

GO Term ID	Definition	Size
GO:0007476	imaginal disc-derived wing morphogenesis	237
GO:0007411	axon guidance	152
GO:0046331	lateral inhibition	144
GO:0006909	phagocytosis	142
GO:0045944	positive regulation of transcription from RNA polymerase II promoter	137
GO:0007163	establishment or maintenance of cell polarity	133
GO:0000398	mRNA splicing, via spliceosome	132
GO:0006259	DNA metabolic process	126
GO:0008283	cell proliferation	120
GO:0034645	cellular macromolecule biosynthetic process	120
GO:0044087	regulation of cellular component biogenesis	117
GO:0009968	negative regulation of signal transduction	115
GO:0048813	dendrite morphogenesis	113
GO:0007067	mitosis	111
GO:0042127	regulation of cell proliferation	102
GO:0009967	positive regulation of signal transduction	96
GO:0010608	posttranscriptional regulation of gene expression	96
GO:0007059	chromosome segregation	95
GO:0008340	determination of adult lifespan	95
GO:0048589	developmental growth	95
GO:0006955	immune response	94
GO:0007626	locomotory behavior	93
GO:0070925	organelle assembly	93
GO:0010927	cellular component assembly involved in morphogenesis	92
GO:0009953	dorsal/ventral pattern formation	89
GO:0007420	brain development	88
GO:0002682	regulation of immune system process	84
GO:0007155	cell adhesion	84
GO:0008360	regulation of cell shape	84
GO:0051656	establishment of organelle localization	84
GO:0001751	compound eye photoreceptor cell differentiation	83
GO:0007298	border follicle cell migration	82
GO:0007015	actin filament organization	80
GO:0042981	regulation of apoptotic process	79
GO:0034613	cellular protein localization	78
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GO:0060341	regulation of cellular localization	77
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GO:0044281	small molecule metabolic process	75
GO:0048638	regulation of developmental growth	75
GO:0000022	mitotic spindle elongation	73
GO:0030534	adult behavior	73
GO:0006403	RNA localization	72
GO:0007519	skeletal muscle tissue development	72
GO:0045664	regulation of neuron differentiation	72
GO:0051298	centrosome duplication	72
GO:1901564	organonitrogen compound metabolic process	72
GO:0000122	negative regulation of transcription from RNA polymerase II promoter	71

GO:0051130	positive regulation of cellular component organization	71
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GO:0007167	enzyme linked receptor protein signaling pathway	70
GO:0007286	spermatid development	70
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GO:0015031	protein transport	70
GO:0022416	chaeta development	70
GO:0032535	regulation of cellular component size	70
GO:0050808	synapse organization	70
GO:0051129	negative regulation of cellular component organization	70
GO:0031399	regulation of protein modification process	69
GO:0044265	cellular macromolecule catabolic process	69
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GO:0010627	regulation of intracellular protein kinase cascade	68
GO:0060249	anatomical structure homeostasis	68
GO:0060562	epithelial tube morphogenesis	68
GO:0016570	histone modification	67
GO:0034654	nucleobase-containing compound biosynthetic process	67
GO:0042742	defense response to bacterium	67
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GO:0016458	gene silencing	63
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GO:0006508	proteolysis	56
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