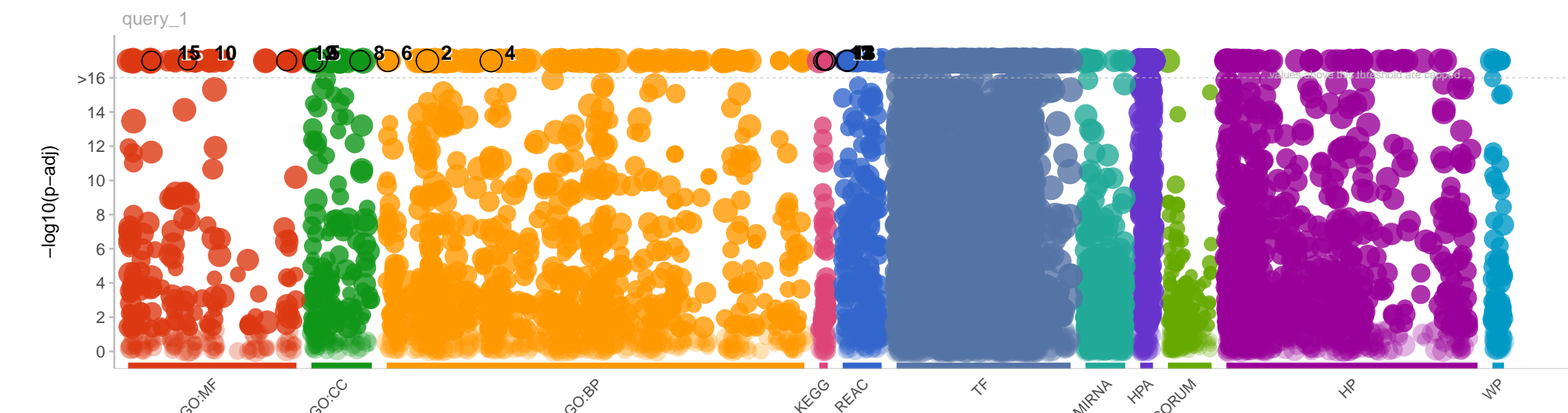
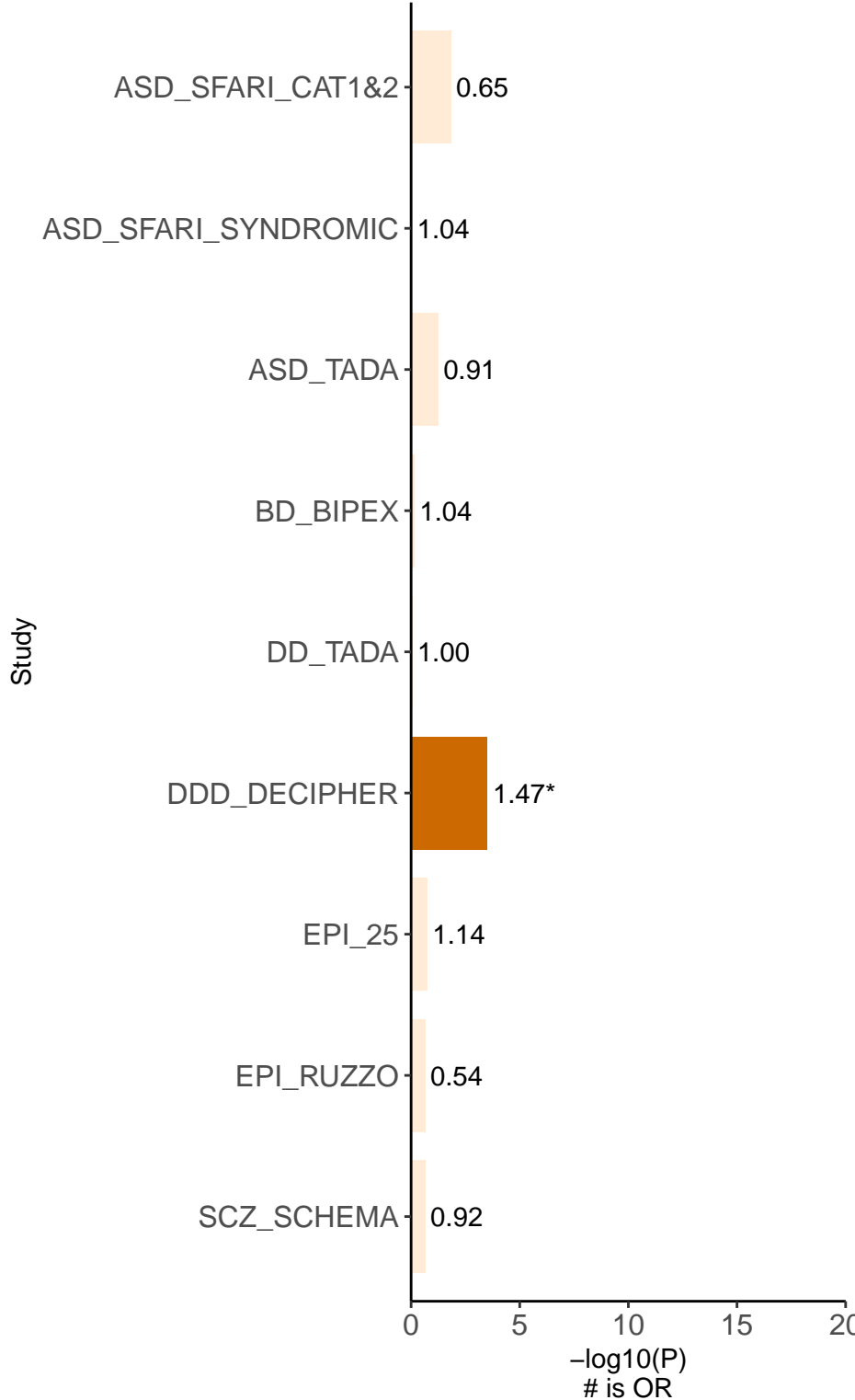
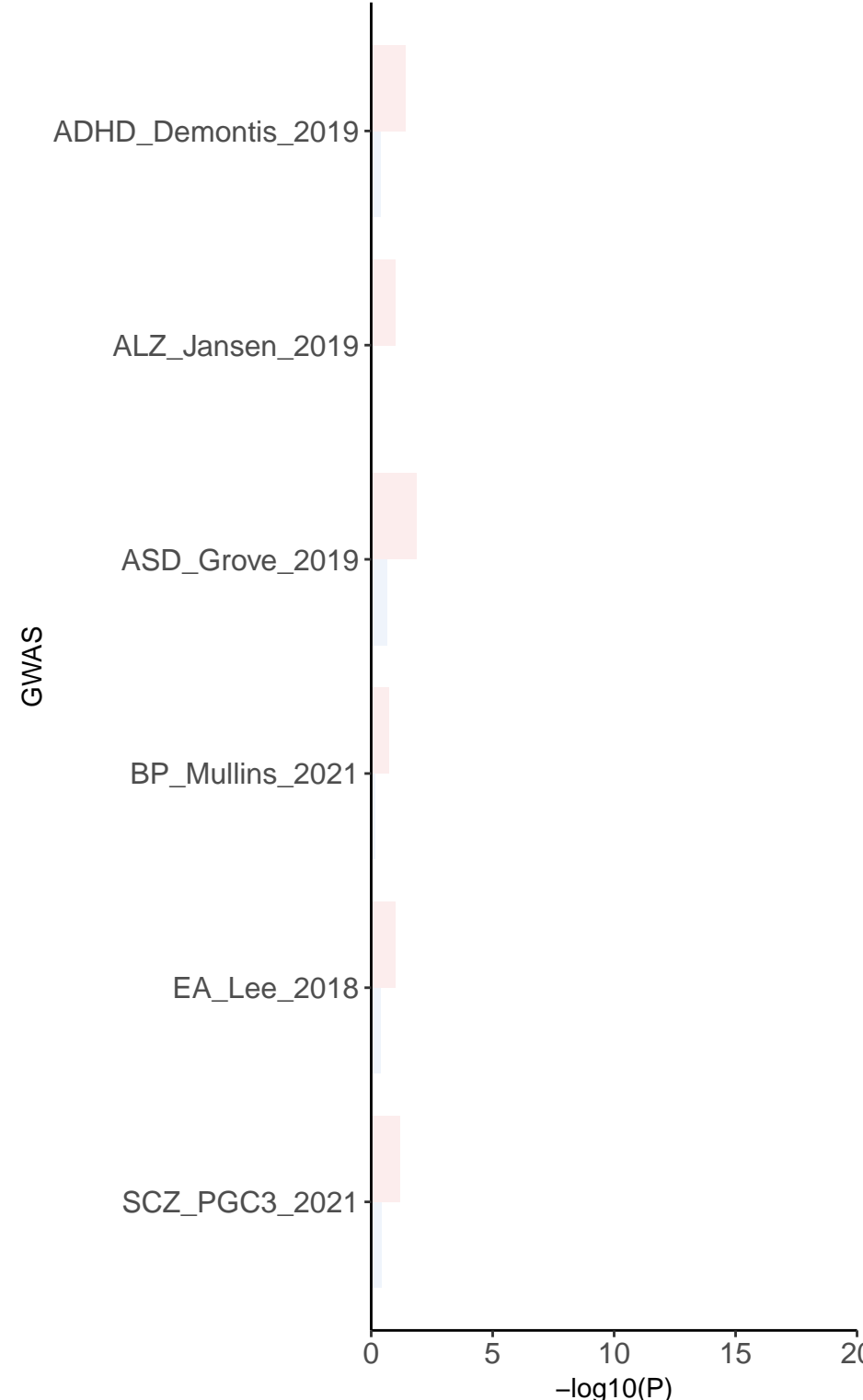
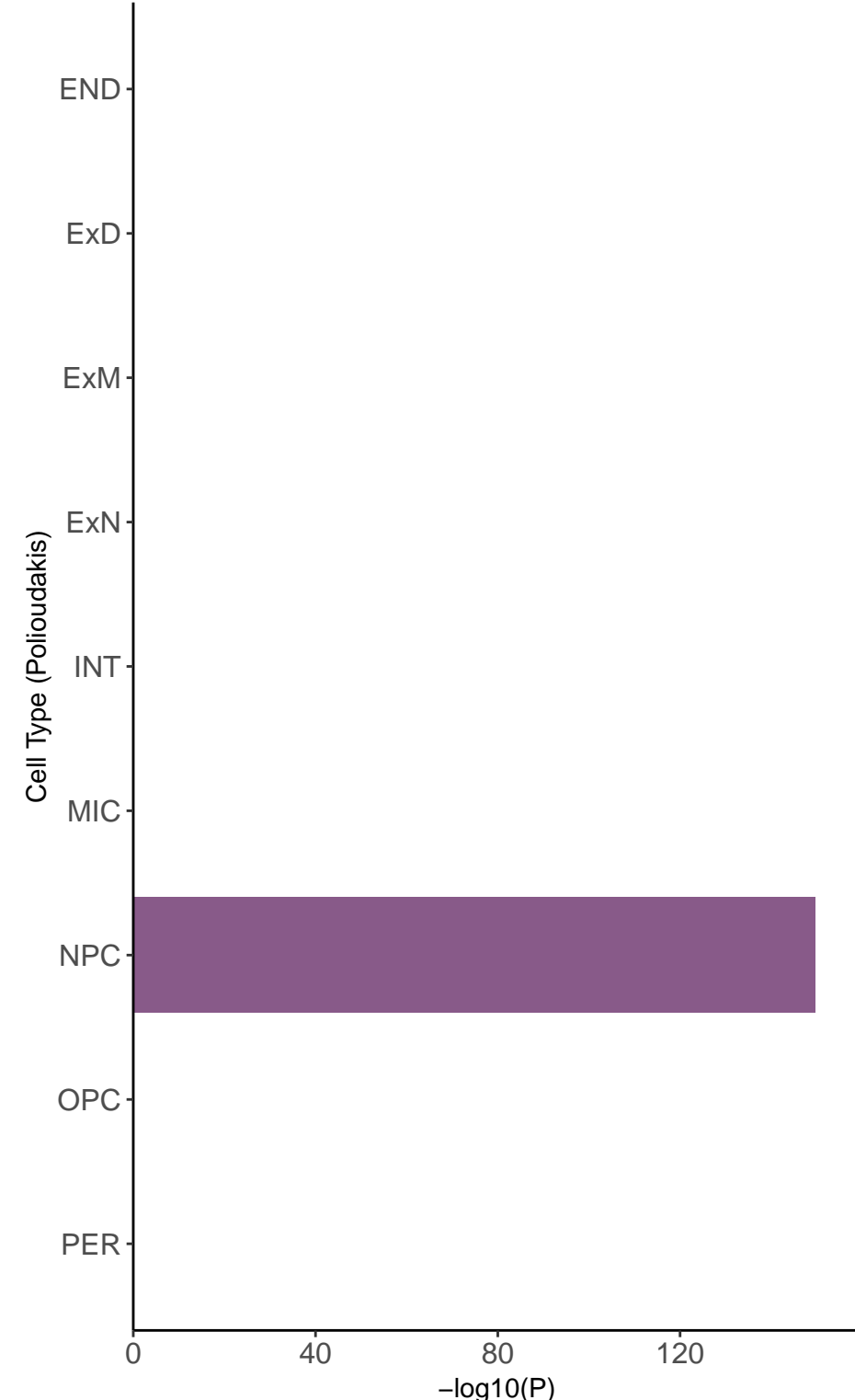
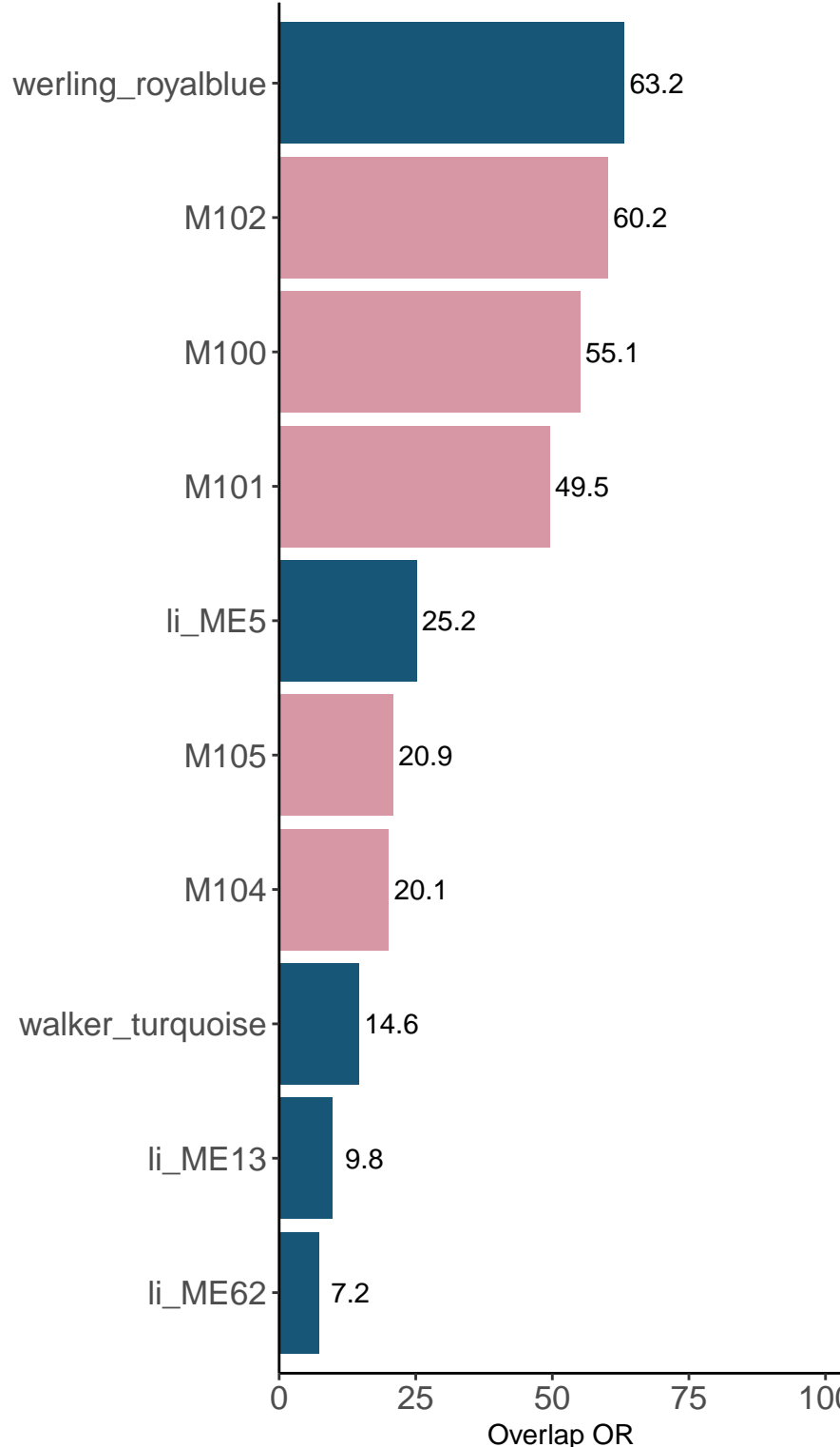
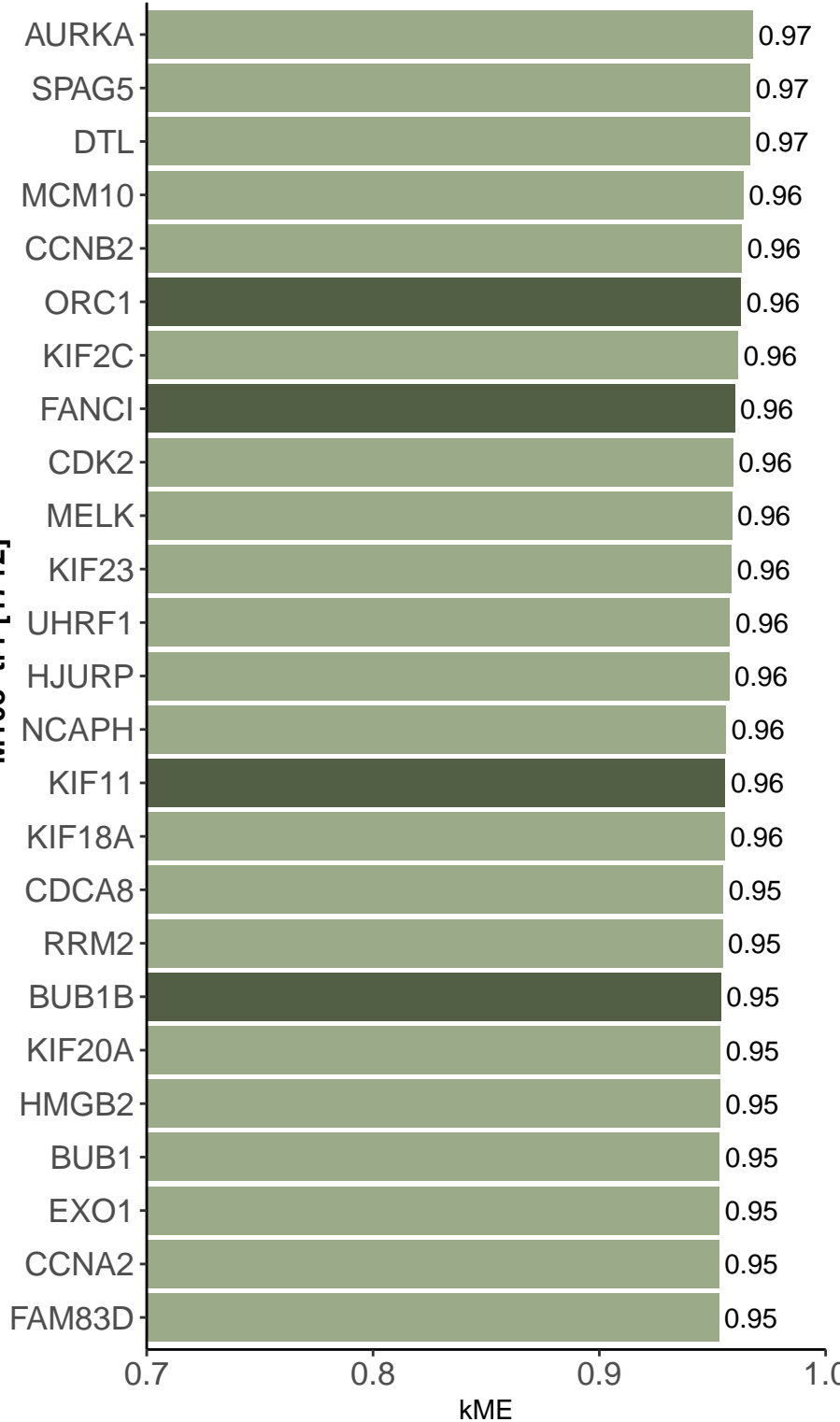


