

source	native	name	p_value	significant	description	term_size	query_size	intersection	effective_c	precision	recall
GO:BP	GO:00995:	trans-syna	1.49E-70	TRUE	"Cell-cell s	772	1172	192	20972	0.163822526	0.248704663
GO:BP	GO:00995:	synaptic si	4E-69	TRUE	"Cell-cell s	803	1172	194	20972	0.16552901	0.241594022
GO:BP	GO:00989:	anterograc	3.12E-68	TRUE	"Cell-cell s	765	1172	188	20972	0.160409556	0.245751634
GO:BP	GO:00072:	chemical s	3.12E-68	TRUE	"The vesic	765	1172	188	20972	0.160409556	0.245751634
GO:BP	GO:00220:	neurogene	4.96E-62	TRUE	"Generatic	1769	1172	285	20972	0.243174061	0.161107971
GO:BP	GO:00508:	synapse or	1.04E-59	TRUE	"A process	561	1172	151	20972	0.12883959	0.26916221
GO:BP	GO:00991:	regulation	7.5E-58	TRUE	"Any proce	504	1172	141	20972	0.120307167	0.279761905
GO:BP	GO:00486:	generator	1.84E-57	TRUE	"The proce	1523	1172	254	20972	0.216723549	0.1667761
GO:BP	GO:00508:	modulatio	3.61E-57	TRUE	"Any proce	503	1172	140	20972	0.119453925	0.27833002
GO:BP	GO:00301:	neuron dif	1.2E-53	TRUE	"The proce	1444	1172	240	20972	0.204778157	0.166204986
GO:BP	GO:00072:	cell-cell sig	2.38E-53	TRUE	"Any proce	1316	1172	227	20972	0.193686007	0.172492401
GO:BP	GO:00343:	cell junctio	2.07E-49	TRUE	"A process	829	1172	170	20972	0.145051195	0.205066345
GO:BP	GO:00486:	neuron de	8.82E-48	TRUE	"The proce	1167	1172	203	20972	0.173208191	0.1739503
GO:BP	GO:00074:	central ner	1.41E-45	TRUE	"The proce	1035	1172	186	20972	0.158703072	0.179710145
GO:BP	GO:00311:	neuron prc	2.43E-45	TRUE	"The proce	1010	1172	183	20972	0.156143345	0.181188119
GO:BP	GO:00511:	regulation	2.75E-45	TRUE	"Any proce	2427	1172	310	20972	0.264505119	0.127729707
GO:BP	GO:00009:	cell morph	8.11E-42	TRUE	"The devel	998	1172	176	20972	0.150170648	0.176352705
GO:BP	GO:00076:	behavior	1.13E-41	TRUE	"The interi	665	1172	140	20972	0.119453925	0.210526316
GO:BP	GO:01200:	plasma me	9.98E-40	TRUE	"The proce	664	1172	137	20972	0.116894198	0.206325301
GO:BP	GO:00488:	cell projec	2.3E-39	TRUE	"The proce	669	1172	137	20972	0.116894198	0.204783259
GO:BP	GO:00507:	regulation	3.03E-39	TRUE	"Any proce	2431	1172	297	20972	0.253412969	0.122171946
GO:BP	GO:00488:	neuron prc	5.86E-39	TRUE	"The proce	648	1172	134	20972	0.114334471	0.206790123
GO:BP	GO:00423:	regulation	1.11E-38	TRUE	"Any proce	443	1172	109	20972	0.093003413	0.246049661
GO:BP	GO:00995:	trans-syna	6.26E-37	TRUE	"Cell-cell s	772	682	110	20972	0.161290323	0.142487047
GO:BP	GO:00995:	synaptic si	3.32E-36	TRUE	"Cell-cell s	803	682	111	20972	0.162756598	0.138231631
GO:BP	GO:00063:	chromatin	4.08E-36	TRUE	"The asser	1216	1172	187	20972	0.159556314	0.153782895
GO:BP	GO:00603:	head deve	1.18E-35	TRUE	"The biolo	787	1172	144	20972	0.122866894	0.182973316
GO:BP	GO:00063:	chromatin	1.55E-35	TRUE	"The asser	1216	294	88	20972	0.299319728	0.072368421
GO:BP	GO:00072:	chemical s	8.83E-35	TRUE	"The vesic	765	682	106	20972	0.15542522	0.138562092
GO:BP	GO:00989:	anterograc	8.83E-35	TRUE	"Cell-cell s	765	682	106	20972	0.15542522	0.138562092
GO:BP	GO:00300:	cell projec	1E-34	TRUE	"A process	1628	1172	221	20972	0.188566553	0.135749386
GO:BP	GO:00074:	brain deve	1.15E-34	TRUE	"The proce	736	1172	137	20972	0.116894198	0.186141304

GO:BP	GO:01200: plasma me	3.07E-34	TRUE	"A process	1582	1172	216	20972	0.184300341	0.13653603
GO:BP	GO:00074: synapse as	6.51E-34	TRUE	"The aggrega	266	1172	79	20972	0.067406143	0.296992481
GO:BP	GO:00063: chromatin	2.07E-33	TRUE	"A dynam	1057	294	80	20972	0.272108844	0.075685904
GO:BP	GO:00486: cell morph	4.05E-33	TRUE	"The process	584	1172	118	20972	0.100682594	0.202054795
GO:BP	GO:00063: chromatin	6.49E-33	TRUE	"A dynam	1057	1172	166	20972	0.141638225	0.15704825
GO:BP	GO:00508: regulation	5.39E-31	TRUE	"Any process	305	1172	81	20972	0.069112628	0.26557377
GO:BP	GO:00343: cell junctio	8.28E-31	TRUE	"A cellular	509	1172	106	20972	0.090443686	0.208251473
GO:BP	GO:00508: cognition	9.95E-31	TRUE	"The operati	315	1172	82	20972	0.06996587	0.26031746
GO:BP	GO:00512: positive re	1.2E-30	TRUE	"Any process	1804	294	98	20972	0.333333333	0.054323725
