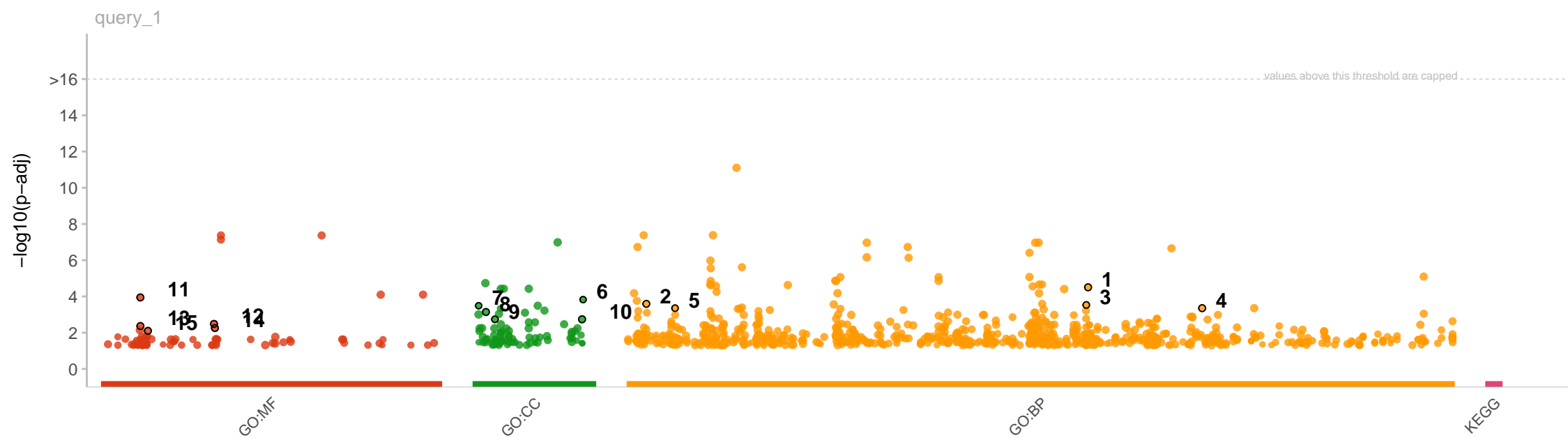


id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0060485	mesenchyme development	316	3.1e-05
2	GO:BP	GO:0002040	sprouting angiogenesis	128	2.6e-04
3	GO:BP	GO:0060421	positive regulation of heart growth	33	3.0e-04
4	GO:BP	GO:0090596	sensory organ morphogenesis	288	4.4e-04
5	GO:BP	GO:0003148	outflow tract septum morphogenesis	24	4.4e-04
6	GO:CC	GO:0170014	ankyrin-1 complex	8	1.5e-04
7	GO:CC	GO:0005579	membrane attack complex	7	3.2e-04
8	GO:CC	GO:0005912	adherens junction	188	7.2e-04
9	GO:CC	GO:0016342	catenin complex	31	1.8e-03
10	GO:CC	GO:0150034	distal axon	279	1.8e-03
11	GO:MF	GO:0005125	cytokine activity	238	1.1e-04
12	GO:MF	GO:0019838	growth factor binding	134	3.3e-03
13	GO:MF	GO:0005126	cytokine receptor binding	278	4.3e-03
14	GO:MF	GO:0019955	cytokine binding	145	5.6e-03
15	GO:MF	GO:0008083	growth factor activity	162	7.9e-03



id	source	term_id	term_name	term_size	p_value
1	GO:BP	GO:0060485	mesenchyme development	316	3.1e-05
2	GO:BP	GO:0002040	sprouting angiogenesis	128	2.6e-04
3	GO:BP	GO:0060421	positive regulation of heart growth	33	3.0e-04
4	GO:BP	GO:0090596	sensory organ morphogenesis	288	4.4e-04
5	GO:BP	GO:0003148	outflow tract septum morphogenesis	24	4.4e-04
6	GO:CC	GO:0170014	ankyrin-1 complex	8	1.5e-04
7	GO:CC	GO:0005579	membrane attack complex	7	3.2e-04
8	GO:CC	GO:0005912	adherens junction	188	7.2e-04
9	GO:CC	GO:0016342	catenin complex	31	1.8e-03
10	GO:CC	GO:0150034	distal axon	279	1.8e-03
11	GO:MF	GO:0005125	cytokine activity	238	1.1e-04
12	GO:MF	GO:0019838	growth factor binding	134	3.3e-03
13	GO:MF	GO:0005126	cytokine receptor binding	278	4.3e-03
14	GO:MF	GO:0019955	cytokine binding	145	5.6e-03
15	GO:MF	GO:0008083	growth factor activity	162	7.9e-03