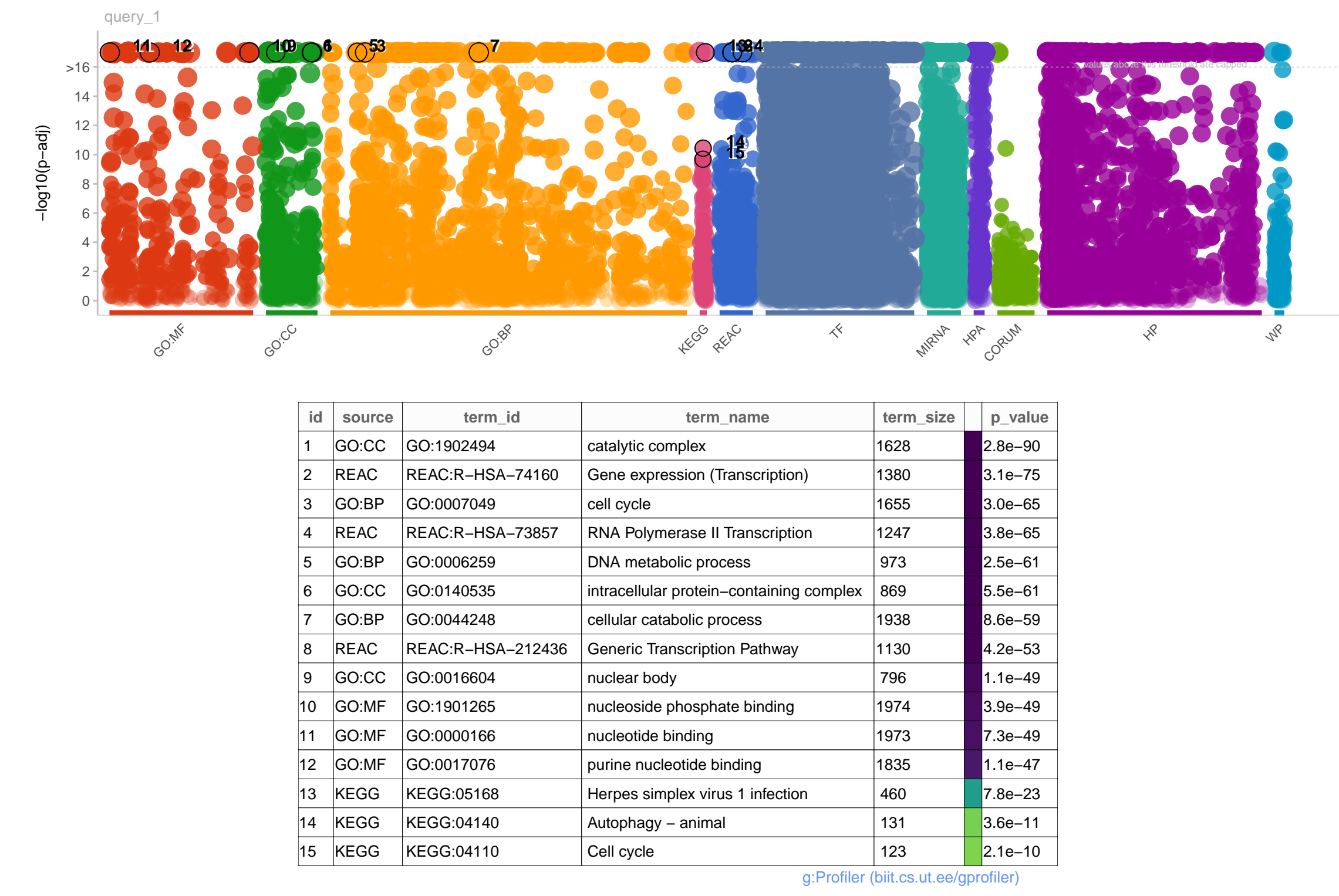
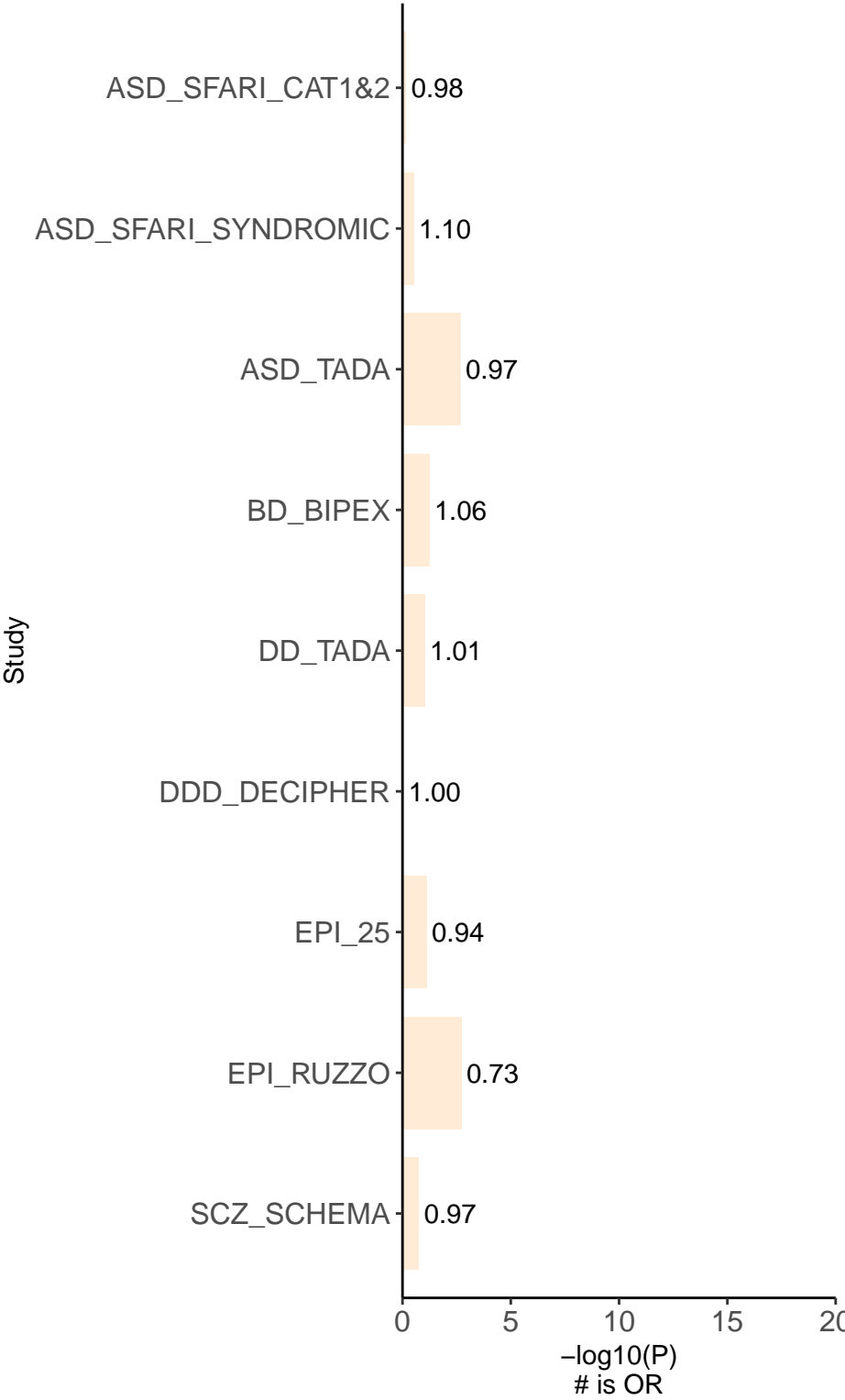
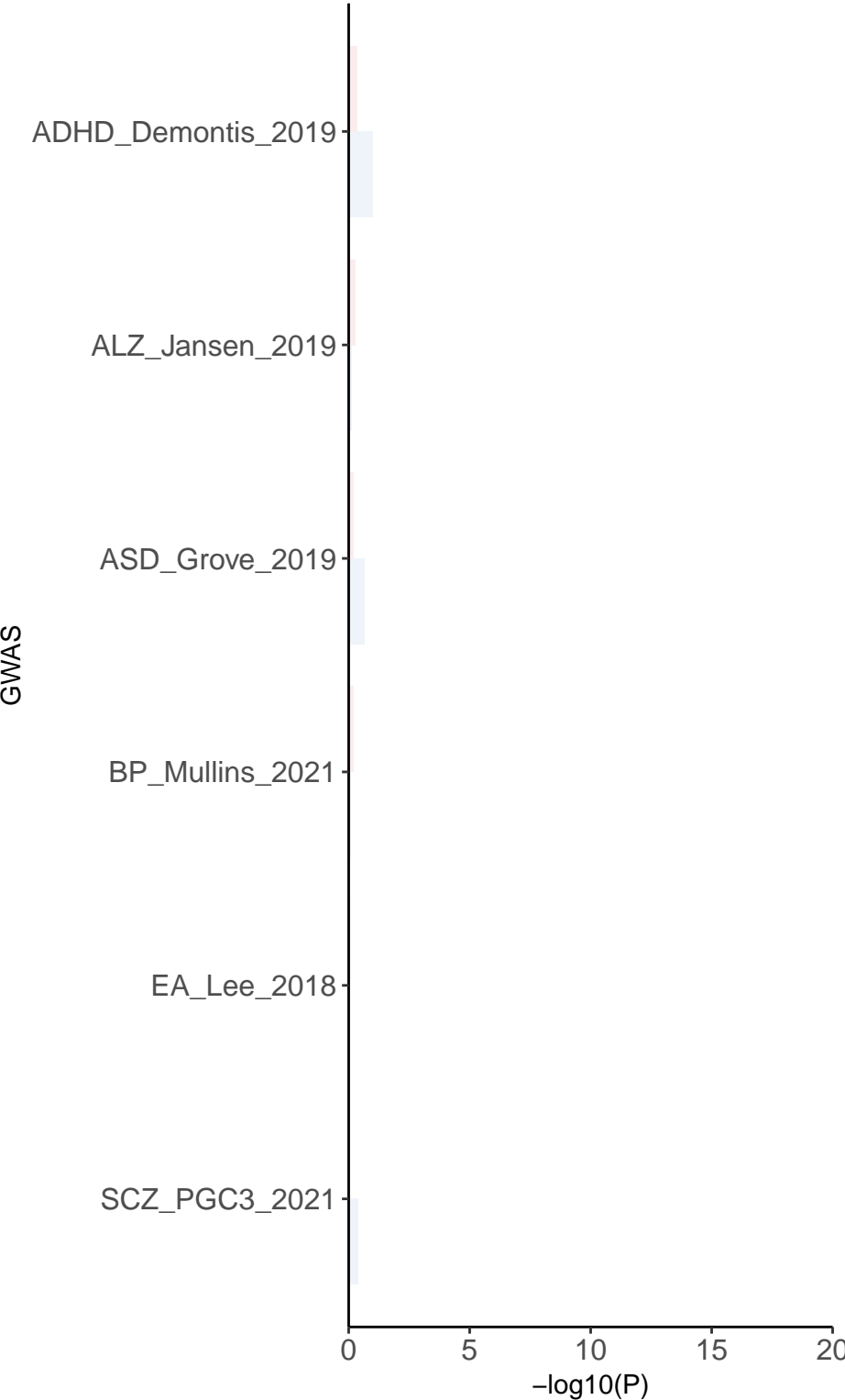
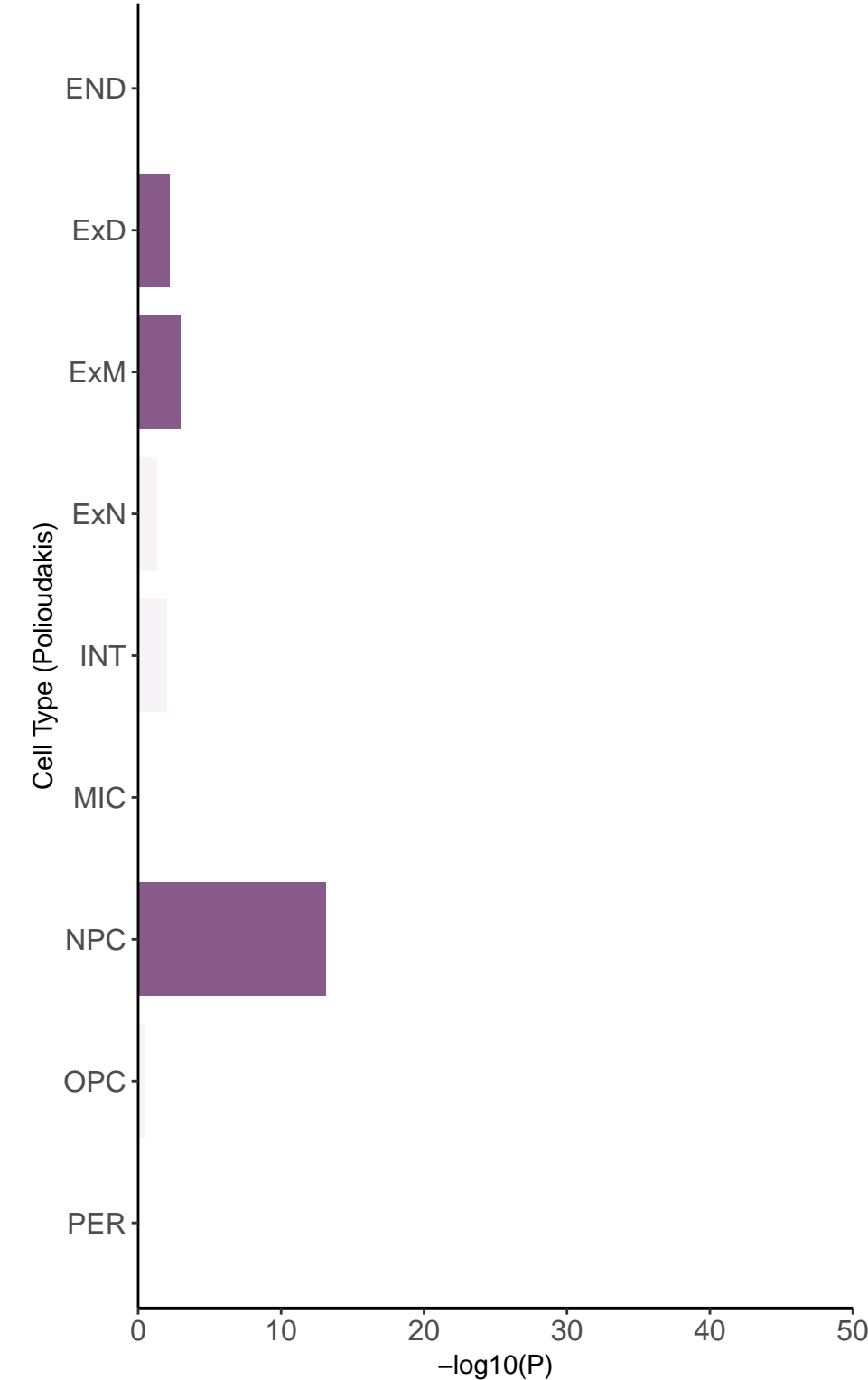
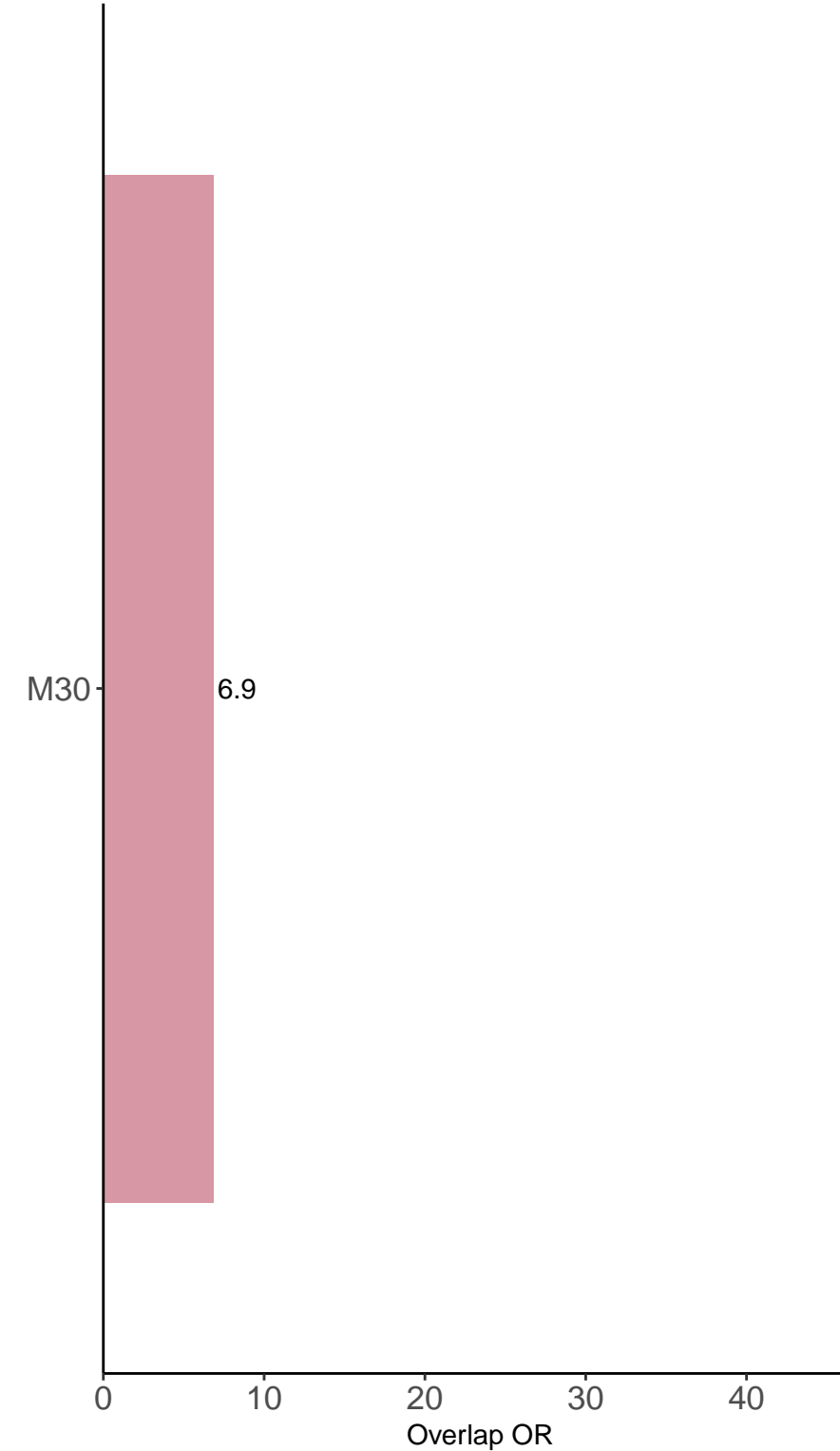
