

id	source	term_id	term_name	term_size	intersection_size	p_value
1	GO:BP	GO:0048731	system development	1535	78	8.5e−08
2	GO:BP	GO:0007275	multicellular organism development	2447	106	1.5e−07
3	GO:BP	GO:0048856	anatomical structure development	2839	117	2.2e−07
4	GO:BP	GO:0048468	cell development	1455	74	3.0e−07
5	GO:BP	GO:0032502	developmental process	3011	120	1.0e−06
6	GO:BP	GO:0030154	cell differentiation	1780	83	1.3e−06
7	GO:BP	GO:0048869	cellular developmental process	1813	83	3.1e−06
8	GO:BP	GO:0071840	cellular component organization or biogenesis	3341	127	5.2e−06
9	GO:BP	GO:0016043	cellular component organization	3160	122	5.2e−06
10	GO:BP	GO:0007399	nervous system development	1184	60	2.7e−05
11	GO:BP	GO:0048523	negative regulation of cellular process	1345	64	9.1e−05
12	GO:BP	GO:0048477	oogenesis	588	37	1.1e−04
13	GO:BP	GO:0007281	germ cell development	724	42	1.4e−04
14	GO:BP	GO:0022412	cellular process involved in reproduction in multicellular organism	895	48	1.7e−04
15	GO:BP	GO:0048513	animal organ development	1254	60	2.1e−04
16	GO:BP	GO:0032501	multicellular organismal process	3855	135	2.7e−04
17	GO:BP	GO:0007292	female gamete generation	638	38	3.1e−04
18	GO:BP	GO:0009653	anatomical structure morphogenesis	1587	70	3.7e−04
19	GO:BP	GO:0009888	tissue development	1015	51	5.0e−04
20	GO:BP	GO:0040007	growth	451	30	6.4e−04
21	GO:BP	GO:0022414	reproductive process	1275	59	8.6e−04
22	GO:BP	GO:0007276	gamete generation	929	47	1.3e−03
23	GO:BP	GO:0009987	cellular process	8640	244	1.4e−03
24	GO:BP	GO:0000902	cell morphogenesis	572	34	1.5e−03
25	GO:BP	GO:0016477	cell migration	326	24	1.6e−03
26	GO:BP	GO:0051128	regulation of cellular component organization	742	40	1.9e−03
27	GO:BP	GO:0003006	developmental process involved in reproduction	943	47	1.9e−03
28	GO:BP	GO:0048699	generation of neurons	896	45	2.7e−03
29	GO:BP	GO:0048609	multicellular organismal reproductive process	957	47	2.9e−03
30	GO:BP	GO:0000904	cell morphogenesis involved in differentiation	486	30	3.0e−03
31	GO:BP	GO:0060429	epithelium development	929	46	3.0e−03
32	GO:BP	GO:0022008	neurogenesis	964	47	3.5e−03
33	GO:BP	GO:0030182	neuron differentiation	823	42	4.1e−03
34	GO:BP	GO:0010256	endomembrane system organization	212	18	4.8e−03
35	GO:BP	GO:0048519	negative regulation of biological process	1701	70	4.9e−03
36	GO:BP	GO:0007154	cell communication	2117	82	6.1e−03
37	GO:BP	GO:0048589	developmental growth	377	25	6.2e−03
38	GO:BP	GO:0048666	neuron development	643	35	7.3e−03
39	GO:BP	GO:0050794	regulation of cellular process	4030	134	7.4e−03
40	GO:BP	GO:0065007	biological regulation	4779	153	7.8e−03
41	GO:BP	GO:0023051	regulation of signaling	969	46	9.3e−03
42	GO:BP	GO:0010646	regulation of cell communication	973	46	1.0e−02
43	GO:BP	GO:0072499	photoreceptor cell axon guidance	32	7	1.1e−02
44	GO:BP	GO:0000003	reproduction	1510	63	1.1e−02
45	GO:BP	GO:0009966	regulation of signal transduction	838	41	1.5e−02
46	GO:BP	GO:0048870	cell motility	397	25	1.5e−02
47	GO:BP	GO:0035295	tube development	724	37	1.6e−02
