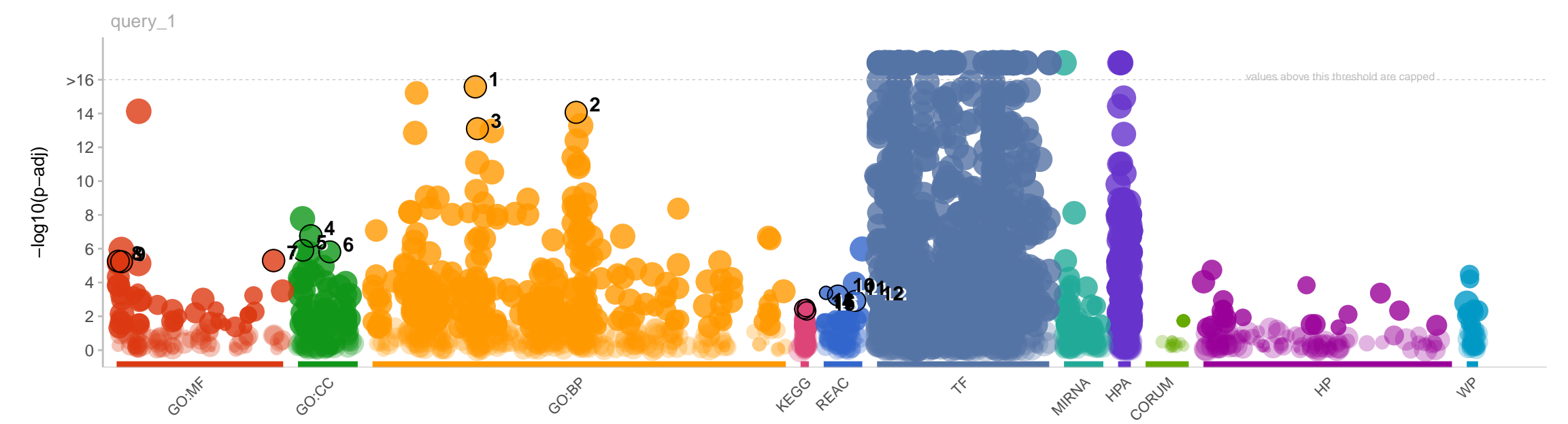
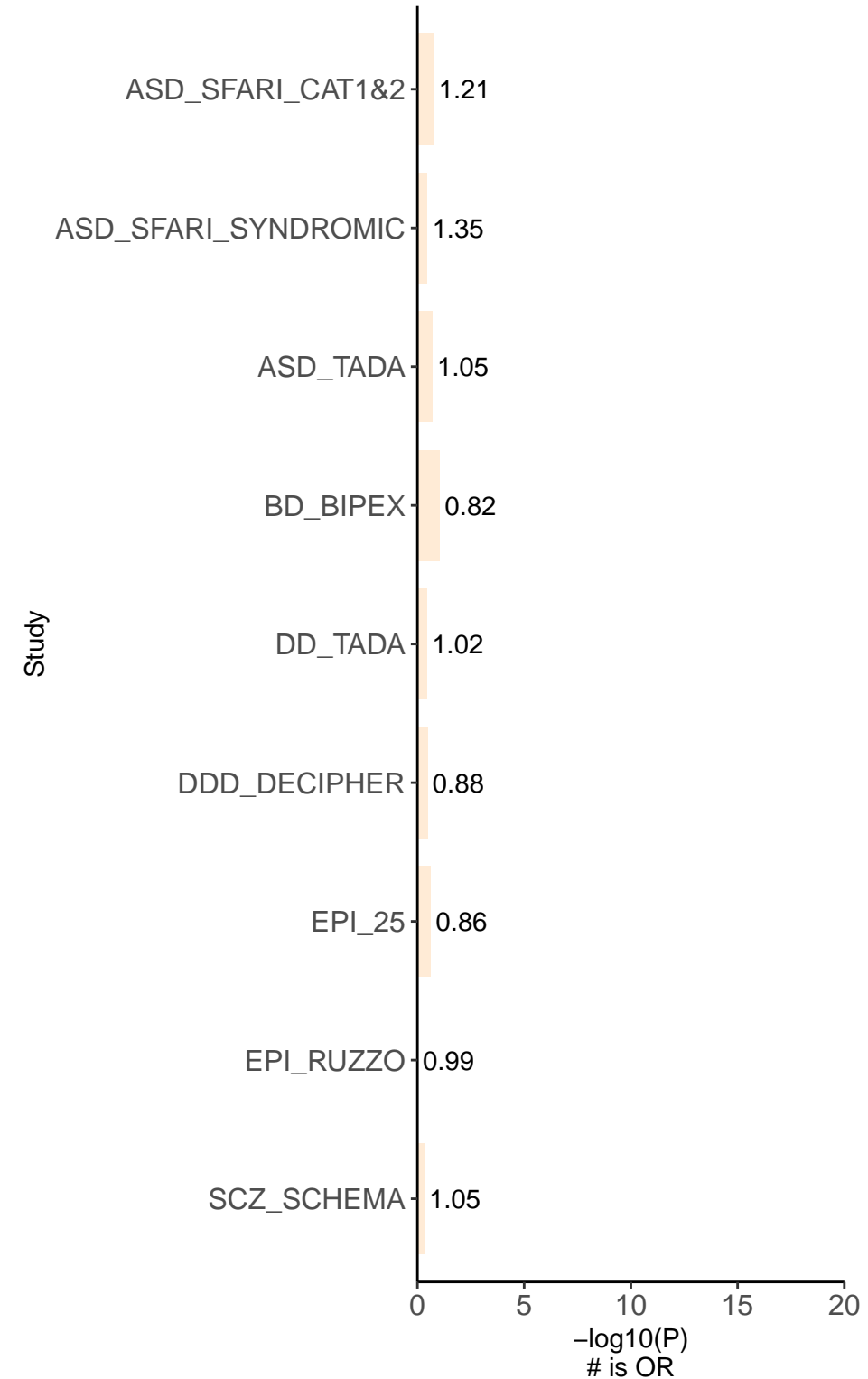
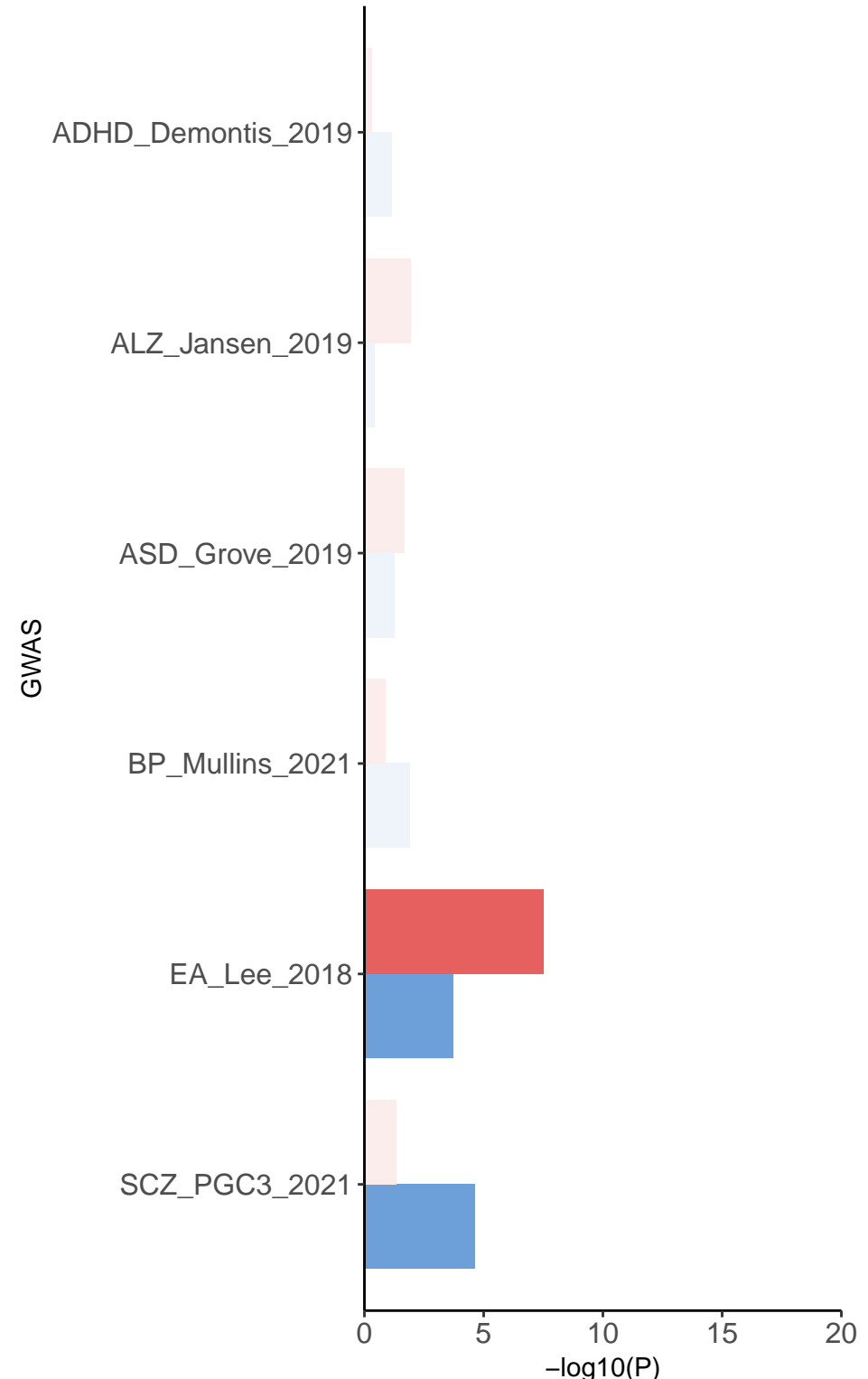
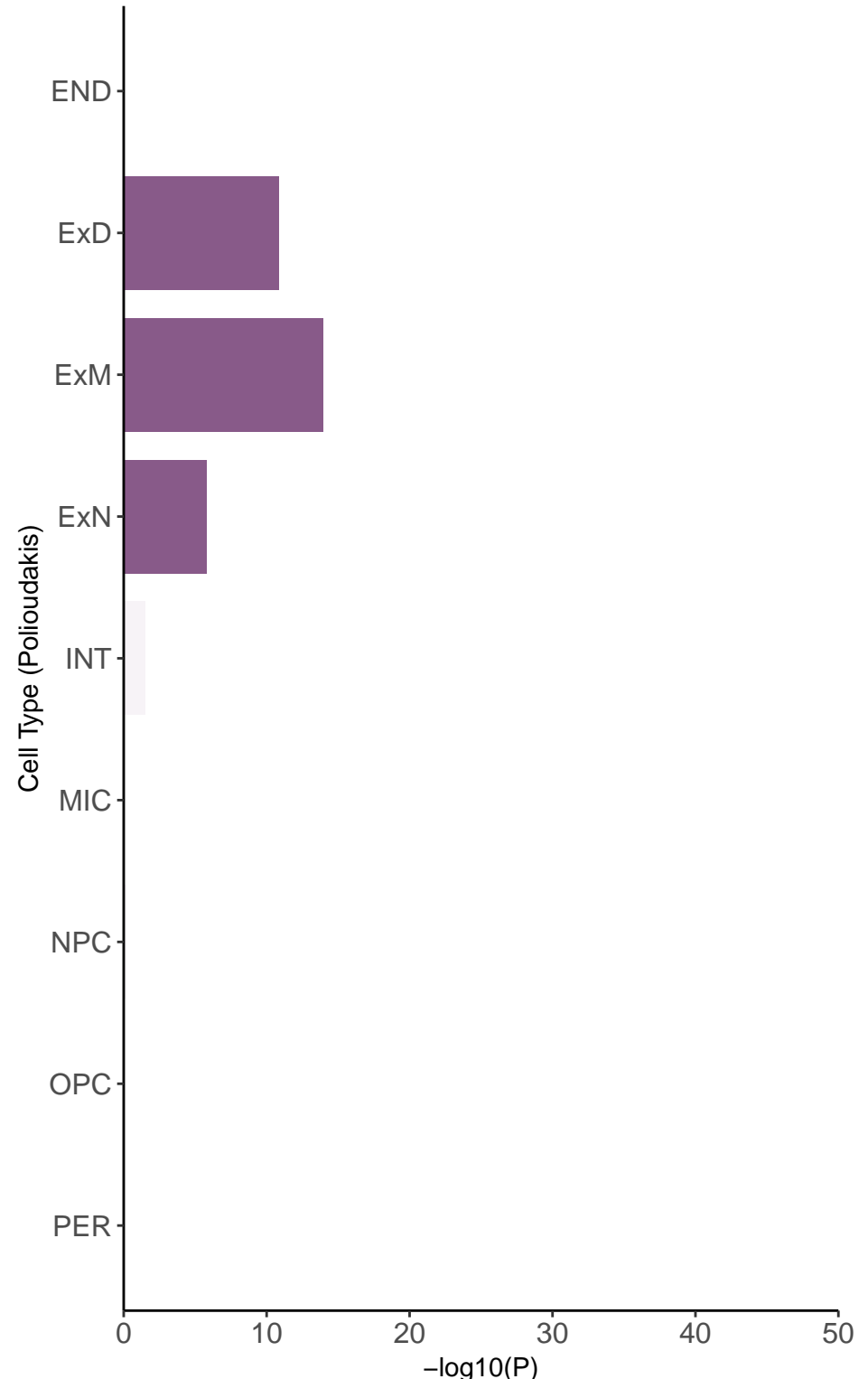
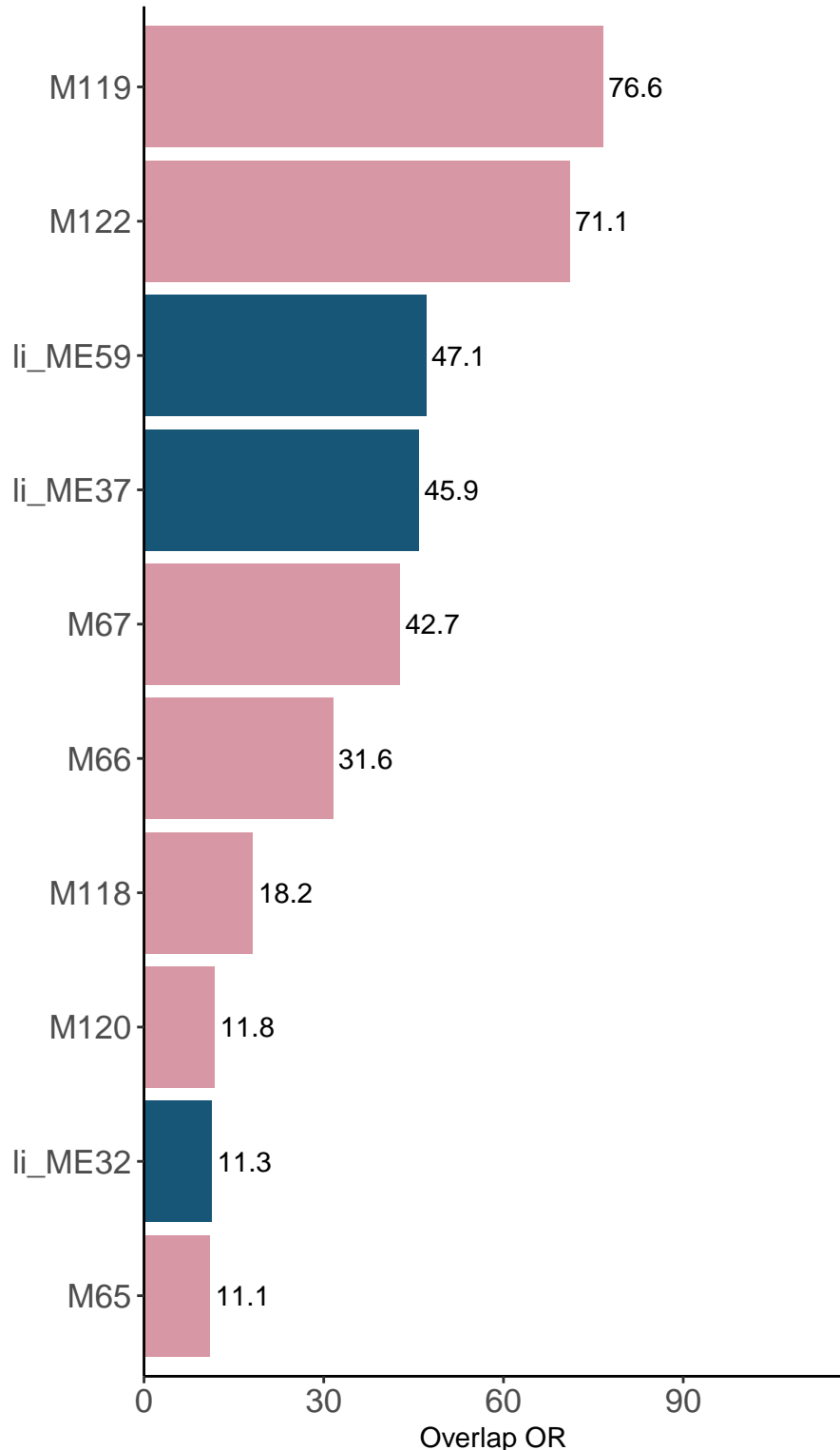
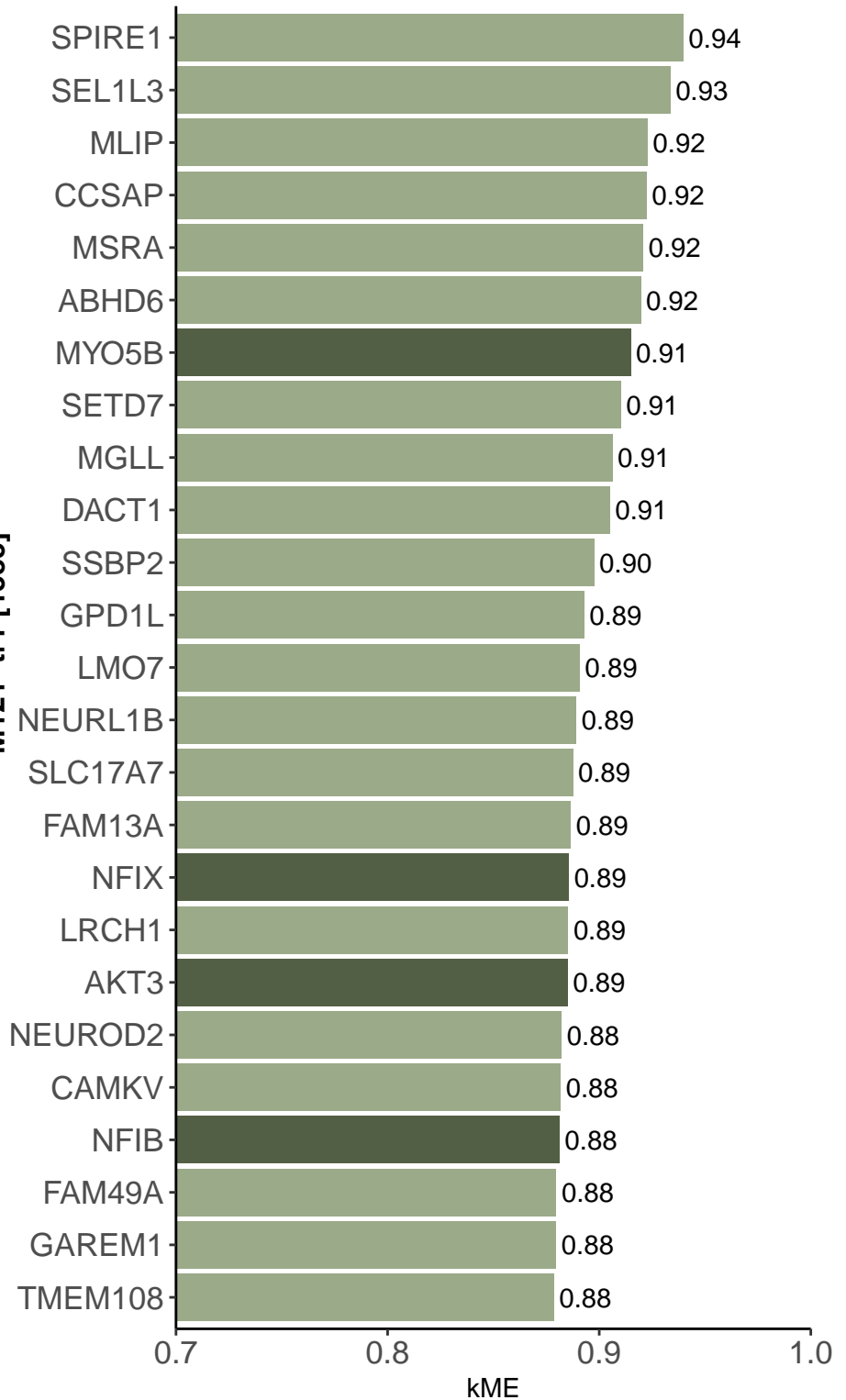


