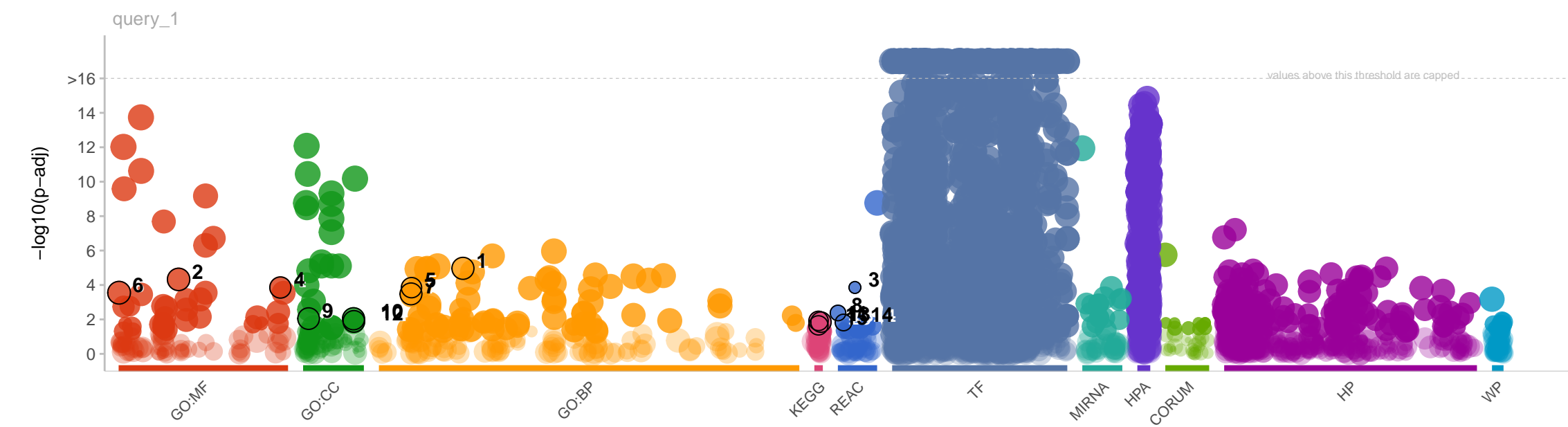
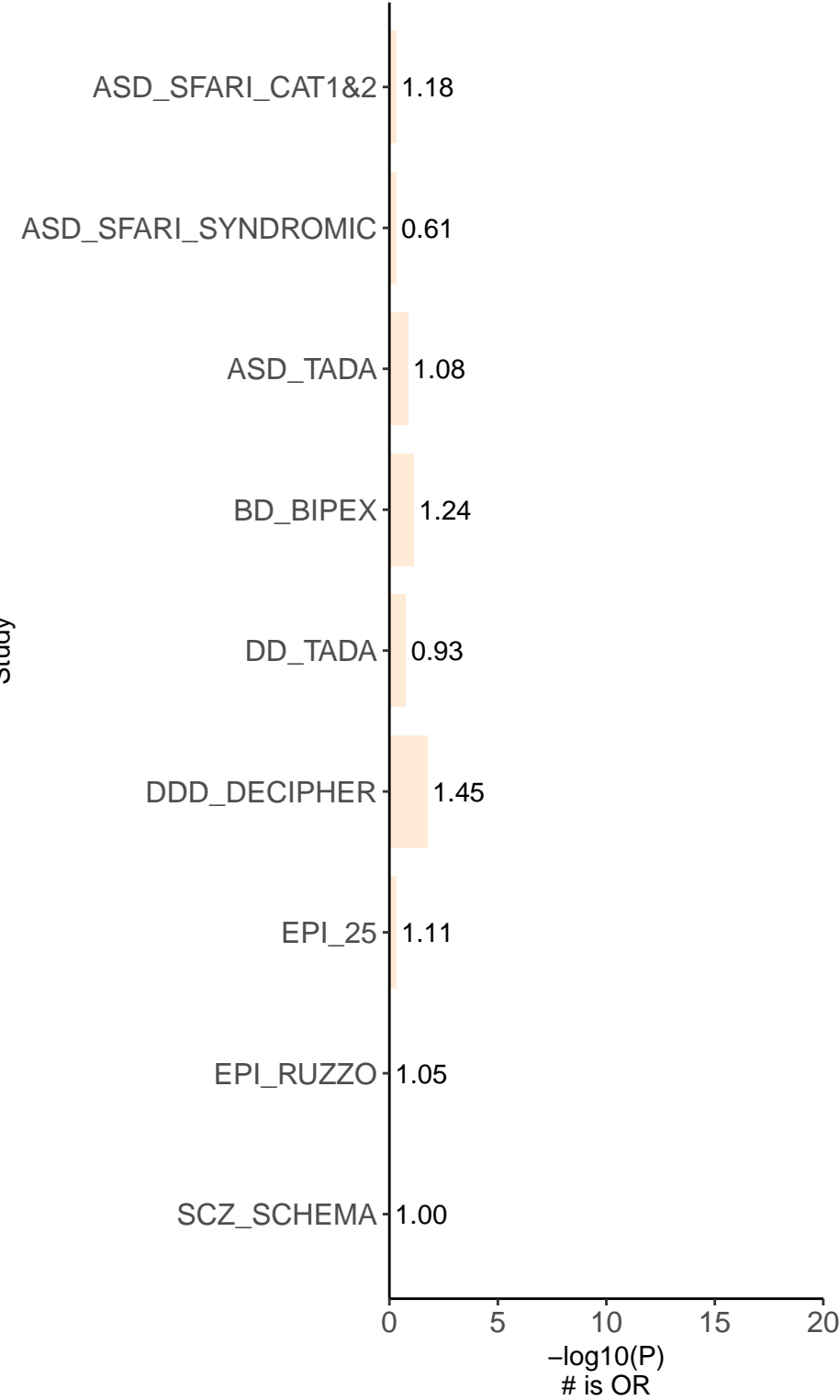
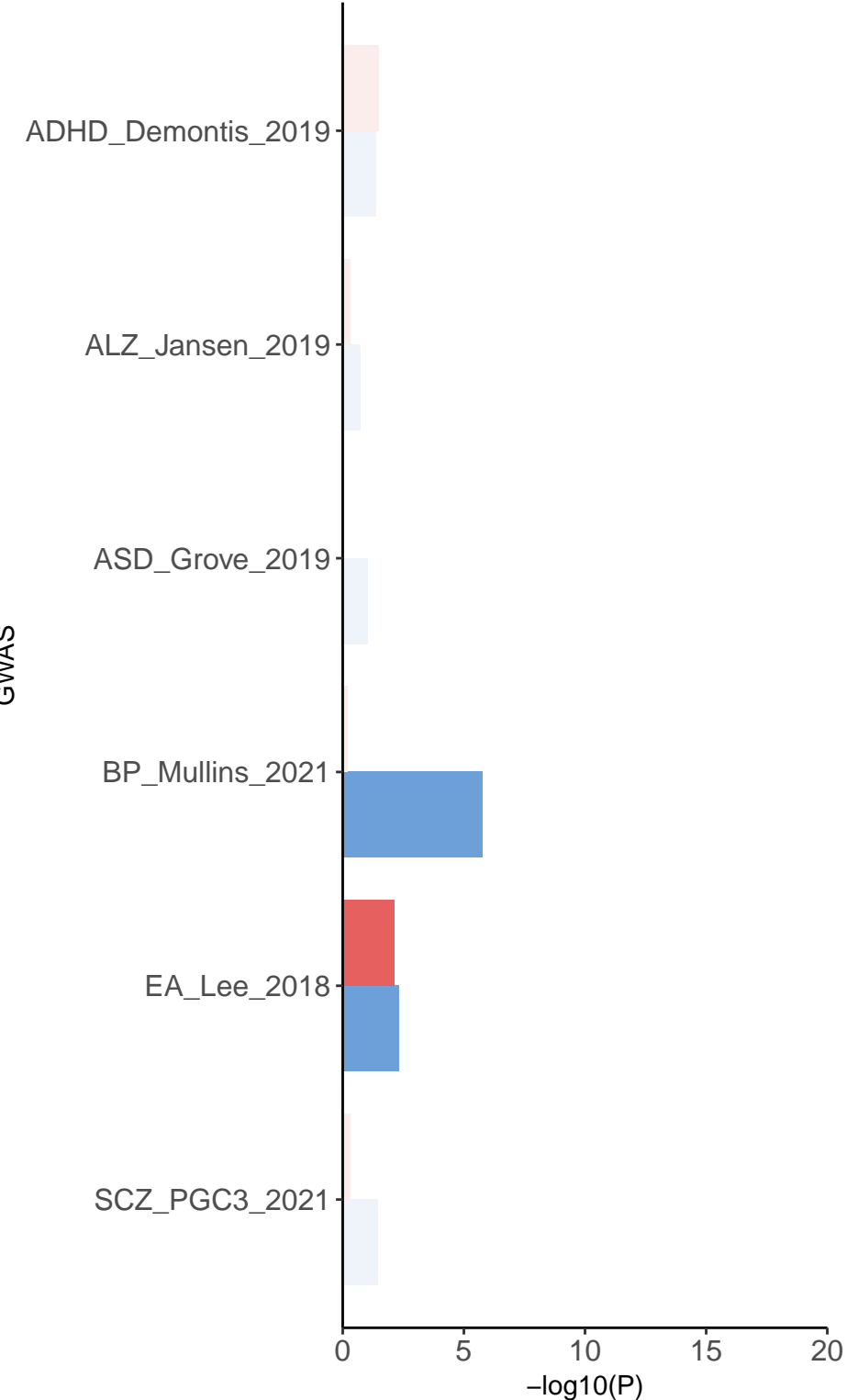
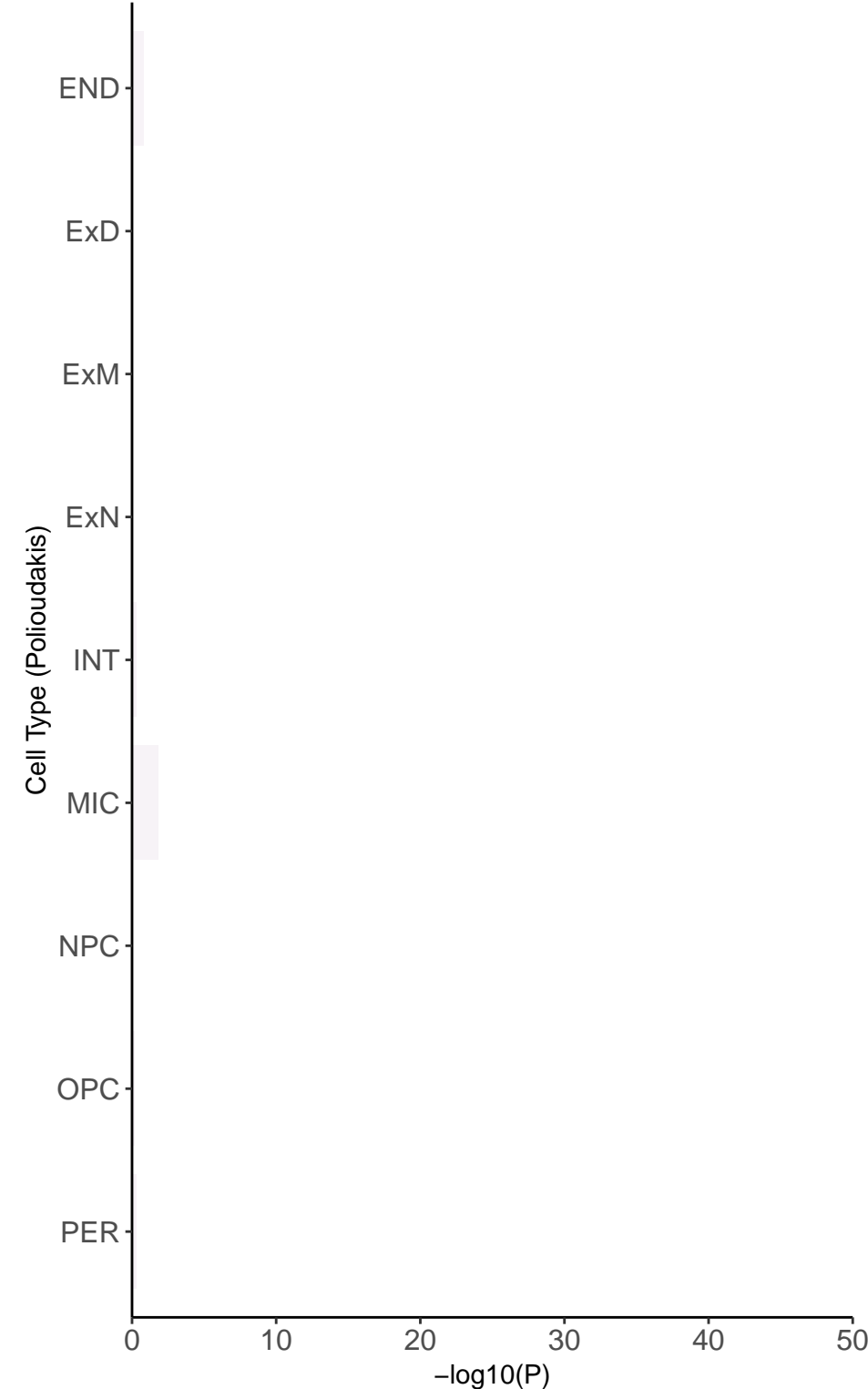
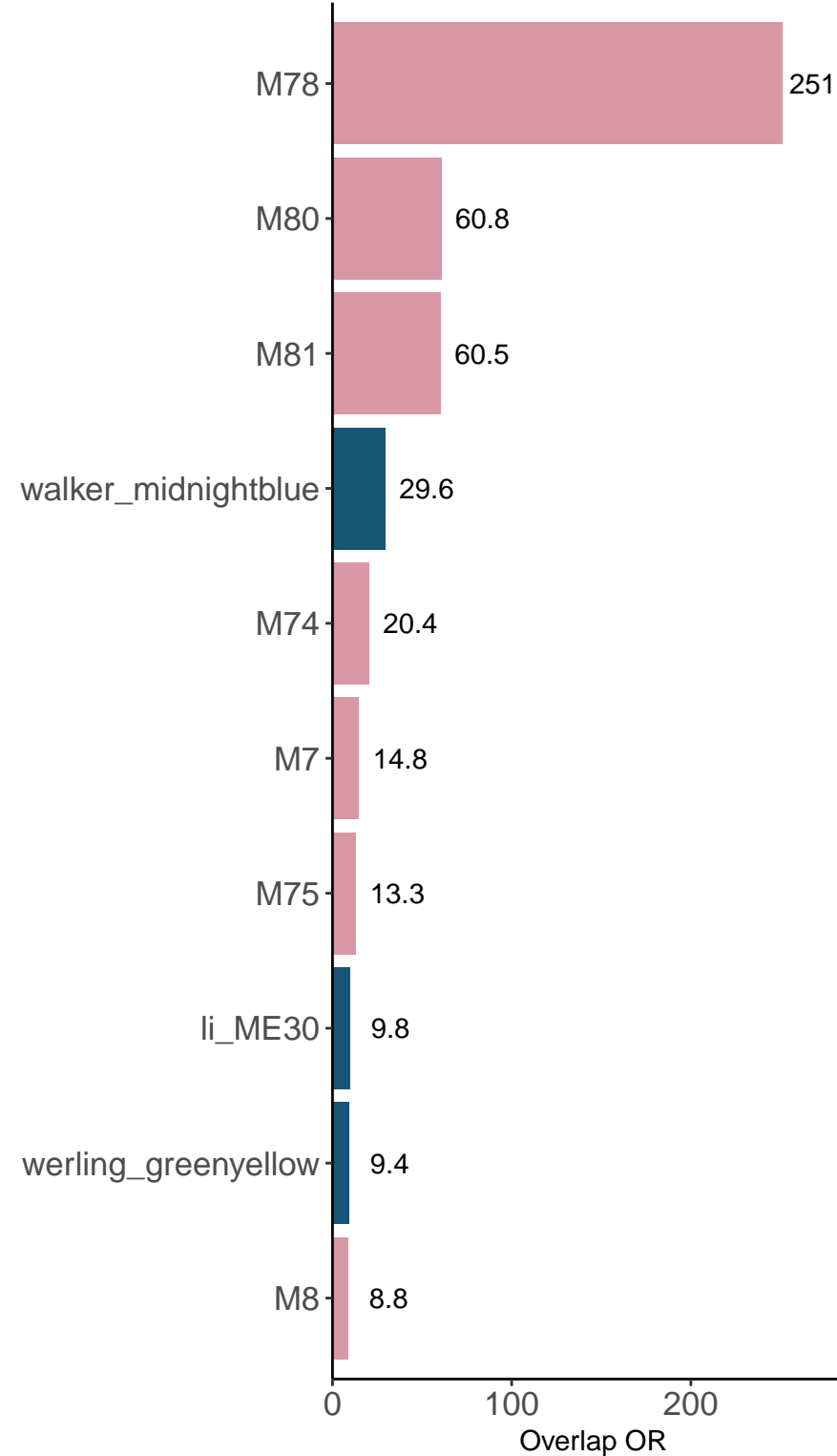
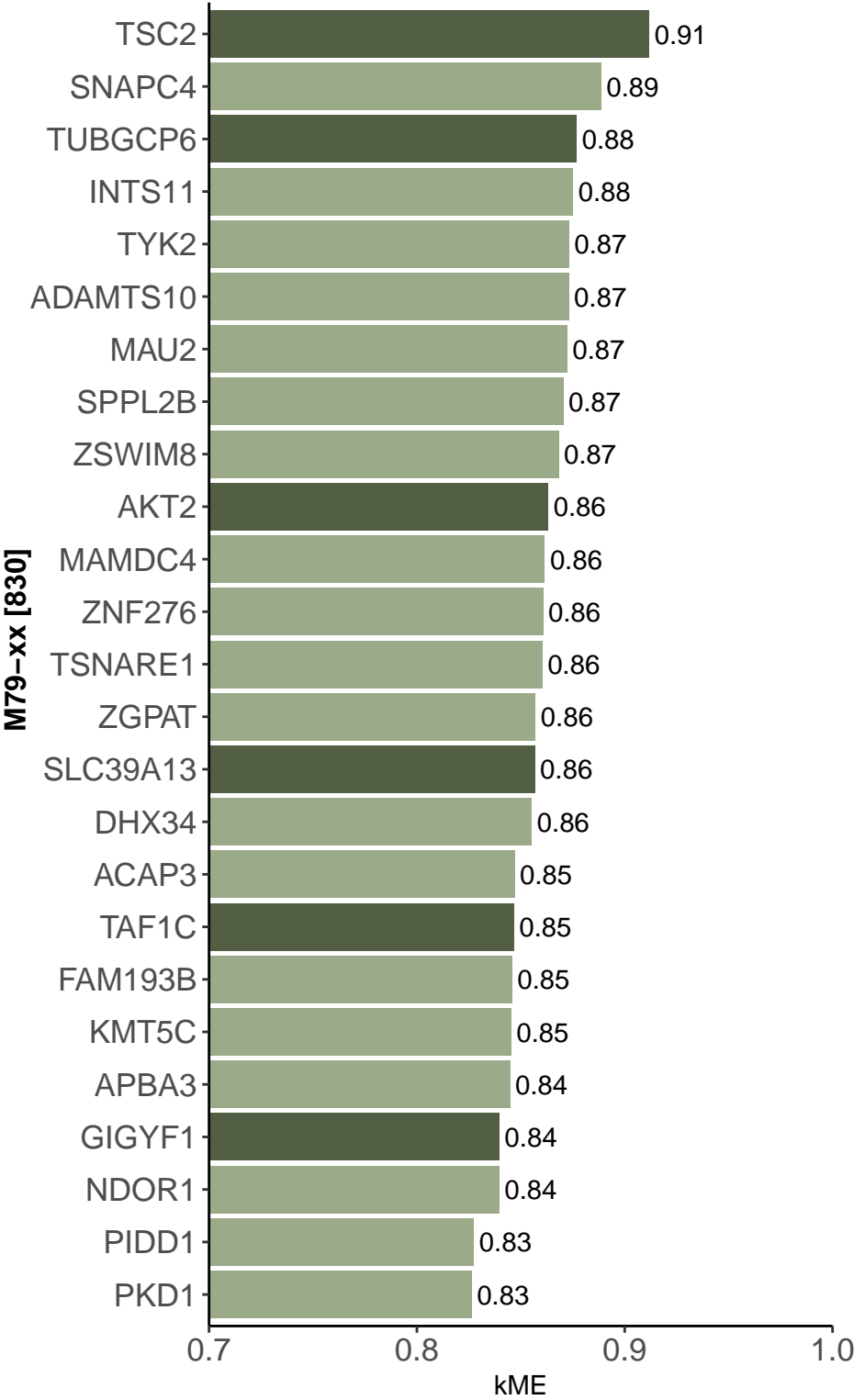


M79-xx [830]



