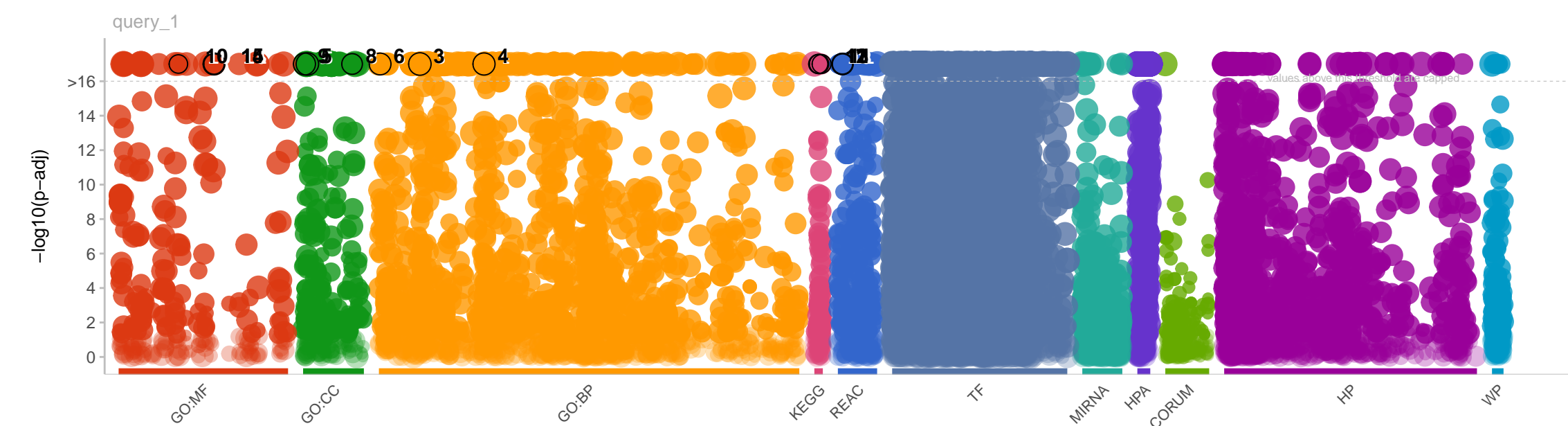
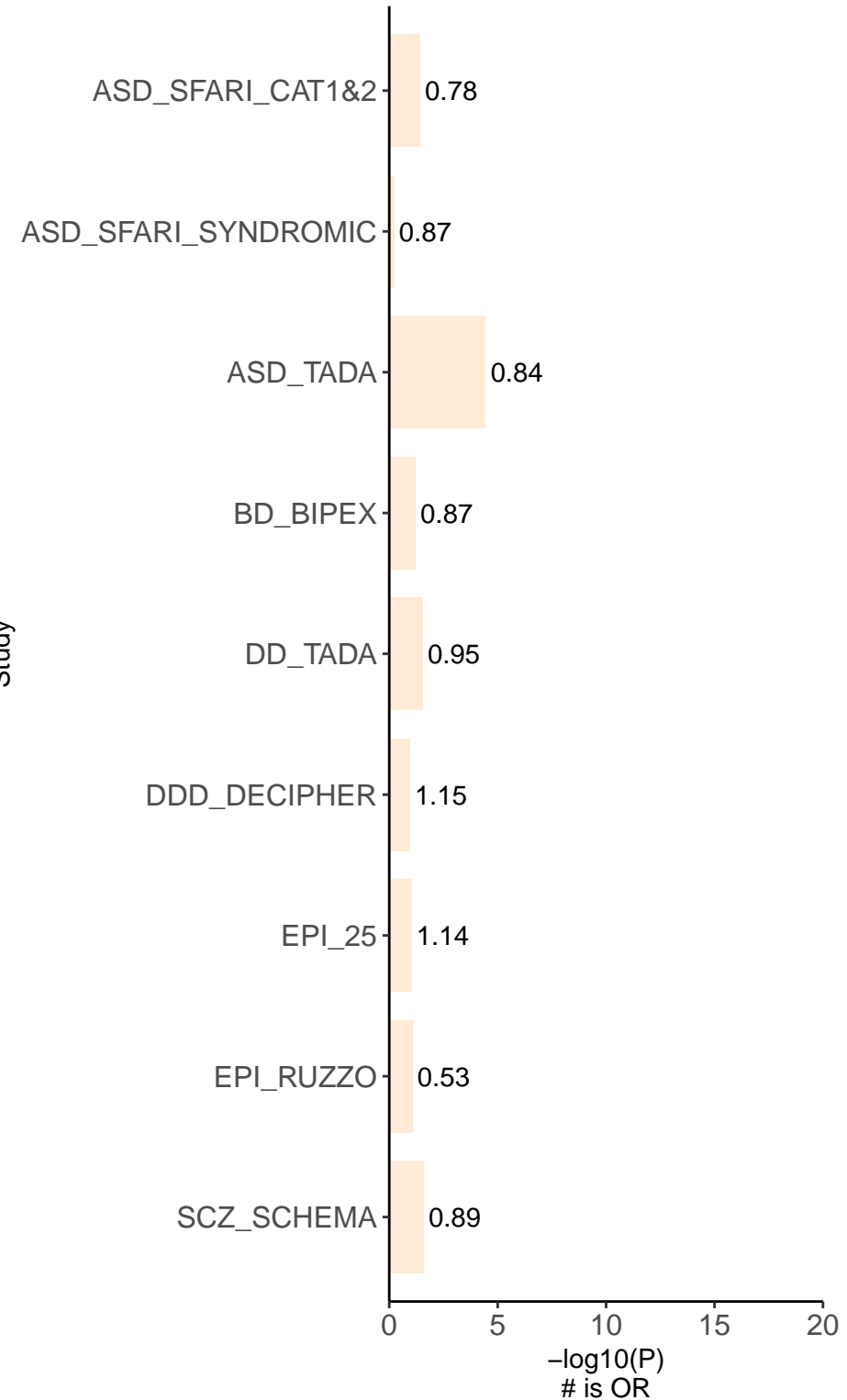
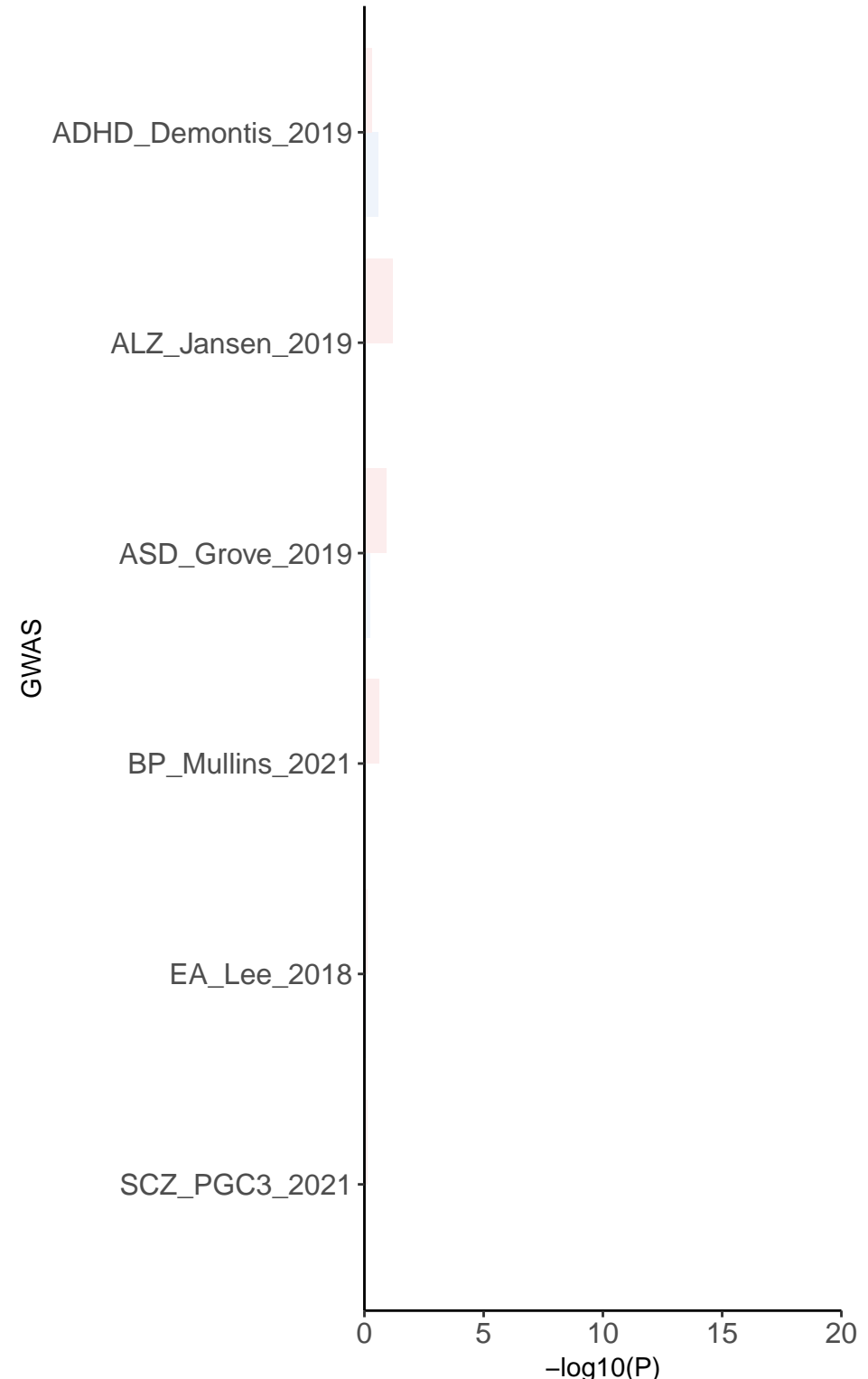
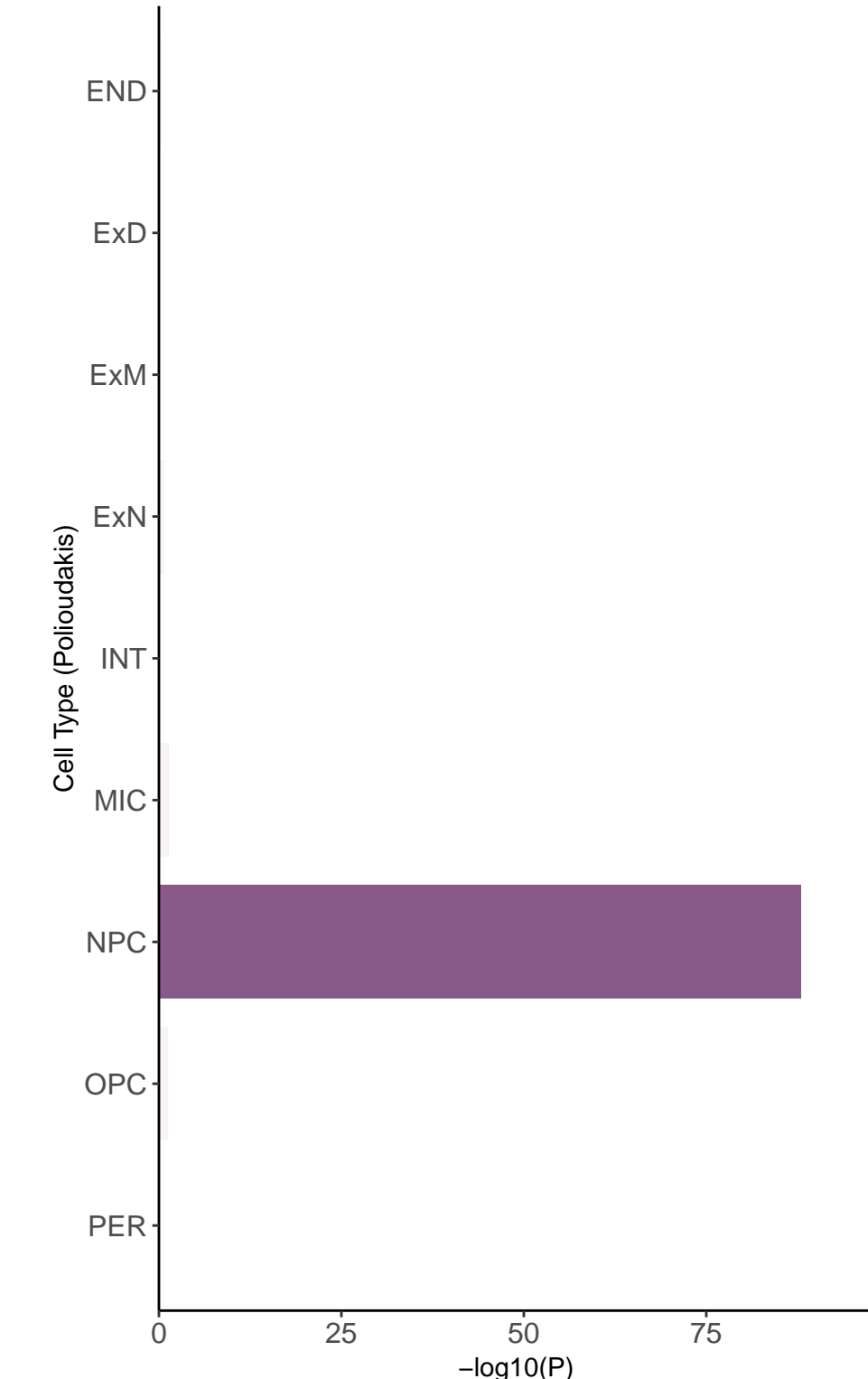
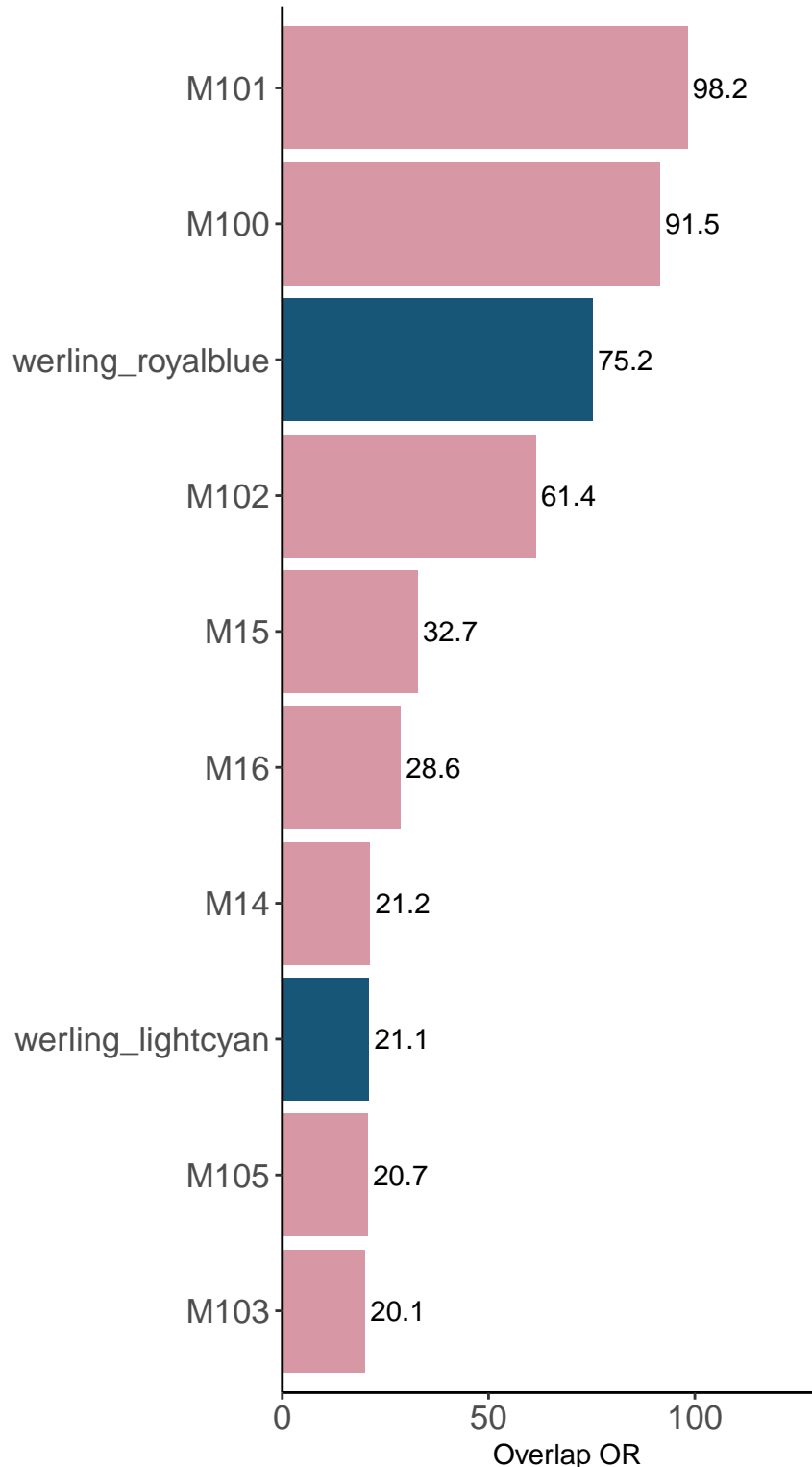
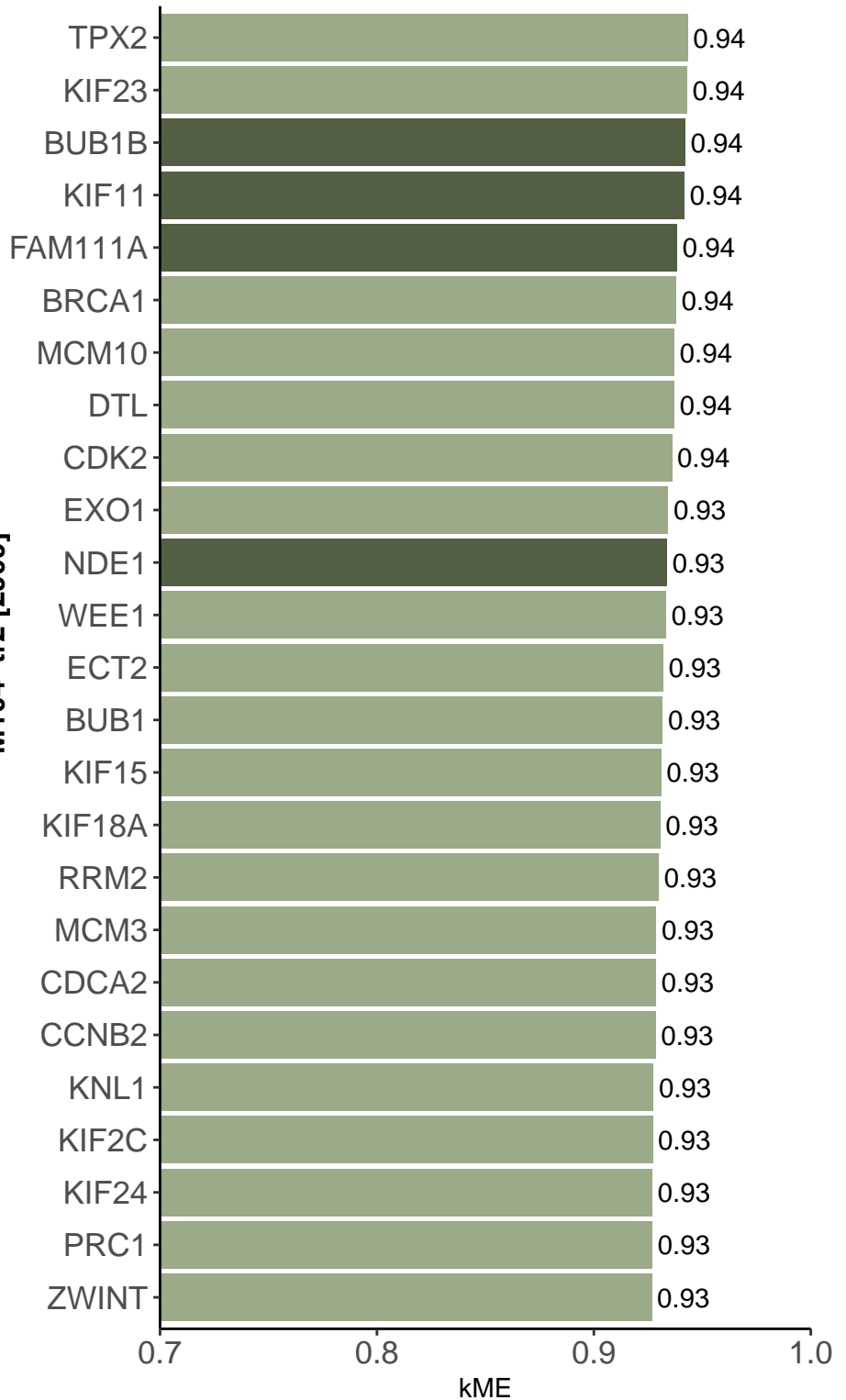


