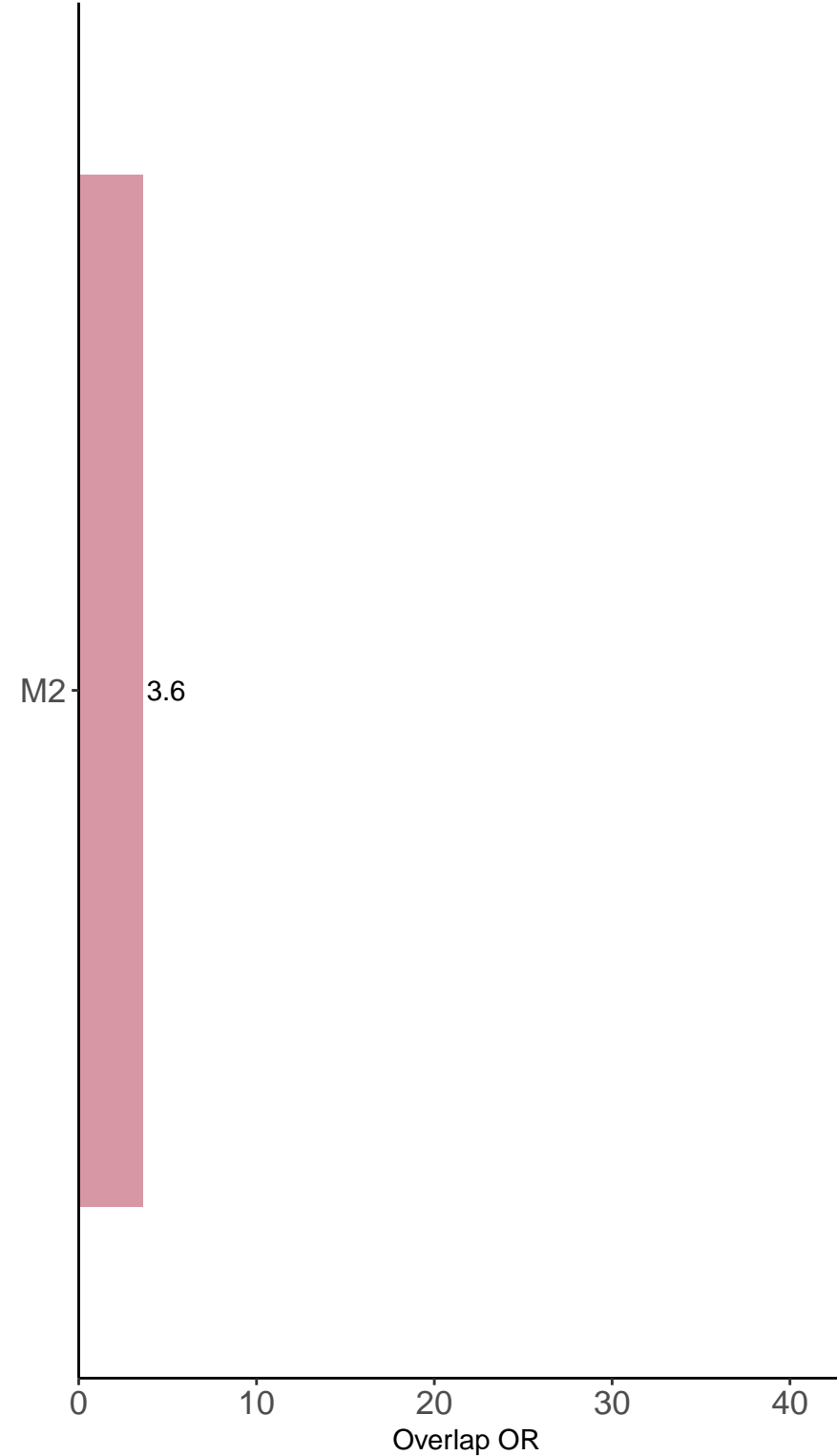
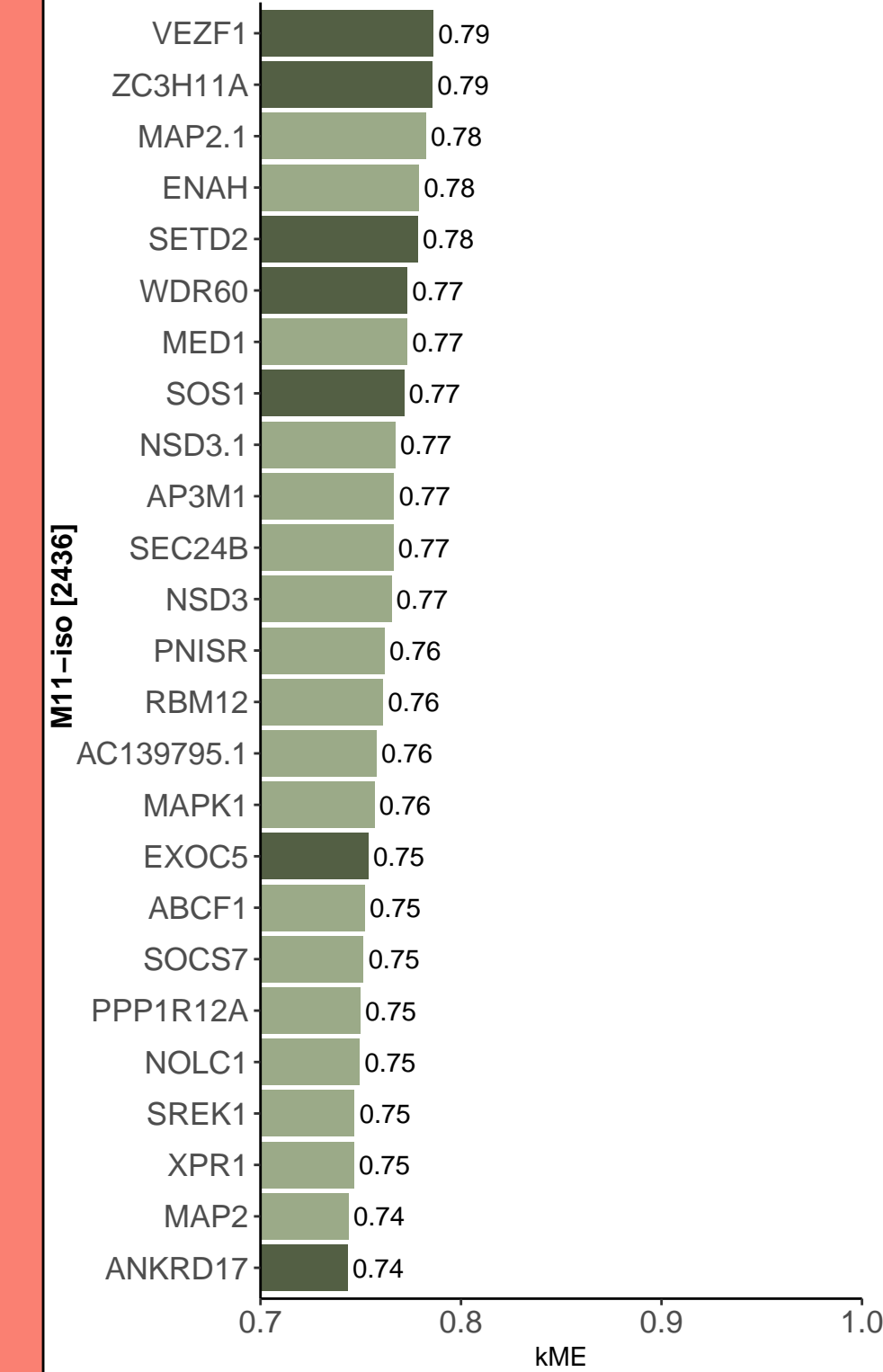


M11-Iso [2436]



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0031324	negative regulation of cellular metabolic process	1963	1.1e-77
2	GO:CC	GO:1902494	catalytic complex	1628	3.9e-66
3	GO:BP	GO:0031327	negative regulation of cellular biosynthetic process	1375	9.6e-60
4	GO:BP	GO:0009890	negative regulation of biosynthetic process	1402	1.1e-58
5	GO:MF	GO:0019899	enzyme binding	1888	2.2e-56
6	GO:CC	GO:0016604	nuclear body	796	4.6e-43
7	REAC	REAC:R-HSA-8953854	Metabolism of RNA	626	1.7e-41
8	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	6.7e-39
9	GO:CC	GO:0140535	intracellular protein-containing complex	869	4.3e-36
10	GO:MF	GO:0000166	nucleotide binding	1973	1.2e-34
11	GO:MF	GO:1901265	nucleoside phosphate binding	1974	1.3e-34
12	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	8.0e-34
13	KEGG	KEGG:04120	Ubiquitin mediated proteolysis	129	2.7e-11
14	KEGG	KEGG:03040	Spliceosome	133	1.1e-10
15	KEGG	KEGG:04140	Autophagy - animal	131	2.6e-08

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