GO:BP	GO:00512: positive re	2.61E-30	TRUE	"Any process	1804	1172	226	20972	0.192832765	0.125277162
GO:BP	GO:00508: regulation	3.55E-30	TRUE	"Any process	298	1172	79	20972	0.067406143	0.265100671
GO:BP	GO:19026: positive re	8.79E-30	TRUE	"Any process	1674	294	93	20972	0.316326531	0.055555556
GO:BP	GO:00458: positive re	8.79E-30	TRUE	"Any process	1672	294	93	20972	0.316326531	0.05562201
GO:BP	GO:19026: positive re	9.12E-30	TRUE	"Any process	1674	1172	214	20972	0.182593857	0.127837515
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GO:BP	GO:00076: learning or	1.78E-29	TRUE	"The acquir	268	1172	74	20972	0.063139932	0.276119403
GO:BP	GO:00458: positive re	2.06E-29	TRUE	"Any process	1672	1172	213	20972	0.181740614	0.127392344
GO:BP	GO:00072: cell-cell sig	2.59E-29	TRUE	"Any process	1316	682	132	20972	0.193548387	0.100303951
GO:BP	GO:00071: cell adhesi	1.07E-28	TRUE	"The attac	1512	1172	198	20972	0.16894198	0.130952381
GO:BP	GO:00600: regulation	1.52E-27	TRUE	"Any process	145	1172	53	20972	0.045221843	0.365517241
GO:BP	GO:00508: synapse or	2.19E-27	TRUE	"A process	561	682	81	20972	0.118768328	0.144385027
GO:BP	GO:00030: system pro	3.58E-26	TRUE	"A multice	2290	1172	254	20972	0.216723549	0.110917031
GO:BP	GO:00991: regulation	2.35E-25	TRUE	"Any process	504	682	74	20972	0.108504399	0.146825397
GO:BP	GO:00352: synaptic tr	2.57E-25	TRUE	"The vesicle	107	1172	44	20972	0.037542662	0.411214953
GO:BP	GO:00481: regulation	4.46E-25	TRUE	"A process	221	1172	62	20972	0.052901024	0.280542986
GO:BP	GO:00508: modulatio	1.09E-24	TRUE	"Any process	503	682	73	20972	0.107038123	0.145129225
GO:BP	GO:00343: cell junctio	1.13E-24	TRUE	"A process	829	682	95	20972	0.139296188	0.114595899
GO:BP	GO:00074: axonogene	1.29E-24	TRUE	"De novo g	452	1172	90	20972	0.076791809	0.199115044
GO:BP	GO:00309: forebrain c	2.44E-24	TRUE	"The process	403	1172	84	20972	0.071672355	0.208436725
GO:BP	GO:00076: learning	4.41E-24	TRUE	"Any process	155	1172	51	20972	0.043515358	0.329032258
GO:BP	GO:00063: chromatin	9.4E-24	TRUE	"The asser	1216	227	63	20972	0.27753304	0.051809211
GO:BP	GO:00991: postsynap	9.84E-24	TRUE	"A process	233	1172	62	20972	0.052901024	0.266094421

GO:BP	GO:00512! positive re	1.94E-23	TRUE	"Any proce	1639	1172	196	20972	0.167235495	0.119585113
GO:BP	GO:00510! regulation	2.1E-23	TRUE	"Any proce	1566	1172	190	20972	0.162116041	0.121328225
GO:BP	GO:19016! response t	3.65E-23	TRUE	"Any proce	941	1172	136	20972	0.116040956	0.144527099
GO:BP	GO:00512! positive re	4.43E-23	TRUE	"Any proce	1804	227	75	20972	0.330396476	0.041574279
GO:BP	GO:00106! positive re	5.11E-23	TRUE	"Any proce	1778	1172	206	20972	0.175767918	0.115860517
GO:BP	GO:00220! neurogene	5.56E-23	TRUE	"Generatic	1769	682	144	20972	0.211143695	0.081401922
GO:BP	GO:00508! cognition	8.25E-23	TRUE	"The opera	315	227	35	20972	0.154185022	0.111111111
GO:BP	GO:00989! anterograc	8.25E-23	TRUE	"Cell-cell s	765	227	50	20972	0.220264317	0.065359477
GO:BP	GO:00991! regulation	8.25E-23	TRUE	"Any proce	504	227	42	20972	0.185022026	0.083333333
GO:BP	GO:00072! chemical s	8.25E-23	TRUE	"The vesic	765	227	50	20972	0.220264317	0.065359477
GO:BP	GO:00508! modulatio	8.25E-23	TRUE	"Any proce	503	227	42	20972	0.185022026	0.083499006
GO:BP	GO:00455! regulation	9.57E-23	TRUE	"Any proce	1549	1172	187	20972	0.159556314	0.120723047
GO:BP	GO:00995! trans-syna	1.18E-22	TRUE	"Cell-cell s	772	227	50	20972	0.220264317	0.064766839
GO:BP	GO:00986! cell-cell ad	1.32E-22	TRUE	"The attac	965	1172	137	20972	0.116894198	0.141968912
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GO:BP	GO:00071! cell adhesi	1.39E-22	TRUE	"The attac	1512	682	130	20972	0.190615836	0.085978836
GO:BP	GO:00615! axon devel	2.06E-22	TRUE	"The progr	513	1172	93	20972	0.079351536	0.18128655
GO:BP	GO:00510! positive re	2.17E-22	TRUE	"Any proce	1305	1172	166	20972	0.141638225	0.127203065
GO:BP	GO:00220! neurogene	2.41E-22	TRUE	"Generatic	1769	227	73	20972	0.321585903	0.041266252
GO:BP	GO:00459! positive re	2.8E-22	TRUE	"Any proce	1237	1172	160	20972	0.136518771	0.12934519
