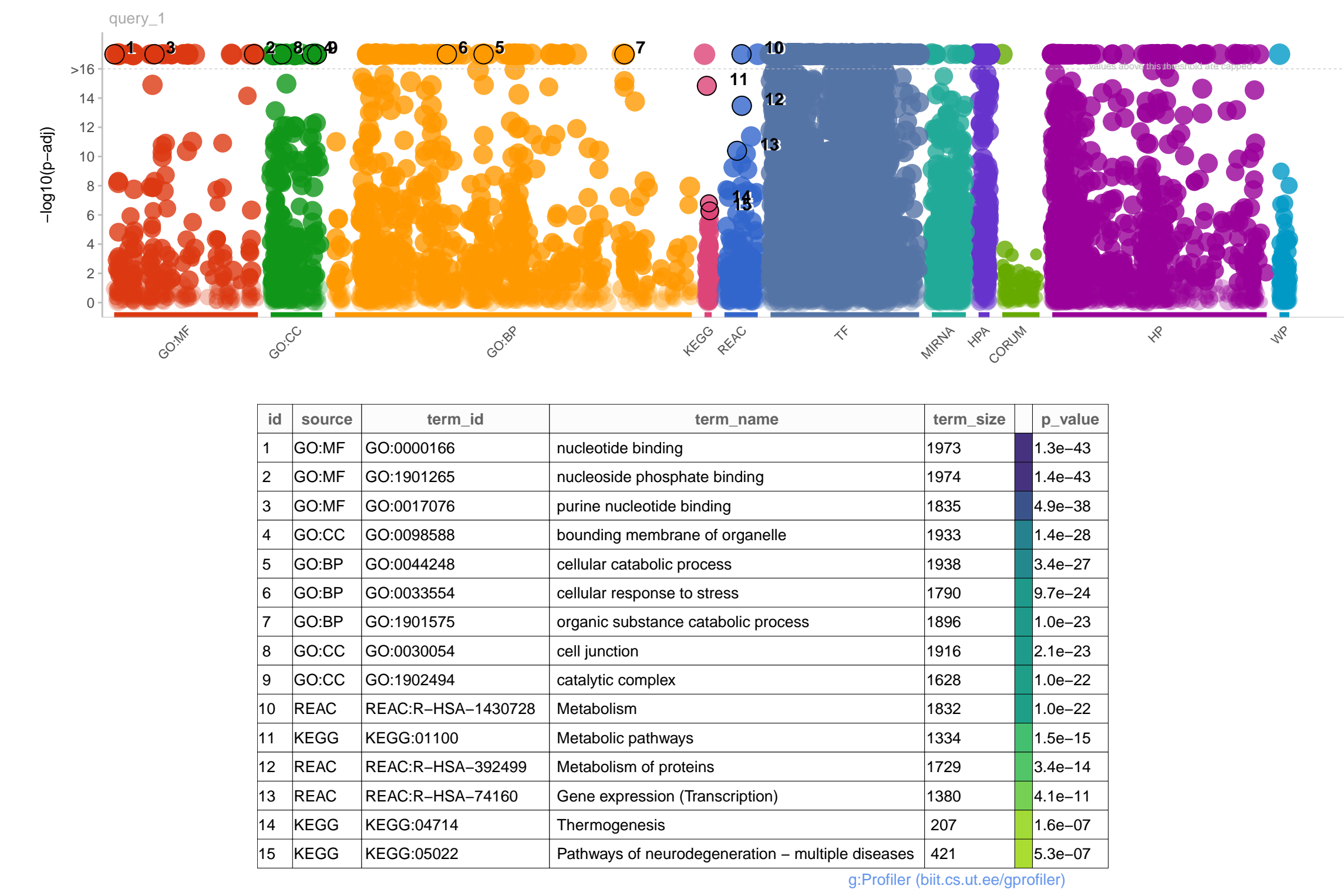
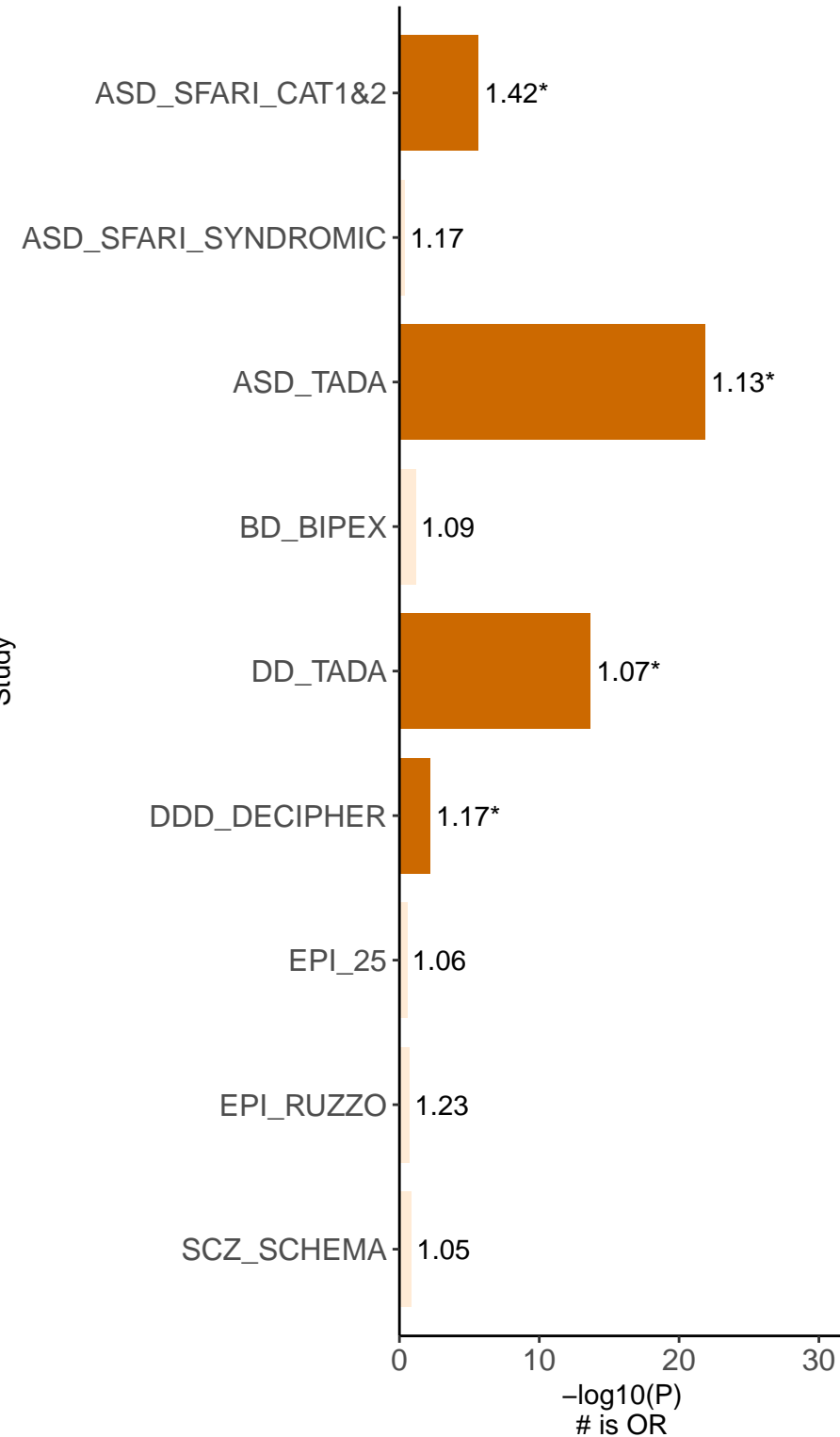
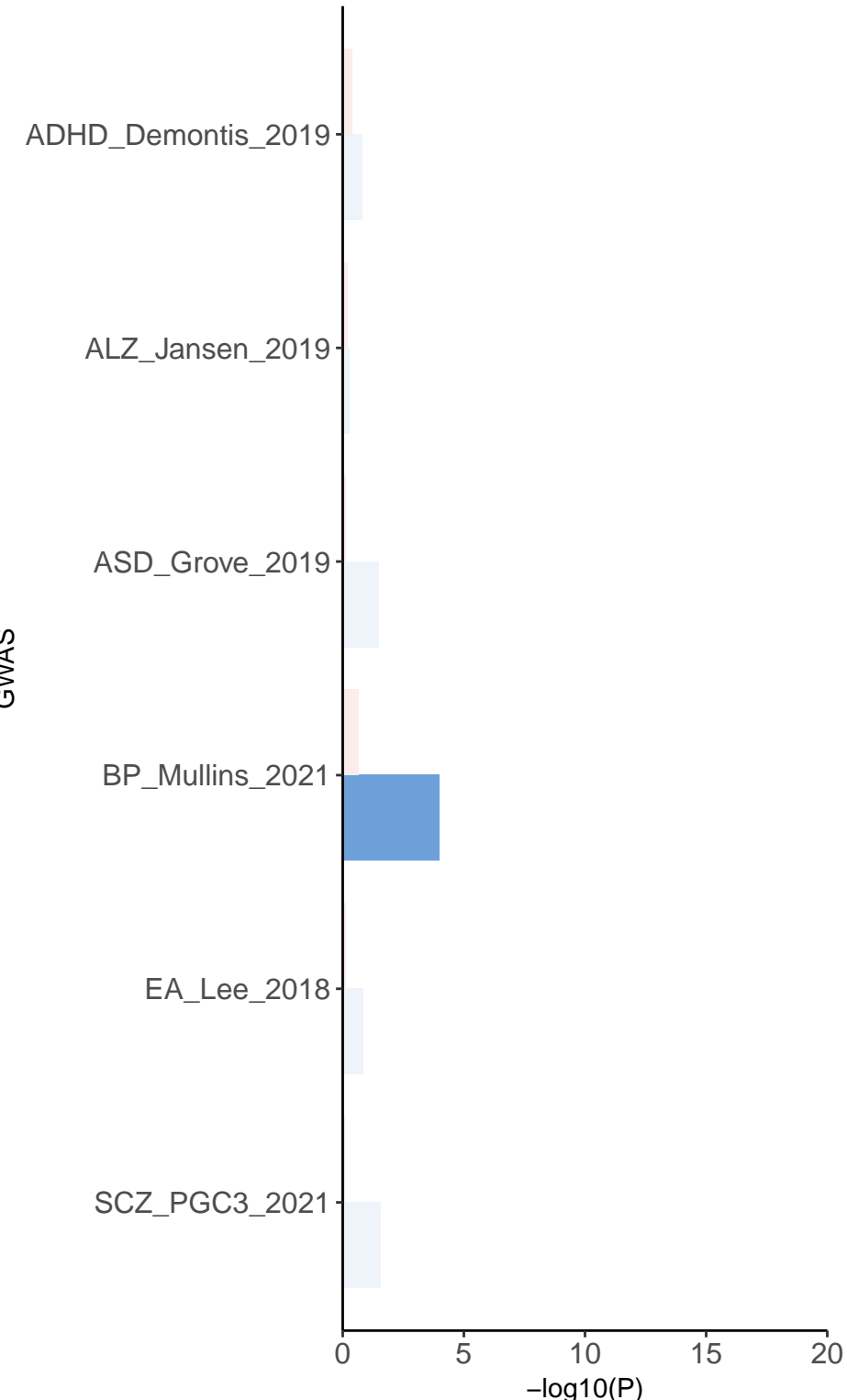
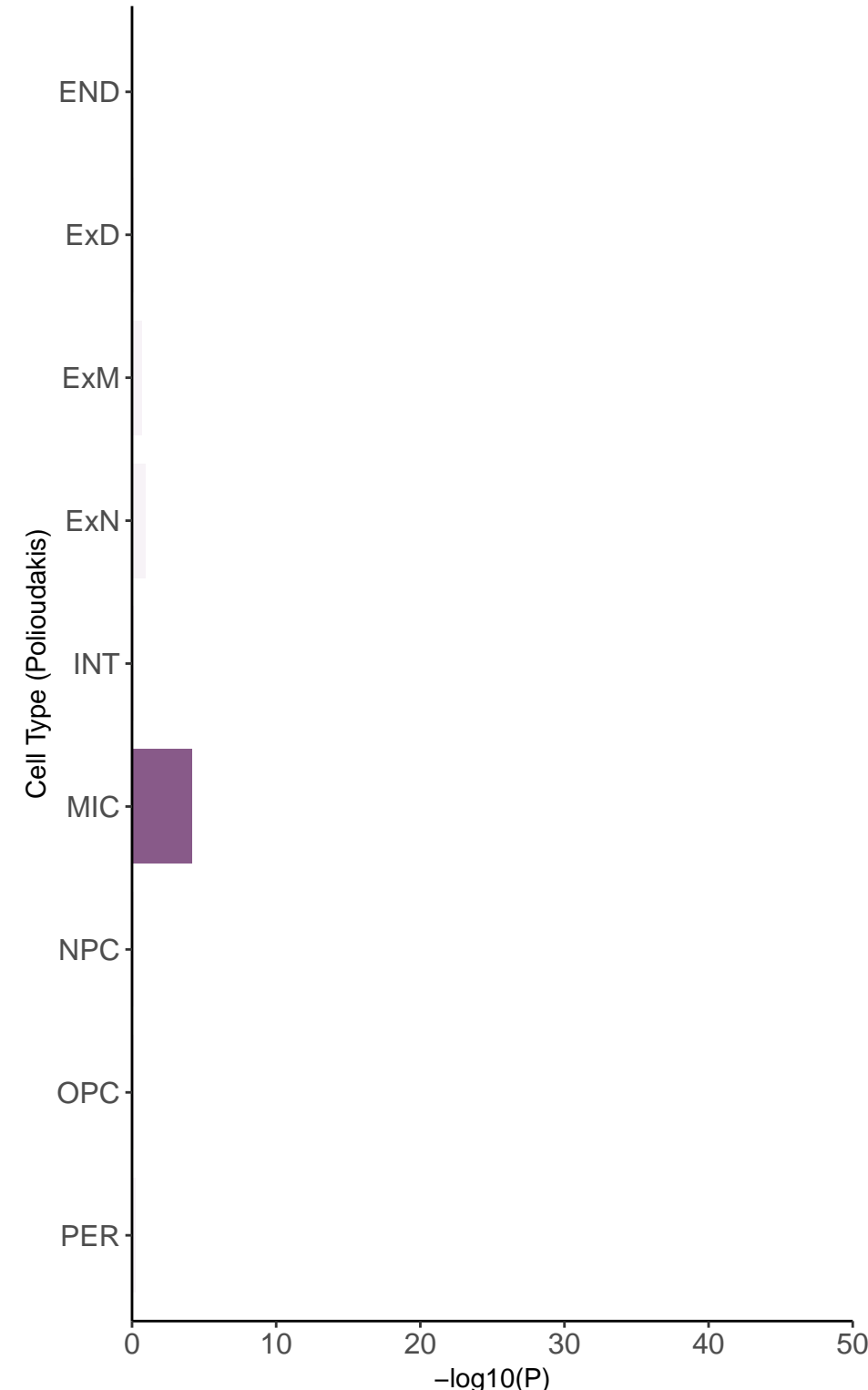
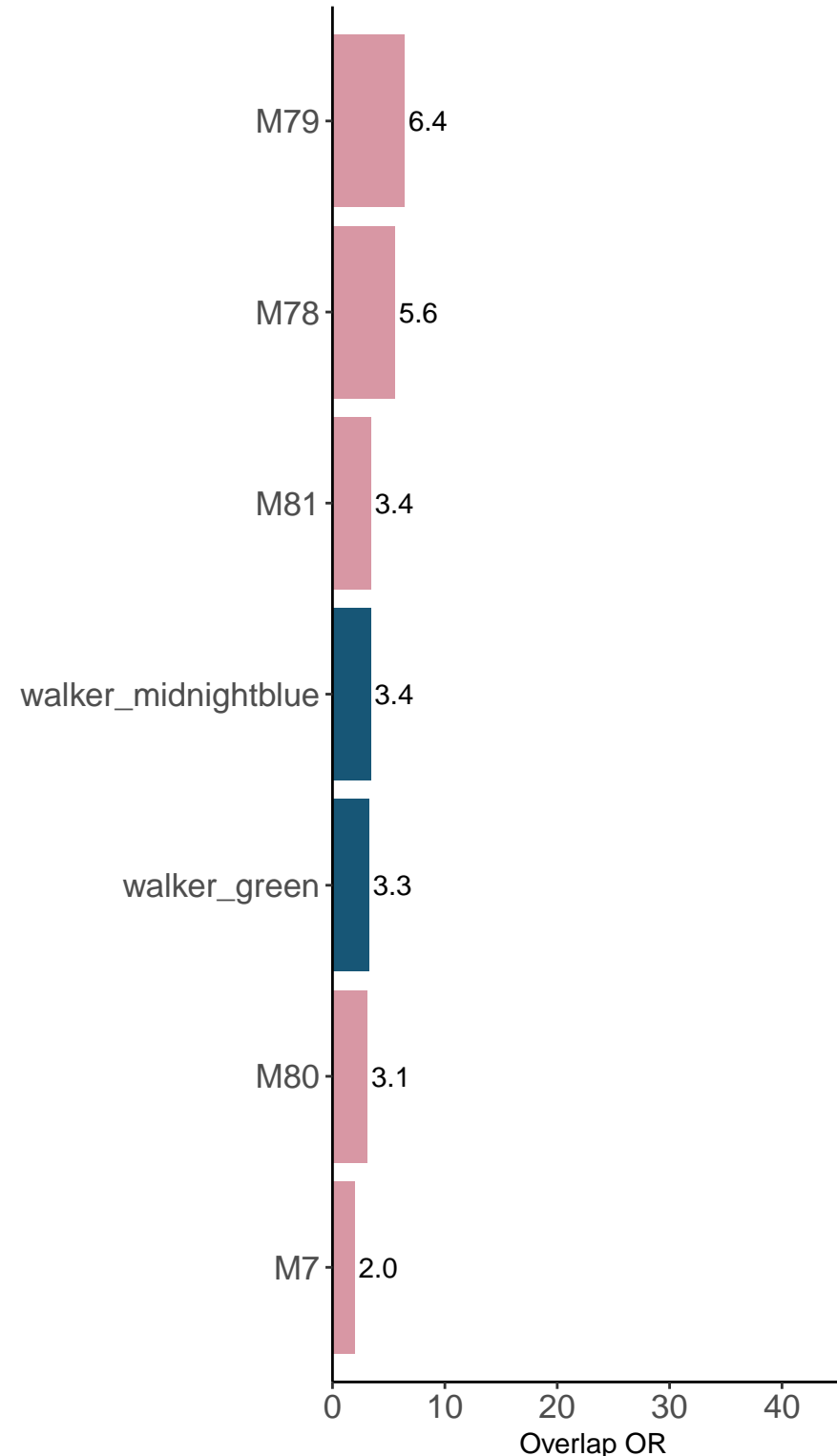
