	p.value	term_goid	term_category	term_name
"1"	8.51915602785e-07	"GO:0008285"	"b"	"negative regulation of cell proliferation"
"2"	2.50967213811e-06	"GO:0050678"	"b"	"regulation of epithelial cell proliferation"
"3"	5.42809452878e-06	"GO:0010631"	"b"	"epithelial cell migration"
"4"	3.38871359311e-05	"GO:0051272"	"b"	"positive regulation of cellular component movement"
"5"	0.000167881717208	"GO:0051174"	"b"	"regulation of phosphorus metabolic process"
"6"	0.000194891886017	"GO:0010594"	"b"	"regulation of endothelial cell migration"
"7"	0.000276258790947	"GO:0051271"	"b"	"negative regulation of cellular component movement"
"8"	0.000361820663383	"GO:0010634"	"b"	"positive regulation of epithelial cell migration"
"9"	0.000563372042009	"GO:0045834"	"b"	"positive regulation of lipid metabolic process"
"10"	0.000588294583857	"GO:0007346"	"b"	"regulation of mitotic cell cycle"
"11"	0.000803167451922	"GO:0005001"	"m"	"transmembrane receptor protein tyrosine phosphatase activity"
"12"	0.000918598763477	"GO:0044409"	"b"	"entry into host"
"13"	0.000918598763477	"GO:0051806"	"b"	"entry into cell of other organism involved in symbiotic interaction"
"14"	0.000929985798515	"GO:0051347"	"b"	"positive regulation of transferase activity"
"15"	0.00101996046229	"GO:0031344"	"b"	"regulation of cell projection organization"
"16"	0.00108811371902	"GO:0007169"	"b"	"transmembrane receptor protein tyrosine kinase signaling pathway"
"17"	0.0010905087542	"GO:1902531"	"b"	"regulation of intracellular signal transduction"
"18"	0.00114745289731	"GO:0010647"	"b"	"positive regulation of cell communication"
"19"	0.00130326218452	"GO:0010942"	"b"	"positive regulation of cell death"
"20"	0.00134740258268	"GO:0090288"	"b"	"negative regulation of cellular response to growth factor stimulus"
"21"	0.00141464236711	"GO:0034394"	"b"	"protein localization to cell surface"
"22"	0.00151398813277	"GO:0006897"	"b"	"endocytosis"
"23"	0.00193487507885	"GO:0071675"	"b"	"regulation of mononuclear cell migration"
"24"	0.00198757789092	"GO:0048015"	"b"	"phosphatidylinositol-mediated signaling"
"25"	0.00214275150265	"GO:0044822"	"m"	"poly(A) RNA binding"
"26"	0.00237467834411	"GO:0010812"	"b"	"negative regulation of cell-substrate adhesion"
"27"	0.00241384544408	"GO:0000904"	"b"	"cell morphogenesis involved in differentiation"
"28"	0.00242230524078	"GO:0071363"	"b"	"cellular response to growth factor stimulus"

"29" 0.00242651114848 "GO:0046501"   "b"   "protoporphyrinogen IX metabolic process"
"30" 0.00247833772829 "GO:0006915"   "b"   "apoptotic process"
"31" 0.00288839807592 "GO:0032268"   "b"   "regulation of cellular protein metabolic process"
"32" 0.00303874933837 "GO:0009967"   "b"   "positive regulation of signal transduction"
"33" 0.00316310293427 "GO:0010648"   "b"   "negative regulation of cell communication"
"34" 0.00318417120962 "GO:0031325"   "b"   "positive regulation of cellular metabolic process"
"35" 0.00320289920517 "GO:0022008"   "b"   "neurogenesis"
"36" 0.00372558165944 "GO:0008284"   "b"   "positive regulation of cell proliferation"
"37" 0.00373918527065 "GO:2000181"   "b"   "negative regulation of blood vessel morphogenesis"
"38" 0.00386417975349 "GO:1901342"   "b"   "regulation of vasculature development"
"39" 0.00419808077457 "GO:0030686"   "c"   "90S preribosome"
"40" 0.00437141053371 "GO:0045765"   "b"   "regulation of angiogenesis"
"41" 0.00457959481892 "GO:0030182"   "b"   "neuron differentiation"
"42" 0.00483230784723 "GO:0051246"   "b"   "regulation of protein metabolic process"
"43" 0.00484612791131 "GO:1900024"   "b"   "regulation of substrate adhesion-dependent cell spreading"
"44" 0.00529707851307 "GO:1901343"   "b"   "negative regulation of vasculature development"
"45" 0.00535362585296 "GO:0046850"   "b"   "regulation of bone remodeling"
"46" 0.00546231517039 "GO:0034660"   "b"   "ncRNA metabolic process"
"47" 0.0054816735841 "GO:0007411"   "b"   "axon guidance"
"48" 0.00572429584462 "GO:0043067"   "b"   "regulation of programmed cell death"
"49" 0.00611638467403 "GO:0003725"   "m"   "double-stranded RNA binding"
"50" 0.00633232990954 "GO:0033594"   "b"   "response to hydroxyisoflavone"
"51" 0.00633232990954 "GO:0003176"   "b"   "aortic valve development"
"52" 0.00633232990954 "GO:0019960"   "m"   "C-X3-C chemokine binding"
"53" 0.00637184260225 "GO:0050801"   "b"   "ion homeostasis"
"54" 0.00685734546832 "GO:0007265"   "b"   "Ras protein signal transduction"
"55" 0.00700692953809 "GO:0060284"   "b"   "regulation of cell development"
"56" 0.00729759870029 "GO:0015696"   "b"   "ammonium transport"
"57" 0.00748139666399 "GO:1900115"   "b"   "extracellular regulation of signal transduction"
"58" 0.00775682799707 "GO:0043202"   "c"   "lysosomal lumen"
"59" 0.00849840611391 "GO:0035296"   "b"   "regulation of tube diameter"
"60" 0.00850647660275 "GO:0032102"   "b"   "negative regulation of response to external stimulus"
"61" 0.00894107247166 "GO:0070670"   "b"   "response to interleukin-4"

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"62"  0.00933650742251 "GO:0080184"     "b"   "response to
phenylpropanoid"
"63"  0.00933650742251 "GO:0010044"     "b"   "response to aluminum ion"
"64"  0.00933650742251 "GO:0048050"     "b"   "post-embryonic eye
morphogenesis"
"65"  0.00933650742251 "GO:0007000"     "b"   "nucleolus organization"
"66"  0.00952957528632 "GO:0006023"     "b"   "aminoglycan biosynthetic
process"
"67"  0.00956856337942 "GO:0006812"     "b"   "cation transport"
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