GO:BP	GO:00063! chromatin	2.91E-22	TRUE	"A dynami	1057	227	57	20972	0.251101322	0.053926206
GO:BP	GO:19016! cellular res	3.91E-22	TRUE	"Any proce	676	1172	109	20972	0.093003413	0.161242604
GO:BP	GO:00215! telenceph	3.91E-22	TRUE	"The proce	272	1172	65	20972	0.055460751	0.238970588
GO:BP	GO:00511! positive re	4.34E-22	TRUE	"Any proce	1091	1172	147	20972	0.125426621	0.134738772
GO:BP	GO:00508! synapse or	4.49E-22	TRUE	"A process	561	227	43	20972	0.189427313	0.076648841
GO:BP	GO:00351! social beh	4.54E-22	TRUE	"Behavior	58	1172	31	20972	0.026450512	0.534482759
GO:BP	GO:00995! synaptic si	5.31E-22	TRUE	"Cell-cell s	803	227	50	20972	0.220264317	0.062266501
GO:BP	GO:00076! learning or	6.54E-22	TRUE	"The acqui	268	227	32	20972	0.140969163	0.119402985
GO:BP	GO:00486! generati	7.88E-22	TRUE	"The proce	1523	227	67	20972	0.295154185	0.043992121
GO:BP	GO:00440! regulation	9.01E-22	TRUE	"Any proce	543	1172	95	20972	0.08105802	0.174953959
GO:BP	GO:00517! biological p	1.74E-21	TRUE	"Any proce	60	1172	31	20972	0.026450512	0.516666667
GO:BP	GO:00068! neurotrans	2.21E-21	TRUE	"The direc	211	1172	56	20972	0.04778157	0.265402844

GO:BP	GO:00486! generation	2.37E-21	TRUE	"The proce	1523	682	128	20972	0.187683284	0.084044649
GO:BP	GO:00459! positive re	2.46E-21	TRUE	"Any proce	1237	294	69	20972	0.234693878	0.055780113
GO:BP	GO:00458! positive re	4.46E-21	TRUE	"Any proce	1672	227	69	20972	0.303964758	0.041267943
GO:BP	GO:19026! positive re	4.63E-21	TRUE	"Any proce	1674	227	69	20972	0.303964758	0.041218638
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GO:BP	GO:00995! chemical s	1.11E-20	TRUE	"The part c	116	1172	41	20972	0.034982935	0.353448276
GO:BP	GO:00163! dendrite d	1.47E-20	TRUE	"The proce	234	1172	58	20972	0.049488055	0.247863248
GO:BP	GO:00507! regulation	1.73E-20	TRUE	"Any proce	2431	294	97	20972	0.329931973	0.039901275
GO:BP	GO:00708! cellular res	2.14E-20	TRUE	"Any proce	2199	1172	231	20972	0.197098976	0.105047749
GO:BP	GO:00301! neuron dif	2.91E-20	TRUE	"The proce	1444	227	63	20972	0.27753304	0.043628809
GO:BP	GO:00600! excitatory	3.2E-20	TRUE	"A process	107	1172	39	20972	0.033276451	0.364485981
GO:BP	GO:20000! regulation	7.48E-20	TRUE	"Any proce	1392	1172	167	20972	0.142491468	0.119971264
GO:BP	GO:00400! growth	7.91E-20	TRUE	"The incre	920	1172	127	20972	0.108361775	0.138043478
GO:BP	GO:19017! response t	8.53E-20	TRUE	"Any proce	1632	1172	186	20972	0.158703072	0.113970588
GO:BP	GO:00485! developme	9.37E-20	TRUE	"The incre	661	1172	103	20972	0.087883959	0.155824508
GO:BP	GO:00486! generatio	1.19E-19	TRUE	"The proce	1523	294	74	20972	0.25170068	0.048588313
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GO:BP	GO:00076! behavior	1.93E-19	TRUE	"The inter	665	227	43	20972	0.189427313	0.064661654
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GO:BP	GO:00313! regulation	2.68E-19	TRUE	"Any proce	639	1172	100	20972	0.085324232	0.156494523
GO:BP	GO:00508! nervous sy	4.02E-19	TRUE	"An organ	1503	1172	174	20972	0.148464164	0.115768463
GO:BP	GO:00072! glutamate	4.55E-19	TRUE	"The serie	60	1172	29	20972	0.024744027	0.483333333
GO:BP	GO:00440! regulation	1.02E-18	TRUE	"Any proce	1006	1172	132	20972	0.112627986	0.131212724
GO:BP	GO:00164! cell migrat	1.22E-18	TRUE	"The contr	1532	1172	175	20972	0.149317406	0.114229765
GO:BP	GO:00300! metal ion t	1.26E-18	TRUE	"The direc	860	1172	119	20972	0.101535836	0.138372093
GO:BP	GO:00519! regulation	1.46E-18	TRUE	"Any proce	72	1172	31	20972	0.026450512	0.430555556
GO:BP	GO:00109! regulation	1.92E-18	TRUE	"Any proce	426	1172	77	20972	0.065699659	0.180751174
GO:BP	GO:00301! neuron dif	1.99E-18	TRUE	"The proce	1444	294	70	20972	0.238095238	0.048476454
GO:BP	GO:00099! positive re	2.54E-18	TRUE	"Any proce	1569	1172	177	20972	0.151023891	0.112810707
GO:BP	GO:00508! positive re	2.71E-18	TRUE	"Any proce	174	1172	47	20972	0.040102389	0.270114943

GO:BP	GO:004861 neuron de	2.83E-18	TRUE	"The proce	1167	682	103	20972	0.151026393	0.088260497
GO:BP	GO:000681 monoatom	3.27E-18	TRUE	"The direc	1260	1172	152	20972	0.129692833	0.120634921
