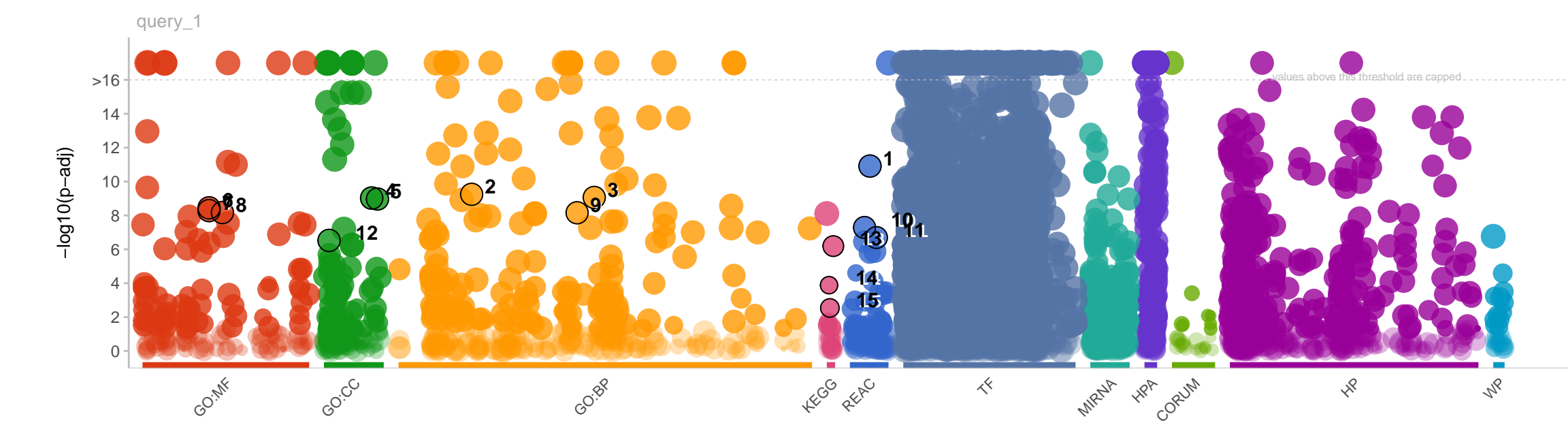
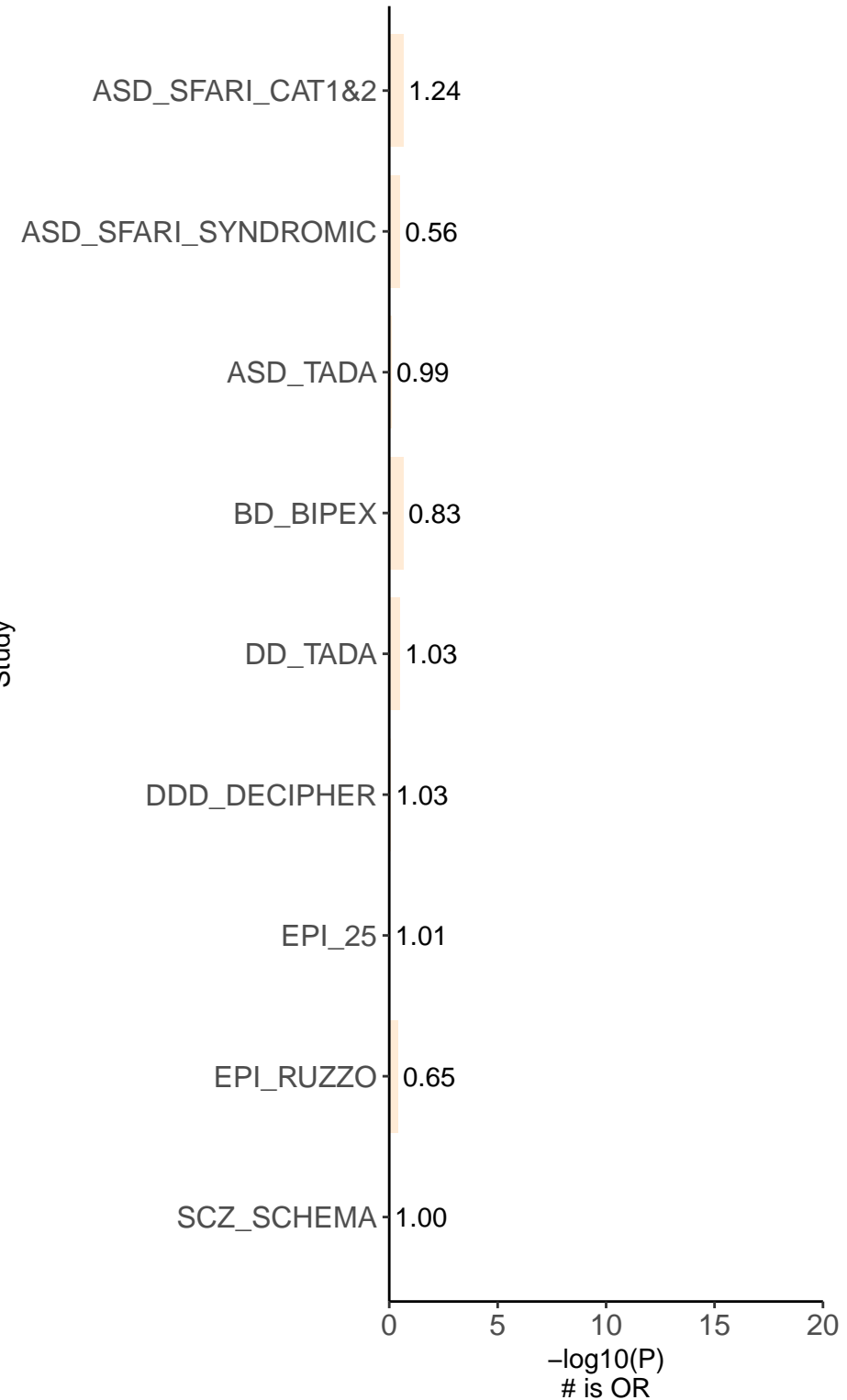
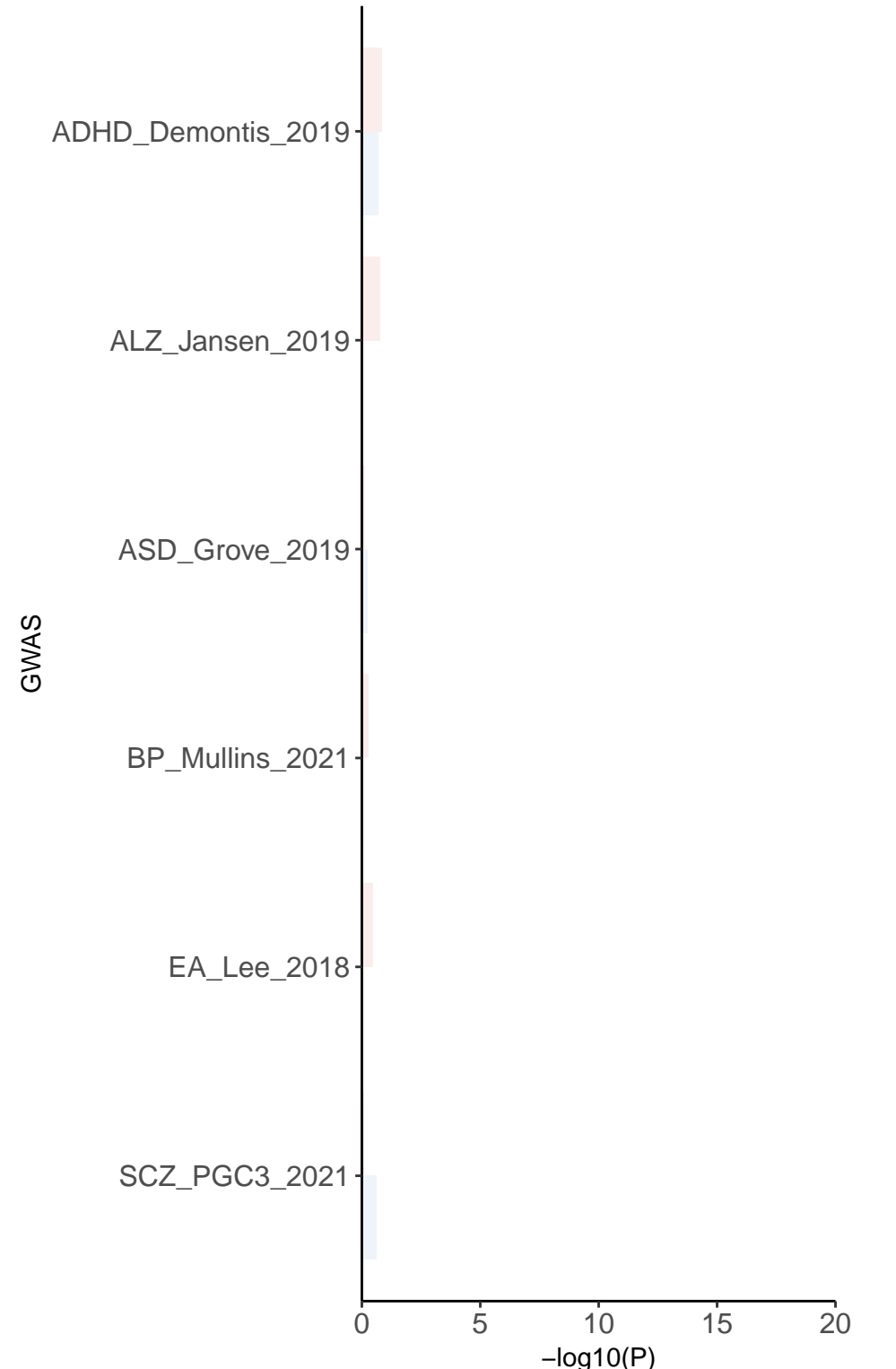
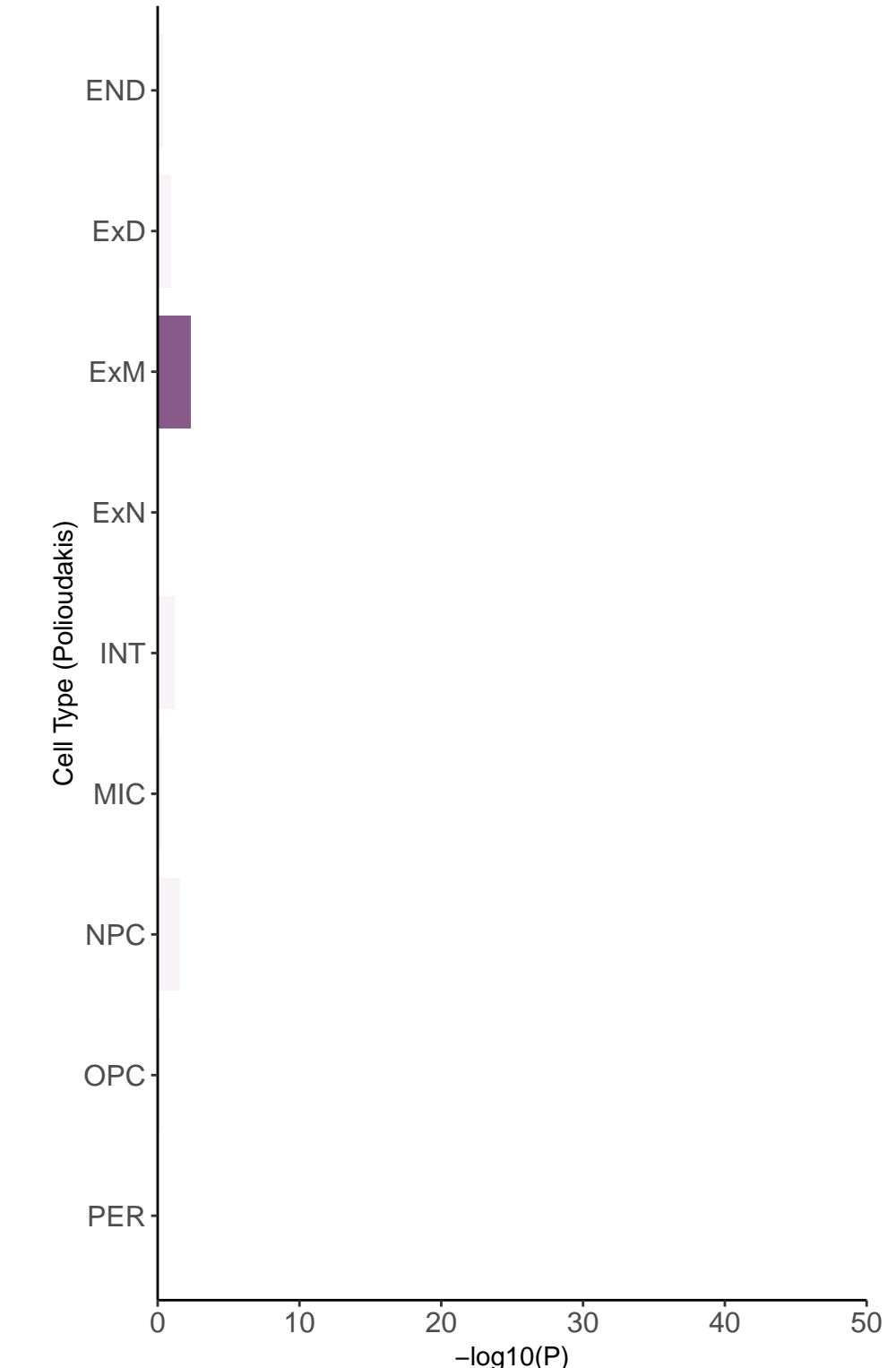
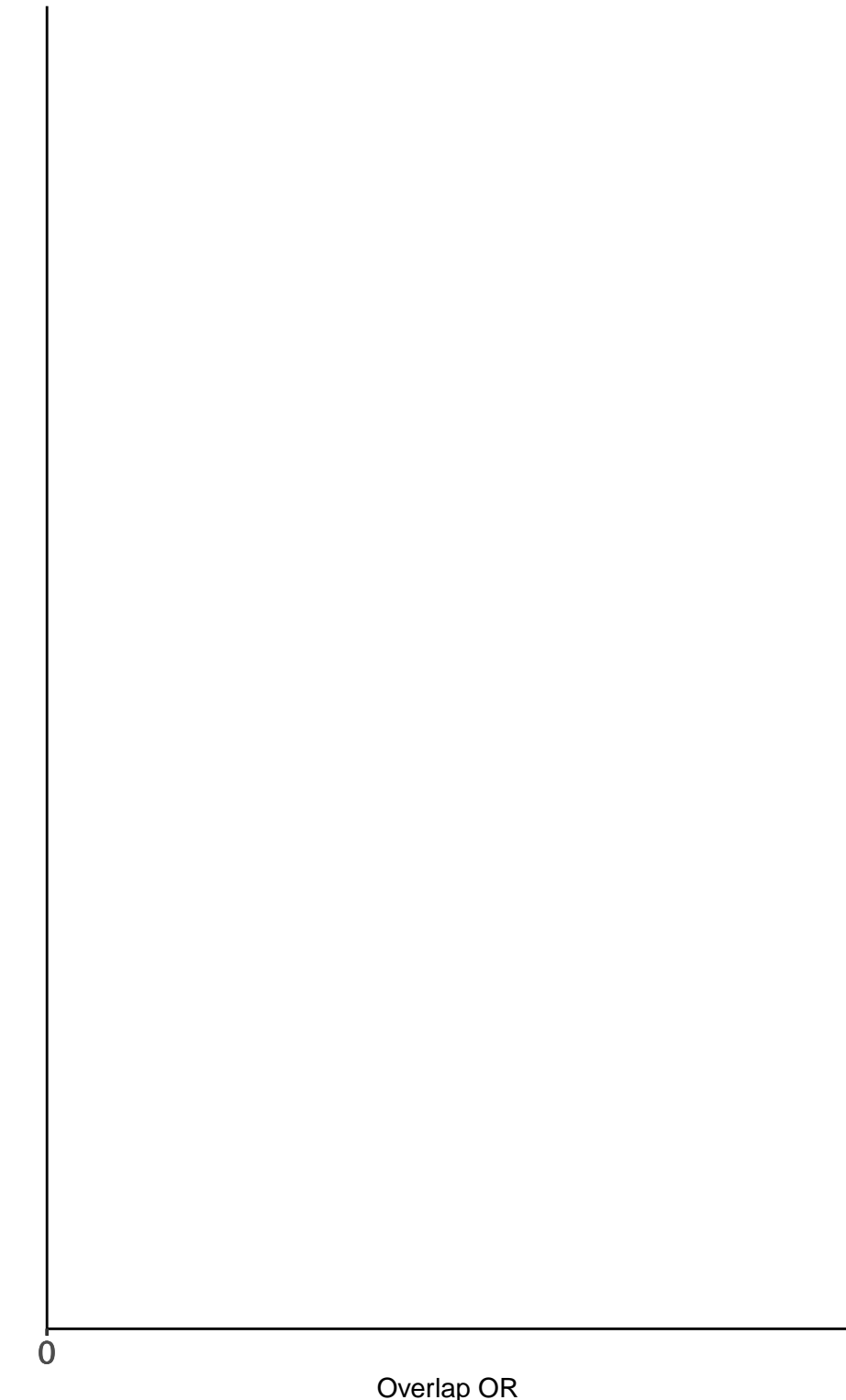