M104-tr2 [2909]



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-1640170	Cell Cycle	640	2.9e-132
2	REAC	REAC:R-HSA-69278	Cell Cycle, Mitotic	517	1.6e-124
3	GO:BP	GO:0007049	cell cycle	1655	1.7e-117
4	GO:BP	GO:0022402	cell cycle process	1173	1.6e-114
5	GO:CC	GO:0005694	chromosome	1638	1.9e-114
6	GO:BP	GO:0000278	mitotic cell cycle	878	9.8e-107
7	REAC	REAC:R-HSA-69620	Cell Cycle Checkpoints	275	2.3e-76
8	GO:CC	GO:0098687	chromosomal region	366	1.4e-60
9	GO:CC	GO:0000785	chromatin	1112	7.9e-57
10	GO:MF	GO:0030527	structural constituent of chromatin	78	2.0e-52
11	KEGG	KEGG:05322	Systemic lupus erythematosus	101	4.7e-38
12	KEGG	KEGG:04110	Cell cycle	123	3.1e-36
13	KEGG	KEGG:04613	Neutrophil extracellular trap formation	149	2.5e-32
14	GO:MF	GO:0046982	protein heterodimerization activity	293	3.8e-27
15	GO:MF	GO:0046983	protein dimerization activity	938	1.2e-21

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