GO:BP	GO:004581 negative re	4E-18	TRUE	"Any proce	1275	1172	153	20972	0.130546075	0.12
GO:BP	GO:014111 intracellul	4.95E-18	TRUE	"An intrac	1866	1172	199	20972	0.169795222	0.10664523
GO:BP	GO:005191 positive re	5.07E-18	TRUE	"Any proce	271	1172	59	20972	0.050341297	0.217712177
GO:BP	GO:000721 cell-cell sig	5.21E-18	TRUE	"Any proce	1316	227	57	20972	0.251101322	0.04331307
GO:BP	GO:005191 regulation	6.32E-18	TRUE	"Any proce	163	1172	45	20972	0.038395904	0.27607362
GO:BP	GO:190261 negative re	7.57E-18	TRUE	"Any proce	1284	1172	153	20972	0.130546075	0.119158879
GO:BP	GO:000761 learning	7.8E-18	TRUE	"Any proce	155	227	23	20972	0.101321586	0.148387097
GO:BP	GO:009861 cell-cell ad	8.36E-18	TRUE	"The attac	965	682	91	20972	0.133431085	0.094300518
GO:BP	GO:005111 regulation	8.6E-18	TRUE	"Any proce	2427	294	92	20972	0.31292517	0.037906881
GO:BP	GO:009901 postsynap	9.55E-18	TRUE	"A process	56	1172	27	20972	0.023037543	0.482142857
GO:BP	GO:190181 regulation	1E-17	TRUE	"Any proce	258	1172	57	20972	0.048634812	0.220930233
GO:BP	GO:005121 negative re	1.11E-17	TRUE	"Any proce	1389	1172	161	20972	0.137372014	0.115910727
GO:BP	GO:005121 negative re	1.22E-17	TRUE	"Any proce	1389	227	58	20972	0.255506608	0.041756659
GO:BP	GO:006031 head deve	1.37E-17	TRUE	"The biolo	787	227	44	20972	0.193832599	0.055908513
GO:BP	GO:012001 regulation	1.4E-17	TRUE	"Any proce	623	1172	95	20972	0.08105802	0.152487961
GO:BP	GO:003111 neuron pr	1.5E-17	TRUE	"The proce	1010	682	93	20972	0.136363636	0.092079208
GO:BP	GO:003431 cell junctio	1.53E-17	TRUE	"A process	829	227	45	20972	0.198237885	0.054282268
GO:BP	GO:000741 central ner	1.53E-17	TRUE	"The proce	1035	227	50	20972	0.220264317	0.048309179
GO:BP	GO:002261 regulation	1.81E-17	TRUE	"Any proce	822	1172	113	20972	0.096416382	0.137469586
GO:BP	GO:004861 neuron de	2.37E-17	TRUE	"The proce	1167	294	61	20972	0.207482993	0.05227078
GO:BP	GO:000811 intracellul	2.88E-17	TRUE	"Any proce	2401	1172	236	20972	0.201365188	0.098292378
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GO:BP	GO:005111 regulation	3.02E-17	TRUE	"Any proce	2427	227	77	20972	0.339207048	0.031726411
GO:BP	GO:003111 neuron pr	3.02E-17	TRUE	"The proce	1010	227	49	20972	0.215859031	0.048514851
GO:BP	GO:003421 monoatom	3.17E-17	TRUE	"A process	1025	1172	130	20972	0.110921502	0.126829268
GO:BP	GO:000171 neuron mi	3.26E-17	TRUE	"The chara	192	1172	48	20972	0.040955631	0.25
GO:BP	GO:000151 action pot	3.62E-17	TRUE	"A process	170	1172	45	20972	0.038395904	0.264705882
GO:BP	GO:190261 negative re	3.76E-17	TRUE	"Any proce	1284	227	55	20972	0.242290749	0.042834891
GO:BP	GO:005111 regulation	4.98E-17	TRUE	"Any proce	2427	682	161	20972	0.236070381	0.066337042
GO:BP	GO:000761 behavior	4.98E-17	TRUE	"The interi	665	682	72	20972	0.105571848	0.108270677

GO:BP	GO:005501:transmembrane transport	6.18E-17	TRUE	"The process of moving substances across the cell membrane"	1555	1172	172	20972	0.146757679	0.110610932
GO:BP	GO:000091:cell morphogenesis	6.81E-17	TRUE	"The development of the shape and structure of the cell"	998	682	91	20972	0.133431085	0.091182365
GO:BP	GO:003511:social behavior	7.68E-17	TRUE	"Behavior of an organism in relation to other organisms of the same species"	58	227	16	20972	0.070484581	0.275862069
GO:BP	GO:009881:modulation of synaptic transmission	7.72E-17	TRUE	"Any process that modulates the transmission of signals across a synapse"	47	227	15	20972	0.066079295	0.319148936
GO:BP	GO:005121:negative regulation of cell growth	7.95E-17	TRUE	"Any process that inhibits the growth of a cell"	1389	294	66	20972	0.224489796	0.047516199
GO:BP	GO:004591:negative regulation of cell cycle	1.11E-16	TRUE	"Any process that inhibits the cell cycle"	1510	227	59	20972	0.259911894	0.039072848
GO:BP	GO:005171:biological process	1.32E-16	TRUE	"Any process that occurs in a living organism"	60	227	16	20972	0.070484581	0.266666667
GO:BP	GO:004881:neuron projection morphogenesis	1.36E-16	TRUE	"The process of the development of the shape and structure of a neuron projection"	648	227	39	20972	0.171806167	0.060185185
