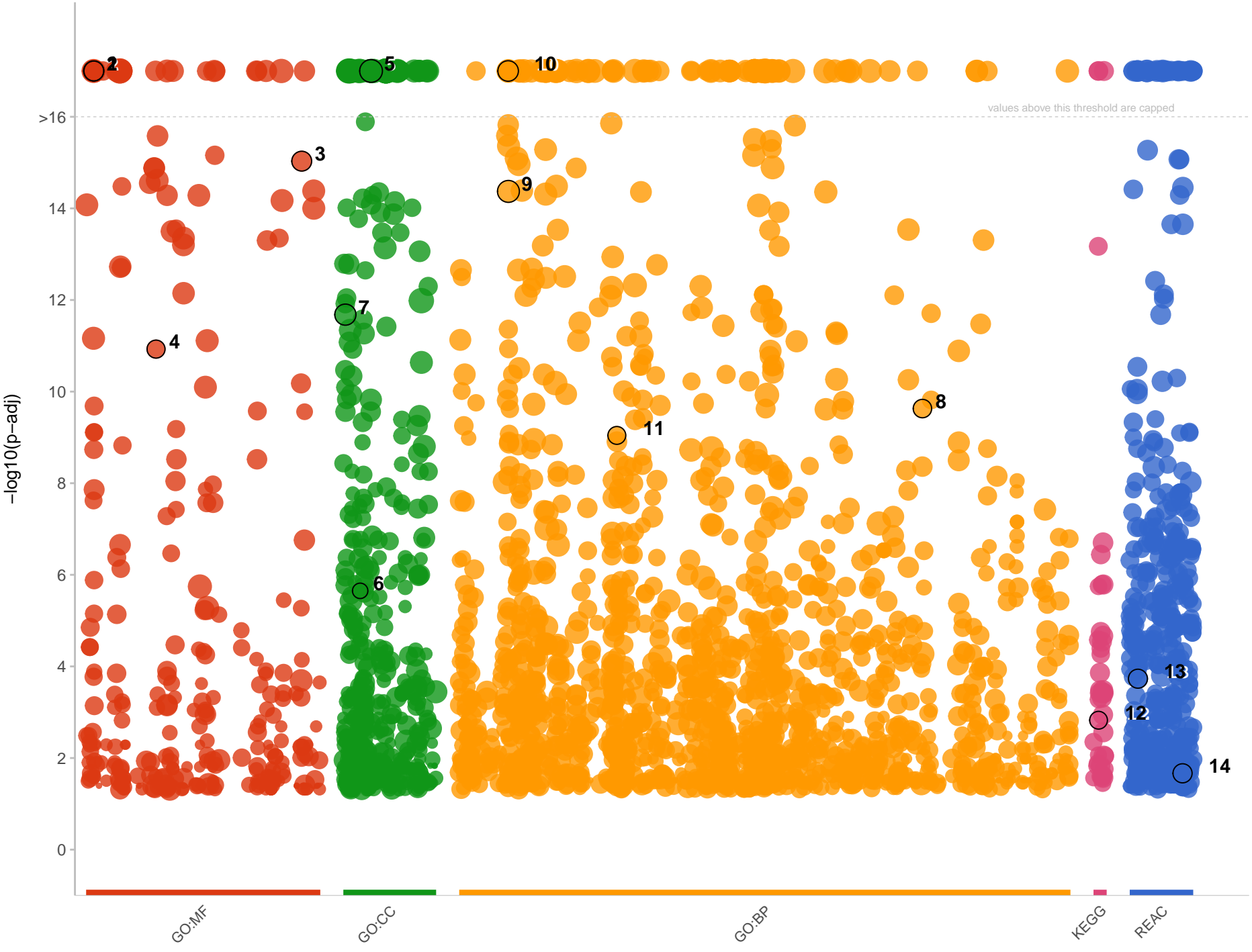
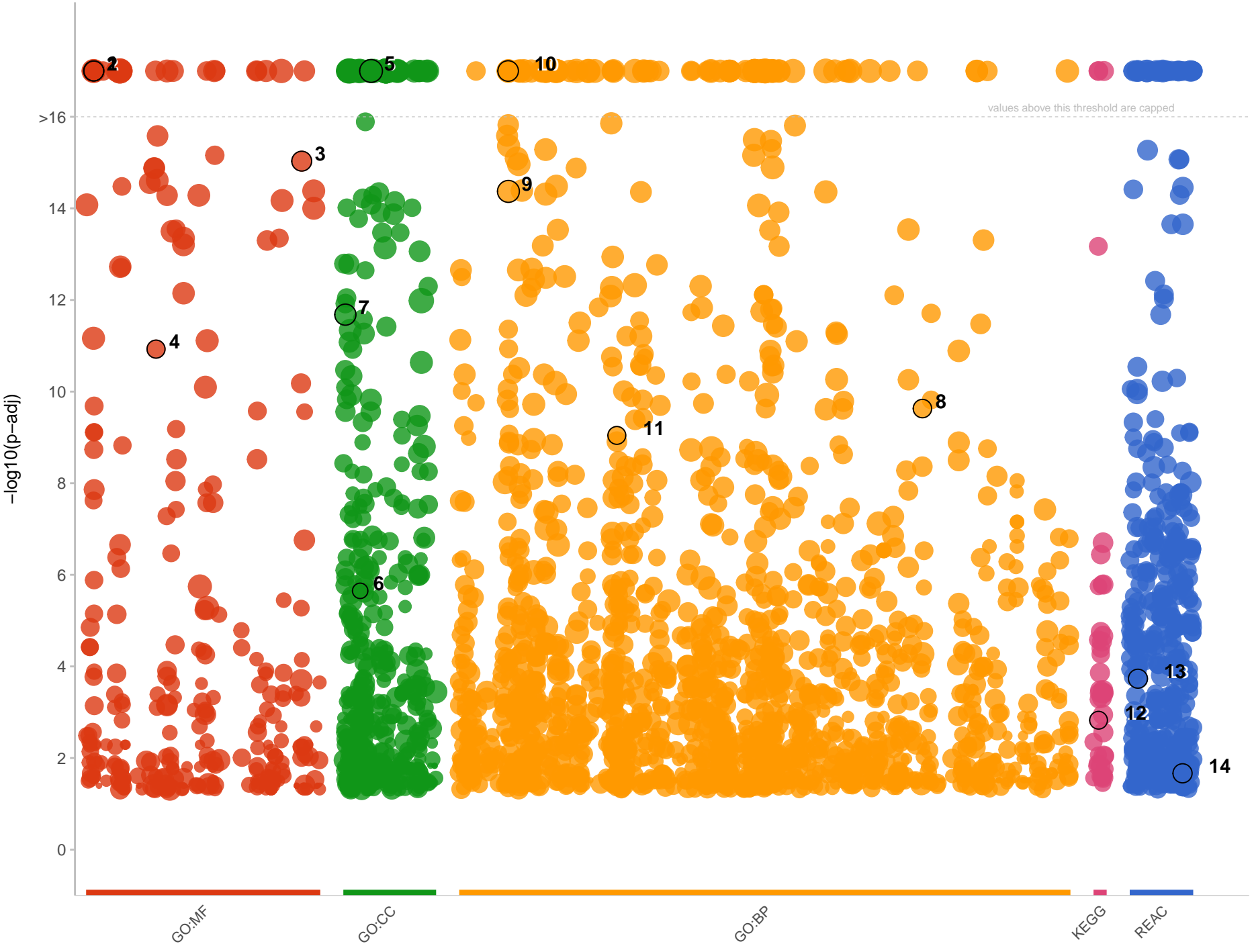