M121-tr1 [1383]



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0022008	neurogenesis	1548	2.6e-16
2	GO:BP	GO:0048699	generation of neurons	1343	8.6e-15
3	GO:BP	GO:0030182	neuron differentiation	1274	7.9e-14
4	GO:CC	GO:0030054	cell junction	1916	1.8e-07
5	GO:CC	GO:0005768	endosome	934	1.2e-06
6	GO:CC	GO:0045202	synapse	1219	1.5e-06
7	GO:MF	GO:0140110	transcription regulator activity	1632	5.0e-06
8	GO:MF	GO:0000981	DNA-binding transcription factor activity, RNA polymerase II-specific	1131	5.4e-06
9	GO:MF	GO:0003700	DNA-binding transcription factor activity	1177	5.9e-06
10	REAC	REAC:R-HSA-426048	Arachidonate production from DAG	5	4.1e-04
11	REAC	REAC:R-HSA-388396	GPCR downstream signalling	485	5.8e-04
12	REAC	REAC:R-HSA-372790	Signaling by GPCR	542	1.2e-03
13	KEGG	KEGG:04923	Regulation of lipolysis in adipocytes	50	3.2e-03
14	KEGG	KEGG:04022	cGMP-PKG signaling pathway	146	3.7e-03
15	KEGG	KEGG:05032	Morphine addiction	81	4.8e-03

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