GO:BP	GO:009911:regulation of gene expression	1.63E-16	TRUE	"Any process that modulates the expression of a gene"	140	1172	40	20972	0.034129693	0.285714286
GO:BP	GO:009881:modulation of synaptic transmission	1.63E-16	TRUE	"Any process that modulates the transmission of signals across a synapse"	47	1172	24	20972	0.020477816	0.510638298
GO:BP	GO:009711:postsynaptic transmission	1.63E-16	TRUE	"A process that occurs at the synapse after the release of neurotransmitters"	47	1172	24	20972	0.020477816	0.510638298
GO:BP	GO:004591:positive regulation of cell cycle	1.69E-16	TRUE	"Any process that promotes the cell cycle"	1237	227	53	20972	0.233480176	0.042845594
GO:BP	GO:000741:central nervous system development	1.74E-16	TRUE	"The process of the development of the central nervous system"	1035	294	56	20972	0.19047619	0.05410628
GO:BP	GO:004861:cell morphology	2.04E-16	TRUE	"The process of the development of the shape and structure of a cell"	584	227	37	20972	0.162995595	0.063356164
GO:BP	GO:000301:circulatory system	2.06E-16	TRUE	"An organ system that transports substances throughout the body"	595	1172	90	20972	0.076791809	0.151260504
GO:BP	GO:190251:regulation of gene expression	2.07E-16	TRUE	"Any process that modulates the expression of a gene"	1945	1172	200	20972	0.170648464	0.102827763
GO:BP	GO:000741:central nervous system development	2.08E-16	TRUE	"The process of the development of the central nervous system"	1035	682	92	20972	0.134897361	0.088888889
GO:BP	GO:004591:negative regulation of cell cycle	2.1E-16	TRUE	"Any process that inhibits the cell cycle"	1510	1172	167	20972	0.142491468	0.110596026
GO:BP	GO:006021:regulation of gene expression	2.24E-16	TRUE	"Any process that modulates the expression of a gene"	816	1172	110	20972	0.093856655	0.134803922
GO:BP	GO:000741:brain development	2.46E-16	TRUE	"The process of the development of the brain"	736	227	41	20972	0.18061674	0.055706522
GO:BP	GO:003111:neuron projection morphogenesis	2.72E-16	TRUE	"The process of the development of the shape and structure of a neuron projection"	1010	294	55	20972	0.18707483	0.054455446
GO:BP	GO:002191:central nervous system development	2.88E-16	TRUE	"The process of the development of the central nervous system"	218	1172	50	20972	0.042662116	0.229357798
GO:BP	GO:005161:establishment of gene expression	2.89E-16	TRUE	"Any process that modulates the expression of a gene"	1938	1172	199	20972	0.169795222	0.102683179
GO:BP	GO:012001:plasma membrane	2.92E-16	TRUE	"The process of the development of the plasma membrane"	664	227	39	20972	0.171806167	0.05873494
GO:BP	GO:004591:negative regulation of cell cycle	3.22E-16	TRUE	"Any process that inhibits the cell cycle"	1510	294	68	20972	0.231292517	0.045033113
GO:BP	GO:004881:cell projection morphogenesis	3.73E-16	TRUE	"The process of the development of the shape and structure of a cell projection"	669	227	39	20972	0.171806167	0.058295964
GO:BP	GO:199081:ligand-gated ion channel activity	3.91E-16	TRUE	"The series of events that lead to the opening of a ligand-gated ion channel"	44	1172	23	20972	0.019624573	0.522727273
GO:BP	GO:200041:positive regulation of gene expression	4.29E-16	TRUE	"Any process that promotes the expression of a gene"	32	1172	20	20972	0.017064846	0.625
GO:BP	GO:000961:response to stress	4.57E-16	TRUE	"Any process that occurs in response to a stressor"	1108	1172	134	20972	0.114334471	0.120938628
GO:BP	GO:000701:cell cycle	5.03E-16	TRUE	"The process of the development of the cell cycle"	1668	1172	178	20972	0.151877133	0.106714628
GO:BP	GO:000301:system process	5.24E-16	TRUE	"A multicomponent system that performs a specific function"	2290	682	152	20972	0.2228739	0.066375546
GO:BP	GO:006031:head development	5.62E-16	TRUE	"The biological process of the development of the head"	787	294	48	20972	0.163265306	0.060991105

GO:BP	GO:00714! cellular res	7.44E-16	TRUE	"Any proce	1238	1172	144	20972	0.122866894	0.11631664
GO:BP	GO:00097! response t	7.85E-16	TRUE	"Any proce	1467	1172	162	20972	0.138225256	0.110429448