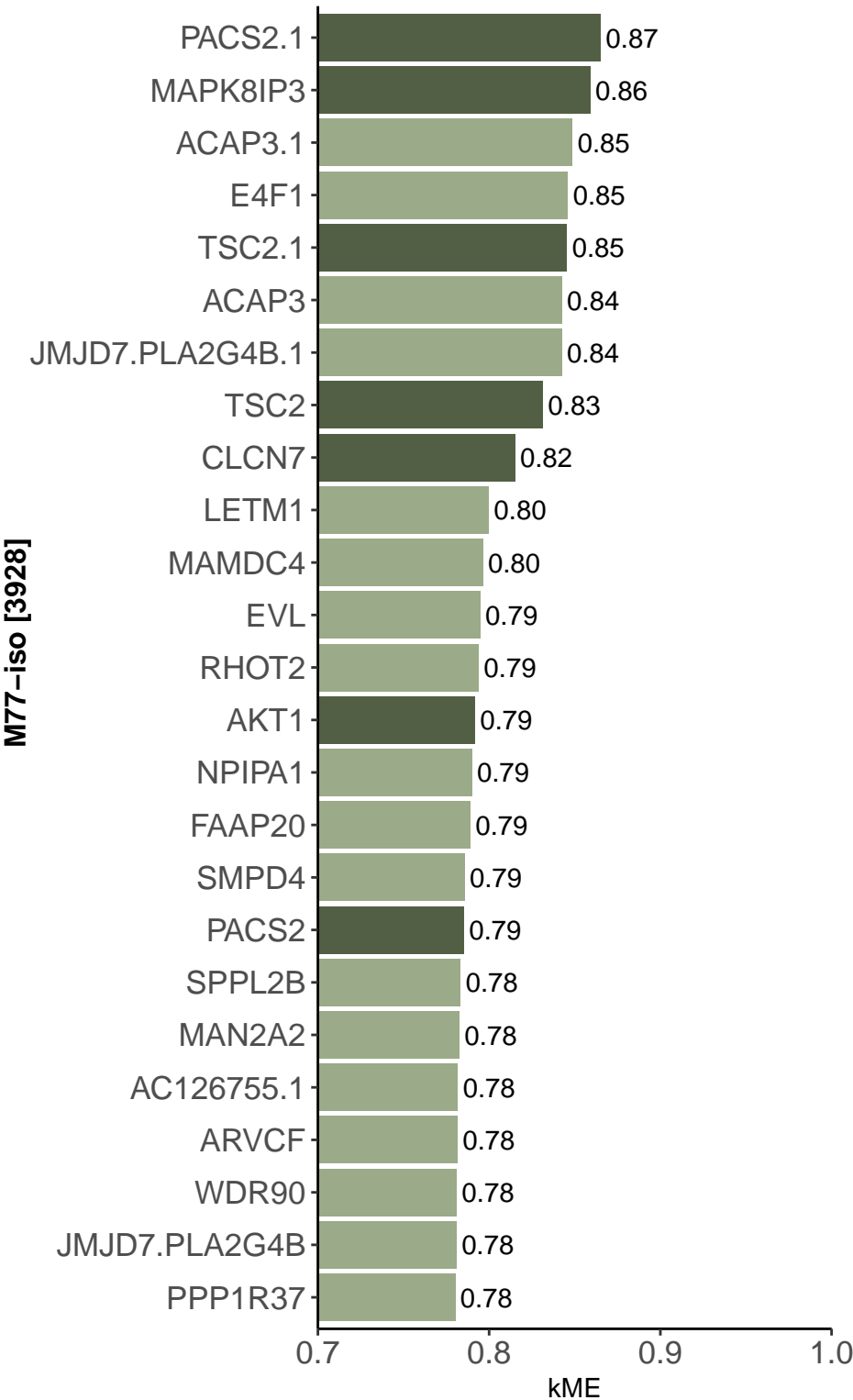


M77-iso [3928]



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0000166	nucleotide binding	1973	1.3e-43
2	GO:MF	GO:1901265	nucleoside phosphate binding	1974	1.4e-43
3	GO:MF	GO:0017076	purine nucleotide binding	1835	4.9e-38
4	GO:CC	GO:0098588	bounding membrane of organelle	1933	1.4e-28
5	GO:BP	GO:0044248	cellular catabolic process	1938	3.4e-27
6	GO:BP	GO:0033554	cellular response to stress	1790	9.7e-24
7	GO:BP	GO:1901575	organic substance catabolic process	1896	1.0e-23
8	GO:CC	GO:0030054	cell junction	1916	2.1e-23
9	GO:CC	GO:1902494	catalytic complex	1628	1.0e-22
10	REAC	REAC:R-HSA-1430728	Metabolism	1832	1.0e-22
11	KEGG	KEGG:01100	Metabolic pathways	1334	1.5e-15
12	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	3.4e-14
13	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	4.1e-11
14	KEGG	KEGG:04714	Thermogenesis	207	1.6e-07
15	KEGG	KEGG:05022	Pathways of neurodegeneration – multiple diseases	421	5.3e-07

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