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COURTINGS Security	GO:0032094	response to food	16	3	0.21	0.00106	0.0102
Commission Temperature Commission Co	GO:0032774	RNA biosynthetic process	2301	46	29.81	0.00109	0.0102
General State General Stat	GO:0007632	visual behavior	37	4	0.48	0.00128	0.0102
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CO-0019159 Peterocycle biosynthetic process 2594 49 33.61 0.00247 0.0107	GO:0019438	aromatic compound biosynthetic process	2585	49	33.49	0.00229	0.0107
GO-0003306 associative learning 45 4 4 0.58 0.00267 0.0107	GO:0021955	central nervous system neuron axonogenes	21	3	0.27	0.00239	0.0107
GO-0004656 nucleobase-containing compound biosynthes. 2537 48 32.87 0.00274 0.0107	GO:0018130	heterocycle biosynthetic process	2594	49	33.61	0.00247	0.0107
GO-0008251 regulation of glial cell proliferation 22 3 0.29 0.00274 0.0107	GO:0008306	associative learning	45	4	0.58	0.00267	0.0107
CO-0004576 single-organism behavior 227 9 2.94 0.00279 0.0107	GO:0034654	nucleobase-containing compound biosynthe	2537	48	32.87	0.00270	0.0107
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GO.0051865 positive regulation of synapse assembly 25 3 0.32 0.00388 0.0114 (SO.1901362 organic cyclic compound biosynthetic pro 2660 49 34.46 0.00420 0.0114 (GO.00323210 leptin-mediated signaling pathway 8 2 0.10 0.00444 0.0114 (GO.0033210 leptin-mediated signaling pathway 8 2 0.10 0.00444 0.0114 (GO.0033210 leptin-mediated signaling pathway 8 2 0.10 0.00444 0.0114 (GO.0007611 learning or memory 120 6 1.55 0.00467 0.0117 (GO.0007611 learning or memory 120 6 1.55 0.00467 0.0117 (GO.00007611 learning or memory 120 6 1.55 0.00467 0.0118 (GO.00007611 regulation of multicellular org 27 3 0.35 0.00497 0.0118 (GO.00007615 regulation of transcription, DNA-templat 2215 42 28.70 0.00526 0.0118 (GO.00003326 regulation of transcription, DNA-templat 2215 42 28.70 0.00566 0.0118 (GO.0003296 negative regulation of response to food 9 2 0.12 0.00566 0.0118 (GO.0003296 negative regulation of JUN kinase activi 9 2 0.12 0.00566 0.0118 (GO.0003367 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 (GO.0004320 cellular response to leptin stimulus 9 2 0.12 0.00566 0.0118 (GO.0006285 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO.0006285 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO.0006285 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO.0006285 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO.0006285 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO.0006285 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO.0003791) post-embryonic development 89 5 1.15 0.00593 0.0119 (GO.00021552 regulation of macromolecule biosynthetic 2565 47 33.23 0.00578 0.0121 (GO.0002154 subpallium development 10 2 0.13 0.00701 0.0121 (GO.0002154 subpallium development 10 2 0.13 0.00701 0.0121 (GO.0003104009 gilal cell proliteration 10 2 0.13 0.00701 0.0121 (GO.0003104009 gilal cell proliteration 10 0.000404 0.000304 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000310 0.000	GO:0001754	eye photoreceptor cell differentiation	24	3	0.31	0.00354	0.0107
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GO:0032099 negative regulation of appetite 8 2 0.10 0.00444 0.0114 (GO:0033210 laptin-mediated signaling pathway 8 2 0.10 0.00444 0.0114 (GO:0034350 regulation of gilal cell apontotic proce 8 2 0.10 0.00444 0.0114 (GO:0034350 regulation of gilal cell apontotic proce 8 2 0.10 0.00444 0.0114 (GO:003651 learning or memory 120 6 1.55 0.00467 0.0117 (GO:0040018 positive regulation of multicellular org 27 3 0.35 0.00497 0.0118 (GO:0006555 regulation of transcription, DNA-templat 2215 42 28.70 0.00566 0.0118 (GO:0001539 cilium of