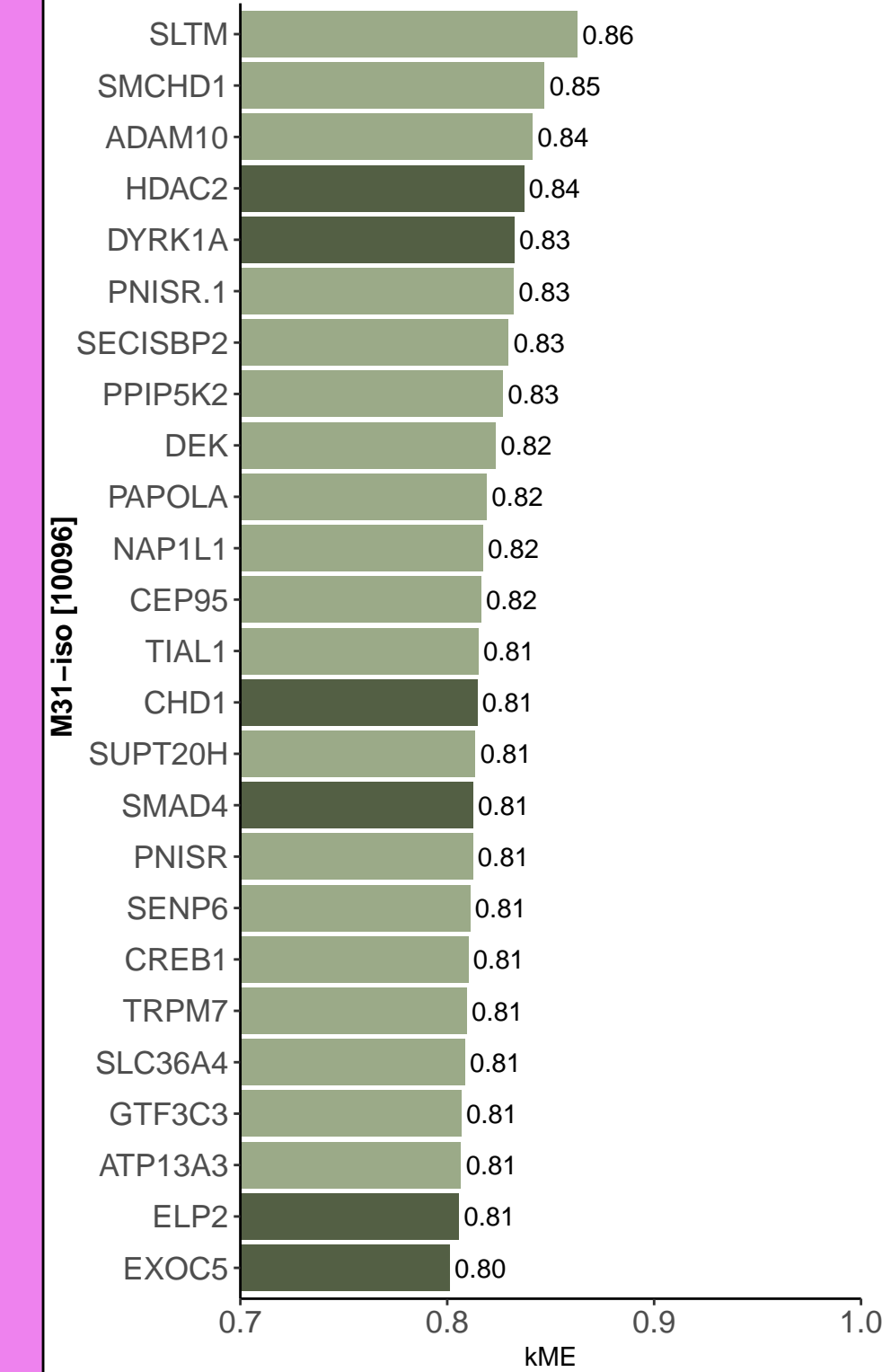


M31-iso [10096]



id	source	term_id	term_name	term_size	p_value
1	GO:CC	GO:1902494	catalytic complex	1628	2.8e-90
2	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	3.1e-75
3	GO:BP	GO:0007049	cell cycle	1655	3.0e-65
4	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	3.8e-65
5	GO:BP	GO:0006259	DNA metabolic process	973	2.5e-61
6	GO:CC	GO:0140535	intracellular protein-containing complex	869	5.5e-61
7	GO:BP	GO:0044248	cellular catabolic process	1938	8.6e-59
8	REAC	REAC:R-HSA-212436	Generic Transcription Pathway	1130	4.2e-53
9	GO:CC	GO:0016604	nuclear body	796	1.1e-49
10	GO:MF	GO:1901265	nucleoside phosphate binding	1974	3.9e-49
11	GO:MF	GO:0000166	nucleotide binding	1973	7.3e-49
12	GO:MF	GO:0017076	purine nucleotide binding	1835	1.1e-47
13	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	7.8e-23
14	KEGG	KEGG:04140	Autophagy – animal	131	3.6e-11
15	KEGG	KEGG:04110	Cell cycle	123	2.1e-10

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