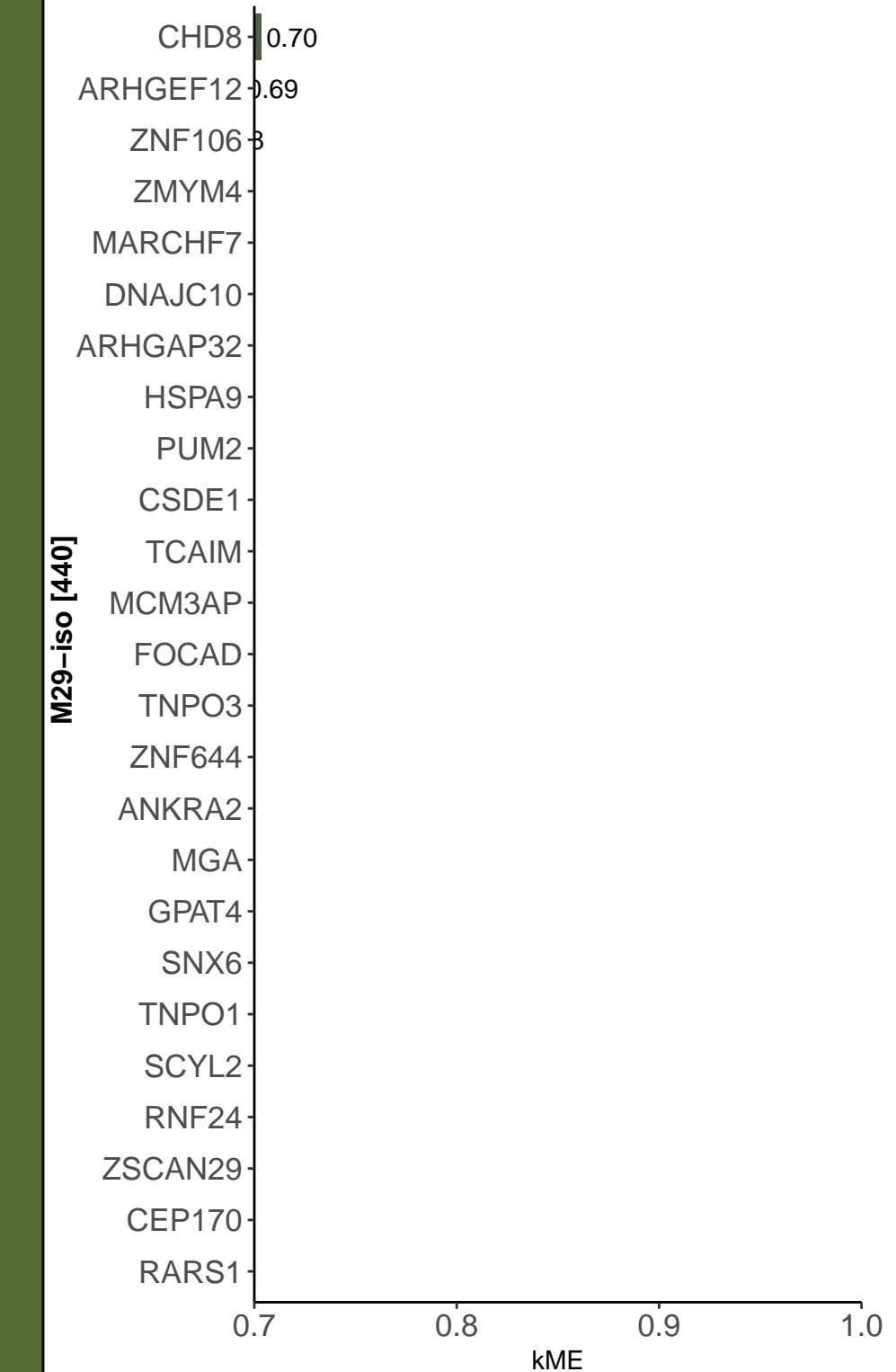


M29-iso [440]



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-392499	Metabolism of proteins	1729	1.2e-11
2	GO:BP	GO:0015031	protein transport	1541	5.7e-10
3	GO:BP	GO:0046907	intracellular transport	1572	8.7e-10
4	GO:CC	GO:0098588	bounding membrane of organelle	1933	9.4e-10
5	GO:CC	GO:1902494	catalytic complex	1628	1.1e-09
6	GO:MF	GO:0032555	purine ribonucleotide binding	1739	3.6e-09
7	GO:MF	GO:0032553	ribonucleotide binding	1754	5.2e-09
8	GO:MF	GO:0035639	purine ribonucleoside triphosphate binding	1676	6.8e-09
9	GO:BP	GO:0045184	establishment of protein localization	1643	7.0e-09
10	REAC	REAC:R-HSA-74160	Gene expression (Transcription)	1380	5.2e-08
11	REAC	REAC:R-HSA-73857	RNA Polymerase II Transcription	1247	2.0e-07
12	GO:CC	GO:0005783	endoplasmic reticulum	1784	3.0e-07
13	KEGG	KEGG:05168	Herpes simplex virus 1 infection	460	6.5e-07
14	KEGG	KEGG:00970	Aminoacyl-tRNA biosynthesis	44	1.3e-04
15	KEGG	KEGG:03013	Nucleocytoplasmic transport	98	2.9e-03

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