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0018193	peptidyl–amino acid modification	1120	1.1e–05
2	GO:MF	GO:0030554	adenyl nucleotide binding	1523	4.7e–05
3	REAC	REAC:R–HSA–165181	Inhibition of TSC complex formation by PKB	3	1.4e–04
4	GO:MF	GO:0140640	catalytic activity, acting on a nucleic acid	574	1.4e–04
5	GO:BP	GO:0006302	double–strand break repair	286	1.4e–04
6	GO:MF	GO:0000166	nucleotide binding	1973	2.8e–04
7	GO:BP	GO:0006259	DNA metabolic process	973	3.3e–04
8	REAC	REAC:R–HSA–198323	AKT phosphorylates targets in the cytosol	14	4.2e–03
9	GO:CC	GO:0005815	microtubule organizing center	787	8.9e–03
10	GO:CC	GO:0099080	supramolecular complex	1150	9.1e–03
11	KEGG	KEGG:04150	mTOR signaling pathway	144	1.2e–02
12	GO:CC	GO:0099512	supramolecular fiber	812	1.3e–02
13	KEGG	KEGG:05165	Human papillomavirus infection	294	1.4e–02
14	REAC	REAC:R–HSA–5674400	Constitutive Signaling by AKT1 E17K in Cancer	25	1.5e–02
15	KEGG	KEGG:04140	Autophagy – animal	131	2.2e–02

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