48	GO:BP	GO:0050896	response to stimulus	3255	112	1.7e−02
49	GO:BP	GO:0048522	positive regulation of cellular process	1733	69	1.8e−02
50	GO:BP	GO:0044085	cellular component biogenesis	1667	67	1.8e−02
51	GO:BP	GO:0023057	negative regulation of signaling	402	25	1.9e−02
52	GO:BP	GO:0010648	negative regulation of cell communication	404	25	2.1e−02
53	GO:BP	GO:0150063	visual system development	405	25	2.1e−02
54	GO:BP	GO:0001654	eye development	405	25	2.1e−02
55	GO:BP	GO:0048880	sensory system development	405	25	2.1e−02
56	GO:BP	GO:0009968	negative regulation of signal transduction	381	24	2.3e−02
57	GO:BP	GO:0022607	cellular component assembly	1451	60	2.5e−02
58	GO:BP	GO:0006897	endocytosis	217	17	2.6e−02
59	GO:BP	GO:0007423	sensory organ development	492	28	3.0e−02
60	GO:BP	GO:0002064	epithelial cell development	290	20	3.2e−02
61	GO:BP	GO:0032504	multicellular organism reproduction	1335	56	3.4e−02
62	GO:BP	GO:0050793	regulation of developmental process	693	35	3.6e−02
63	GO:BP	GO:0048518	positive regulation of biological process	1941	74	3.7e−02
64	GO:BP	GO:0051641	cellular localization	1277	54	3.8e−02
65	GO:BP	GO:0022604	regulation of cell morphogenesis	84	10	4.0e−02
66	GO:BP	GO:0097435	supramolecular fiber organization	297	20	4.5e−02
67	GO:BP	GO:0007010	cytoskeleton organization	702	35	4.7e−02
68	GO:BP	GO:0050789	regulation of biological process	4441	141	4.8e−02
69	GO:CC	GO:0005737	cytoplasm	5042	167	5.4e−06
70	GO:CC	GO:0005622	intracellular anatomical structure	7212	216	1.3e−05
71	GO:CC	GO:0045177	apical part of cell	194	19	4.0e−05
72	GO:CC	GO:0043228	non−membrane−bounded organelle	1939	80	6.4e−05
73	GO:CC	GO:0043232	intracellular non−membrane−bounded organelle	1939	80	6.4e−05
74	GO:CC	GO:0110165	cellular anatomical entity	10590	277	6.4e−04
75	GO:CC	GO:0030054	cell junction	602	33	1.5e−03
76	GO:CC	GO:0005829	cytosol	1119	48	9.6e−03
77	GO:CC	GO:0043226	organelle	6265	183	1.2e−02
78	GO:CC	GO:0098793	presynapse	239	17	1.2e−02
79	GO:CC	GO:0043229	intracellular organelle	6167	180	1.6e−02
80	GO:CC	GO:0031594	neuromuscular junction	99	10	2.5e−02
81	GO:CC	GO:0045202	synapse	442	24	3.2e−02
82	GO:CC	GO:0045172	germline ring canal	23	5	3.5e−02
83	GO:CC	GO:0035324	female germline ring canal	13	4	3.8e−02
84	GO:CC	GO:0101031	chaperone complex	13	4	3.8e−02
85	GO:CC	GO:0005700	polytene chromosome	239	16	4.1e−02
86	GO:CC	GO:0045169	fusome	37	6	4.3e−02
87	GO:MF	GO:0005488	binding	6892	224	1.6e−10
88	GO:MF	GO:0005515	protein binding	3670	138	2.0e−07
89	GO:MF	GO:0043167	ion binding	2855	105	3.6e−04
90	GO:MF	GO:1901363	heterocyclic compound binding	3131	107	9.2e−03
91	GO:MF	GO:0097159	organic cyclic compound binding	3156	107	1.3e−02
92	GO:MF	GO:0097367	carbohydrate derivative binding	1240	51	3.4e−02
93	GO:MF	GO:0005509	calcium ion binding	205	15	4.8e−02
94	MIRNA	MIRNA:dme−miR−318−3p	dme−miR−318−3p	1	1	5.0e−02
95	TF	TF:M12357	Factor: BEAF32; motif: NTATCGATAKTTNN	1085	45	1.9e−02
96	TF	TF:M07142	Factor: BEAF−32A; motif: NNNMNAMTATCGATA	1025	43	2.2e−02