query\_1



id	source	term_id	term_name	term_size	p_value
1	GO:MF	GO:0003712	transcription coregulator activity	527	2.8e-32
2	GO:MF	GO:0003682	chromatin binding	603	5.1e-24
3	GO:MF	GO:0140297	DNA-binding transcription factor binding	490	9.3e-16
4	GO:MF	GO:0016922	nuclear receptor binding	138	1.2e-11
5	GO:CC	GO:0031981	nuclear lumen	5524	1.6e-79
6	GO:CC	GO:0016514	SWI/SNF complex	30	2.2e-06
7	GO:CC	GO:0000785	chromatin	1371	2.1e-12
8	GO:BP	GO:0141193	nuclear receptor-mediated signaling pathway	179	2.3e-10
9	GO:BP	GO:0006357	regulation of transcription by RNA polymerase II	2572	4.3e-15
10	GO:BP	GO:0006325	chromatin organization	843	7.3e-18
11	GO:BP	GO:0030518	nuclear receptor-mediated steroid hormone signaling pathway	123	9.0e-10
12	KEGG	KEGG:03082	ATP-dependent chromatin remodeling	116	1.5e-03
13	REAC	REAC:R-HSA-4839726	Chromatin organization	271	1.9e-04
14	REAC	REAC:R-HSA-9006931	Signaling by Nuclear Receptors	290	2.2e-02

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