

## Cluster 1

term	N-term	N-query	overlap	p value	domain
mitochondrion	1900	1800	420	5.9e-109	GO:CC
ribosomal subunit	200	1800	140	1.1e-108	GO:CC
mitochondrial envelope	840	1800	260	2.3e-99	GO:CC
mitochondrial membrane	780	1800	240	3.4e-92	GO:CC
mitochondrial inner membrane	530	1800	200	1e-91	GO:CC
organelle envelope	1300	1800	310	2.1e-91	GO:CC
organelle inner membrane	580	1800	200	1.3e-85	GO:CC
translation	720	1800	210	3.9e-76	GO:BP
cytosolic ribosome	130	1800	92	1.8e-75	GO:CC
cytoplasmic translation	160	1800	99	1.7e-70	GO:BP
parkinson disease	260	950	140	4.8e-67	KEGG
ribosome	430	1800	150	6.3e-64	GO:CC
large ribosomal subunit	130	1800	81	4.1e-60	GO:CC
aerobic respiration	200	1800	100	1.1e-59	GO:BP
huntington disease	300	950	140	2.3e-59	KEGG

## Cluster 2

term	N-term	N-query	overlap	p value	domain
nervous system development	2500	1700	430	1.3e-83	GO:BP
cell junction	2300	1700	370	1.9e-67	GO:CC
neuron development	1300	1700	270	1.2e-65	GO:BP
generation of neurons	1600	1700	310	1.4e-65	GO:BP
neuron projection development	1100	1700	250	2.1e-64	GO:BP
neuron projection	1500	1700	290	3.5e-64	GO:CC
synapse	1600	1700	300	4.9e-64	GO:CC
neurogenesis	1900	1700	330	2e-63	GO:BP
neuron differentiation	1600	1700	290	4e-62	GO:BP
cell projection	2600	1700	390	2e-58	GO:CC
plasma membrane bounded cell projection ...	1700	1700	290	1.1e-53	GO:BP
cell projection organization	1700	1700	290	1.9e-52	GO:BP
axon	800	1700	180	2.1e-51	GO:CC
somatodendritic compartment	1100	1700	220	7.1e-51	GO:CC
plasma membrane bounded cell projection	2500	1700	360	7.5e-51	GO:CC