GO:BP	GO:00343! cell junctio	8.33E-16	TRUE	"A cellular	509	682	60	20972	0.08797654	0.117878193
GO:BP	GO:00070! cytoskeleti	9.06E-16	TRUE	"A process	1534	1172	167	20972	0.142491468	0.108865711
GO:BP	GO:00328! regulation	1.09E-15	TRUE	"Any proce	1914	682	134	20972	0.196480938	0.070010449
GO:BP	GO:00328! cellular res	1.13E-15	TRUE	"Any proce	677	1172	96	20972	0.081911263	0.141802068
GO:BP	GO:00507! regulation	1.34E-15	TRUE	"Any proce	350	1172	64	20972	0.054607509	0.182857143
GO:BP	GO:00458! negative re	1.35E-15	TRUE	"Any proce	1275	294	61	20972	0.207482993	0.047843137
GO:BP	GO:00072! neurotran	1.63E-15	TRUE	"The regul	149	1172	40	20972	0.034129693	0.268456376
GO:BP	GO:00996! signal rele	1.63E-15	TRUE	"Any signa	149	1172	40	20972	0.034129693	0.268456376
GO:BP	GO:00230! signal rele	1.82E-15	TRUE	"The proce	490	1172	78	20972	0.066552901	0.159183673
GO:BP	GO:19026! negative re	1.86E-15	TRUE	"Any proce	1284	294	61	20972	0.207482993	0.047507788
GO:BP	GO:00080! blood circ	1.9E-15	TRUE	"The flow	511	1172	80	20972	0.068259386	0.156555773
GO:BP	GO:00423! regulation	1.99E-15	TRUE	"Any proce	443	682	55	20972	0.080645161	0.124153499
GO:BP	GO:00507! regulation	3.28E-15	TRUE	"Any proce	2431	682	156	20972	0.228739003	0.064171123
GO:BP	GO:00485! positive re	3.98E-15	TRUE	"Any proce	2320	1172	223	20972	0.190273038	0.09612069
GO:BP	GO:00455! regulation	4.14E-15	TRUE	"Any proce	1549	294	67	20972	0.227891156	0.043253712
GO:BP	GO:00001! negative re	4.14E-15	TRUE	"Any proce	933	294	51	20972	0.173469388	0.054662379
GO:BP	GO:00300! actin filam	4.28E-15	TRUE	"Any cellul	816	1172	107	20972	0.091296928	0.131127451
GO:BP	GO:00508! negative re	4.89E-15	TRUE	"Any proce	53	1172	24	20972	0.020477816	0.452830189
GO:BP	GO:00330! regulation	5.2E-15	TRUE	"Any proce	1167	1172	136	20972	0.116040956	0.116538132
GO:BP	GO:00030! circulatory	5.28E-15	TRUE	"An organ	595	682	64	20972	0.093841642	0.107563025
GO:BP	GO:00600! heart cont	5.29E-15	TRUE	"The multi	242	1172	51	20972	0.043515358	0.210743802
GO:BP	GO:00001! negative re	5.85E-15	TRUE	"Any proce	933	227	44	20972	0.193832599	0.0471597
GO:BP	GO:00107! positive re	6.61E-15	TRUE	"Any proce	430	1172	71	20972	0.060580205	0.165116279
GO:BP	GO:00488! cell motilit	6.67E-15	TRUE	"Any proce	1794	1172	184	20972	0.156996587	0.102564103
GO:BP	GO:00423! regulation	7.96E-15	TRUE	"Any proce	443	227	31	20972	0.136563877	0.069977427
GO:BP	GO:00995! synaptic ve	8.49E-15	TRUE	"A biologic	211	1172	47	20972	0.040102389	0.222748815
GO:BP	GO:00068! monoaton	9.19E-15	TRUE	"The direc	1052	1172	126	20972	0.107508532	0.119771863
GO:BP	GO:00001! negative re	1.05E-14	TRUE	"Any proce	933	1172	116	20972	0.098976109	0.124330118
GO:BP	GO:00215! pallium de	1.09E-14	TRUE	"The proce	188	1172	44	20972	0.037542662	0.234042553
GO:BP	GO:00986! import int	1.13E-14	TRUE	"The direc	958	1172	118	20972	0.100682594	0.123173278

GO:BP	GO:004551: positive re	1.19E-14	TRUE	"Any proce	852	1172	109	20972	0.093003413	0.127934272
GO:BP	GO:000091: cell morph	1.26E-14	TRUE	"The devel	998	227	45	20972	0.198237885	0.04509018
GO:BP	GO:000741: synapse as	1.43E-14	TRUE	"The aggre	266	682	41	20972	0.060117302	0.154135338
GO:BP	GO:014031: export froi	1.51E-14	TRUE	"The direc	914	1172	114	20972	0.097269625	0.124726477
GO:BP	GO:004811: regulation	1.59E-14	TRUE	"A process	221	227	23	20972	0.101321586	0.104072398
GO:BP	GO:003281: regulation	2.11E-14	TRUE	"Any proce	859	1172	109	20972	0.093003413	0.126891735
GO:BP	GO:012001: plasma me	2.32E-14	TRUE	"The proce	664	682	67	20972	0.098240469	0.100903614
GO:BP	GO:000301: heart proc	2.46E-14	TRUE	"A circulat	251	1172	51	20972	0.043515358	0.203187251
GO:BP	GO:005101: regulation	2.66E-14	TRUE	"Any proce	1566	682	114	20972	0.167155425	0.072796935
GO:BP	GO:001061: negative re	2.88E-14	TRUE	"Any proce	1449	1172	156	20972	0.133105802	0.107660455
GO:BP	GO:002301: negative re	2.88E-14	TRUE	"Any proce	1449	1172	156	20972	0.133105802	0.107660455
GO:BP	GO:000741: brain deve	2.89E-14	TRUE	"The proce	736	294	44	20972	0.149659864	0.059782609