M103-tr1 [1712]



id	source	term_id	term_name	term_size	p_value
1	REAC	REAC:R-HSA-1640170	Cell Cycle	640	5.0e-169
2	GO:BP	GO:0007049	cell cycle	1655	3.1e-151
3	REAC	REAC:R-HSA-69278	Cell Cycle, Mitotic	517	2.6e-149
4	GO:BP	GO:0022402	cell cycle process	1173	6.5e-146
5	GO:CC	GO:0005694	chromosome	1638	1.3e-137
6	GO:BP	GO:0000278	mitotic cell cycle	878	1.3e-130
7	REAC	REAC:R-HSA-69620	Cell Cycle Checkpoints	275	4.1e-90
8	GO:CC	GO:0098687	chromosomal region	366	6.8e-90
9	GO:CC	GO:0000775	chromosome, centromeric region	235	1.6e-66
10	GO:MF	GO:0030527	structural constituent of chromatin	78	1.8e-55
11	KEGG	KEGG:04110	Cell cycle	123	1.5e-41
12	GO:MF	GO:0140097	catalytic activity, acting on DNA	223	1.0e-39
13	KEGG	KEGG:05322	Systemic lupus erythematosus	101	2.4e-36
14	KEGG	KEGG:05034	Alcoholism	154	6.4e-33
15	GO:MF	GO:0008094	ATP-dependent activity, acting on DNA	114	8.0e-30

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