

GO:1903306	regulation of nucleic acid-templated tra...	2221	43	28.78	0.00317	0.0101
GO:2001141	regulation of RNA biosynthetic process	2226	43	28.84	0.00331	0.0101
GO:0003352	regulation of cilium movement	7	2	0.09	0.00336	0.0101
GO:0014889	muscle atrophy	7	2	0.09	0.00336	0.0101
GO:0034351	negative regulation of glial cell apopto...	7	2	0.09	0.00336	0.0101
GO:2000112	regulation of cellular macromolecule bio...	2501	47	32.41	0.00350	0.0101
GO:0001754	eye photoreceptor cell differentiation	24	3	0.31	0.00354	0.0101
GO:0001570	vasculogenesis	50	4	0.65	0.00393	0.0108
GO:0051965	positive regulation of synapse assembly	25	3	0.32	0.00398	0.0108
GO:1901362	organic cyclic compound biosynthetic pro...	2660	49	34.47	0.00420	0.0108
GO:0032099	negative regulation of appetite	8	2	0.10	0.00444	0.0108
GO:0033210	leptin-mediated signaling pathway	8	2	0.10	0.00444	0.0108
GO:0034350	regulation of glial cell apoptotic proce...	8	2	0.10	0.00444	0.0108
GO:0007611	learning or memory	120	6	1.55	0.00467	0.0111
GO:0040018	positive regulation of multicellular org...	27	3	0.35	0.00497	0.0111
GO:0006355	regulation of transcription, DNA-templat...	2215	42	28.70	0.00527	0.0111
GO:0072659	protein localization to plasma membrane	164	7	2.13	0.00547	0.0111
GO:0001539	cilium or flagellum-dependent cell motil...	9	2	0.12	0.00566	0.0111
GO:0032096	negative regulation of response to food	9	2	0.12	0.00566	0.0111
GO:0043508	negative regulation of JUN kinase activi...	9	2	0.12	0.00566	0.0111
GO:0043567	regulation of insulin-like growth factor...	9	2	0.12	0.00566	0.0111
GO:0044320	cellular response to leptin stimulus	9	2	0.12	0.00566	0.0111
GO:0060285	cilium-dependent cell motility	9	2	0.12	0.00566	0.0111
GO:0010556	regulation of macromolecule biosynthetic...	2565	47	33.24	0.00579	0.0111
GO:0009391	post-embryonic development	89	5	1.15	0.00594	0.0111