GO:BP	GO:004881: cell projec	3.19E-14	TRUE	"The proce	669	682	67	20972	0.098240469	0.100149477
GO:BP	GO:000971: response t	4.41E-14	TRUE	"Any proce	904	1172	112	20972	0.09556314	0.123893805
GO:BP	GO:003521: ionotropic	4.54E-14	TRUE	"The serie	30	1172	18	20972	0.015358362	0.6
GO:BP	GO:009901: vesicle-me	4.58E-14	TRUE	"Any vesic	246	1172	50	20972	0.042662116	0.203252033
GO:BP	GO:000801: blood circ	5.32E-14	TRUE	"The flow	511	682	57	20972	0.083577713	0.111545988
GO:BP	GO:000831: associative	5.85E-14	TRUE	"Learning l	87	1172	29	20972	0.024744027	0.333333333
GO:BP	GO:002191: cerebral cc	7.21E-14	TRUE	"The progr	128	1172	35	20972	0.029863481	0.2734375
GO:BP	GO:000711: homophilic	7.41E-14	TRUE	"The attac	169	682	32	20972	0.046920821	0.189349112
GO:BP	GO:006001: regulation	8.83E-14	TRUE	"Any proce	145	227	19	20972	0.083700441	0.131034483
GO:BP	GO:000741: axonogene	9.51E-14	TRUE	"De novo g	452	227	30	20972	0.13215859	0.066371681
GO:BP	GO:006001: excitatory	1.01E-13	TRUE	"A process	107	227	17	20972	0.074889868	0.158878505
GO:BP	GO:000971: embryo de	1.18E-13	TRUE	"The proce	1126	1172	129	20972	0.110068259	0.114564831
GO:BP	GO:009901: postsynap	1.25E-13	TRUE	"The aggre	83	1172	28	20972	0.023890785	0.337349398
GO:BP	GO:012001: plasma me	1.25E-13	TRUE	"A process	1582	682	113	20972	0.16568915	0.071428571
GO:BP	GO:009861: monoatom	1.35E-13	TRUE	"The proce	859	1172	107	20972	0.091296928	0.124563446
GO:BP	GO:003001: cell projec	1.45E-13	TRUE	"A process	1628	682	115	20972	0.168621701	0.070638821
GO:BP	GO:000301: muscle sys	1.91E-13	TRUE	"An organ	428	1172	68	20972	0.058020478	0.158878505
GO:BP	GO:003001: cell projec	1.95E-13	TRUE	"A process	1628	227	56	20972	0.246696035	0.034398034
GO:BP	GO:005071: regulation	2.37E-13	TRUE	"Any proce	2431	227	70	20972	0.308370044	0.028794735
GO:BP	GO:005071: positive re	2.43E-13	TRUE	"Any proce	221	1172	46	20972	0.039249147	0.208144796

GO:BP	GO:00488: neuron pr	2.44E-13	TRUE	"The proce	648	294	40	20972	0.136054422	0.061728395
GO:BP	GO:00009: cell morph	2.56E-13	TRUE	"The devel	998	294	50	20972	0.170068027	0.0501002
GO:BP	GO:00488: neuron pr	2.58E-13	TRUE	"The proce	648	682	64	20972	0.093841642	0.098765432
GO:BP	GO:00313: positive re	3.07E-13	TRUE	"Any proce	342	1172	59	20972	0.050341297	0.17251462
GO:BP	GO:19016: response t	3.35E-13	TRUE	"Any proce	941	682	80	20972	0.117302053	0.08501594
GO:BP	GO:00329: secretion l	3.35E-13	TRUE	"The contr	835	1172	104	20972	0.088737201	0.124550898
GO:BP	GO:00995: chemical s	3.81E-13	TRUE	"The part c	116	227	17	20972	0.074889868	0.146551724
GO:BP	GO:00509: neuromusi	4.31E-13	TRUE	"Any proce	174	1172	40	20972	0.034129693	0.229885057
GO:BP	GO:00510: regulation	4.61E-13	TRUE	"Any proce	608	1172	84	20972	0.071672355	0.138157895
GO:BP	GO:01200: plasma me	5.24E-13	TRUE	"A process	1582	294	64	20972	0.217687075	0.04045512
GO:BP	GO:01200: plasma me	5.24E-13	TRUE	"The proce	664	294	40	20972	0.136054422	0.060240964
GO:BP	GO:00164: cell migrat	5.33E-13	TRUE	"The contr	1532	682	109	20972	0.159824047	0.071148825
GO:BP	GO:00224: cell cycle p	5.34E-13	TRUE	"The cellul	1290	1172	140	20972	0.119453925	0.108527132
GO:BP	GO:00082: cell death	5.34E-13	TRUE	"Any biolo	1962	1172	190	20972	0.162116041	0.096839959
GO:BP	GO:00300: cell projec	5.63E-13	TRUE	"A process	1628	294	65	20972	0.221088435	0.03992629
GO:BP	GO:00508: regulation	6.05E-13	TRUE	"Any proce	298	682	41	20972	0.060117302	0.137583893
GO:BP	GO:00481: regulation	6.22E-13	TRUE	"A process	58	1172	23	20972	0.019624573	0.396551724
GO:BP	GO:19035: regulation	6.23E-13	TRUE	"Any proce	556	1172	79	20972	0.067406143	0.142086331
GO:BP	GO:00300: actin filam	6.48E-13	TRUE	"Movemer	137	1172	35	20972	0.029863481	0.255474453
GO:BP	GO:00488: cell projec	6.49E-13	TRUE	"The proce	669	294	40	20972	0.136054422	0.059790732
GO:BP	GO:00069: apoptotic	6.81E-13	TRUE	"A prograr	1898	1172	185	20972	0.157849829	0.097471022
GO:BP	GO:00074: synapse as	7.72E-13	TRUE	"The aggre	266	227	23	20972	0.101321586	0.086466165
GO:BP	GO:00125: programm	8.5E-13	TRUE	"A process	1958	1172	189	20972	0.161262799	0.096527068