flagellum-dependent cell motil 9 2 0.12 0.00566 0.0118 (GO:0032096 negative regulation of response to food 9 2 0.12 0.00566 0.0118 (GO:0034508 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 (GO:0043508 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 (GO:0043507 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 (GO:0043507 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 (GO:004043507 regulation of macromolecule biosynthetic 2555 47 33.23 0.00578 0.0118 (GO:006565 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO:0006565 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO:0006565 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO:0006565 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO:0006565 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO:0006565 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 (GO:0006565 cilium-dependent 189 5 1.15 0.00593 0.0118 (GO:0006565 cilium-dependent 189 5 1.15 0.00593 0.0118 (GO:0006565 cilium-dependent 189 5 1.15 0.00593 0.0118 (GO:0006565 cilium-dependent 170 7 2.20 0.00663 0.0121 (GO:0006565 cilium-dependent 170 7 2.20 0.00666 0.0121 (GO:0006565 cilium-dependent 170 7 2.20 0.00666 0.0121 (GO:0006565 cilium-dependent 170 7 2.20 0.00666 0.0121 (GO:0006565 cilium-dependent 170 0.0121 (GO:0006565 cilium-dependent 170 0.0121 (GO:0006565 cilium-dependent 170 0.0121 (GO:0006565 cilium-dependent 170 0.0121 (GO:0006565 cilium	GO:0051965	positive regulation of synapse assembly	25	3	0.32	0.00398	0.0114
GO:0033210 leptin-mediated signaling pathway 8 2 0.10 0.00444 0.0114 GO:0034350 regulation of glial cell apoptotic proce 8 2 0.10 0.00444 0.0114 GO:00034350 regulation of glial cell projection proces 8 2 0.10 0.00444 0.0114 GO:0007611 learning or memory 120 6 1.55 0.00467 0.0117 GO:0040018 positive regulation of multicellular org 27 3 0.35 0.00467 0.0118 GO:000555 regulation of transcription, DNA-templat 2215 42 28.70 0.00526 0.0118 GO:0005153 cilium or flagellum-dependent cell motil 9 2 0.12 0.00566 0.0118 GO:0005153 regulation of response to food 9 2 0.12 0.00566 0.0118 GO:0032098 regulation of JUN kinasa activit 9 2 0.12 0.00566 0.0118 GO:0043508 negative regulation of JUN kinasa activit 9 2 0.12 0.00566 0.0118 GO:0043507 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 GO:0043507 regulation of insulin-like growth factor 9 2 0.12 0.00566 0.0118 GO:0043508 cilium-dependent cell motility 9 2 0.12 0.00566 0.0118 GO:0006356 dilum-dependent cell motility 9 2 0.12 0.00566 0.0118 GO:00069791 post-embryonic development 89 5 1.15 0.00566 0.0118 GO:0009791 post-embryonic development 89 5 1.15 0.00566 0.0118 GO:00072559 protein localization to plasma membrane 170 7 2.20 0.00663 0.0121 GO:0072559 protein localization to plasma membrane 170 7 2.20 0.00663 0.0121 GO:00014009 glial cell proliferation 30 3 0.39 0.00670 0.0121 GO:0001756 striatum development 10 2 0.13 0.00701 0.0121 GO:0001756 regulation of response to extra 32 3 0.41 0.00804 0.0121 GO:0001756 regulation of response to extra 32 3 0.41 0.00804 0.0121 GO:0001756 regulation of response to extra 32 3 0.41 0.00804 0.0121 GO:0000175 regulation of transcription fro 11 2 0.14 0.00850 0.0121 GO:0000181 regulation of transcription fro 11 2 0.14 0.00850 0.0121 GO:0000181 regulation of transcription fro 11 2 0.14 0.00850 0.0121 GO:0000181 reg	GO:1901362		2660	49	34.46		0.0114
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GO:0050890 cognition 138 6 1.79 0.00909 0.0121	GO:0032990	cell part morphogenesis	530	14	6.87	0.00891	0.0121
	GO:0009416	response to light stimulus	180	7	2.33	0.00896	0.0121
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	GO:0035264	multicellular organism growth	140	6	1.81	0.00972	0.0128