GO:BP	GO:00519: positive re	8.66E-13	TRUE	"Any proce	32	227	11	20972	0.04845815	0.34375
GO:BP	GO:20004: positive re	8.66E-13	TRUE	"Any proce	32	227	11	20972	0.04845815	0.34375
GO:BP	GO:01200: plasma me	8.66E-13	TRUE	"A process	1582	227	54	20972	0.237885463	0.034134008
GO:BP	GO:00019: startle res	9.8E-13	TRUE	"An action	30	1172	17	20972	0.014505119	0.566666667
GO:BP	GO:00469: secretion	9.8E-13	TRUE	"The contr	971	1172	114	20972	0.097269625	0.117404737
GO:BP	GO:00508: regulation	1.31E-12	TRUE	"Any proce	305	682	41	20972	0.060117302	0.13442623
GO:BP	GO:00510: positive re	1.39E-12	TRUE	"Any proce	818	1172	101	20972	0.086177474	0.123471883
GO:BP	GO:00070: cell cycle	1.71E-12	TRUE	"The progr	1668	294	65	20972	0.221088435	0.038968825
GO:BP	GO:00508: synapse or	1.85E-12	TRUE	"A process	561	294	36	20972	0.12244898	0.064171123

GO:BP	GO:00076: locomotor	1.98E-12	TRUE	"The speci	216	1172	44	20972	0.037542662	0.203703704
GO:BP	GO:00603: head deve	2.11E-12	TRUE	"The biolo	787	682	70	20972	0.102639296	0.088945362
GO:BP	GO:00335: cellular res	2.17E-12	TRUE	"Any proce	1856	294	69	20972	0.234693878	0.037176724
GO:BP	GO:00508: modulatio	2.22E-12	TRUE	"Any proce	503	294	34	20972	0.115646259	0.067594433
GO:BP	GO:00991: regulation	2.32E-12	TRUE	"Any proce	504	294	34	20972	0.115646259	0.067460317
GO:BP	GO:00615: axon devel	2.35E-12	TRUE	"The progr	513	227	30	20972	0.13215859	0.058479532
GO:BP	GO:00425: homeostat	2.5E-12	TRUE	"Any biolo	1677	1172	167	20972	0.142491468	0.099582588
GO:BP	GO:00601: prepulse ir	2.56E-12	TRUE	"The proce	14	1172	12	20972	0.010238908	0.857142857
GO:BP	GO:00074: brain deve	2.6E-12	TRUE	"The proce	736	682	67	20972	0.098240469	0.091032609
GO:BP	GO:00485: rhythmic p	2.68E-12	TRUE	"Any proce	291	1172	52	20972	0.044368601	0.178694158
GO:BP	GO:00069: muscle cor	2.87E-12	TRUE	"A process	340	1172	57	20972	0.048634812	0.167647059
GO:BP	GO:00076: memory	2.87E-12	TRUE	"The activi	121	1172	32	20972	0.027303754	0.26446281
GO:BP	GO:00076: circadian r	2.9E-12	TRUE	"Any biolo	201	1172	42	20972	0.035836177	0.208955224
GO:BP	GO:00161: vesicle-me	2.92E-12	TRUE	"A cellular	1572	1172	159	20972	0.135665529	0.101145038
GO:BP	GO:19030: mitotic cel	3.48E-12	TRUE	"A process	747	1172	94	20972	0.080204778	0.12583668
GO:BP	GO:00071: homophilic	3.83E-12	TRUE	"The attac	169	1172	38	20972	0.032423208	0.224852071
GO:BP	GO:00002: mitotic cel	3.94E-12	TRUE	"Progressi	893	1172	106	20972	0.090443686	0.118701008
GO:BP	GO:00352: tube devel	4.35E-12	TRUE	"The proce	1094	1172	122	20972	0.104095563	0.111517367
GO:BP	GO:00316: regulation	4.47E-12	TRUE	"Any proce	114	227	16	20972	0.070484581	0.140350877
GO:BP	GO:00991: postsynap	4.65E-12	TRUE	"A process	233	227	21	20972	0.092511013	0.090128755
GO:BP	GO:00069: DNA dama	4.97E-12	TRUE	"Any proce	929	294	46	20972	0.156462585	0.049515608
GO:BP	GO:00488: dendrite r	5.57E-12	TRUE	"The proce	139	1172	34	20972	0.029010239	0.244604317
GO:BP	GO:00486: cell morph	5.68E-12	TRUE	"The proce	584	294	36	20972	0.12244898	0.061643836
GO:BP	GO:00508: nervous sy	6.64E-12	TRUE	"An organ	1503	227	51	20972	0.224669604	0.033932136
GO:BP	GO:00515: regulation	6.98E-12	TRUE	"Any proce	103	1172	29	20972	0.024744027	0.281553398
GO:BP	GO:00860: cardiac mu	7.65E-12	TRUE	"An action	83	1172	26	20972	0.0221843	0.313253012
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GO:BP	GO:00160! synaptic ve	1.6E-10	TRUE	"Fusion of	101	1172	27	20972	0.023037543	0.267326733
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GO:BP	GO:003091 forebrain c	3.2E-09	TRUE	"The proce	403	227	23	20972	0.101321586	0.05707196
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GO:BP	GO:005161 vesicle loc	3.31E-09	TRUE	"Any proce	220	1172	39	20972	0.033276451	0.177272727
GO:BP	GO:006101 muscle str	3.34E-09	TRUE	"The progr	673	1172	80	20972	0.068259386	0.118870728
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GO:BP	GO:009711 postsynap	4.27E-09	TRUE	"The aggre	30	1172	14	20972	0.011945392	0.466666667
GO:BP	GO:190301 mitotic cel	4.7E-09	TRUE	"A process	747	294	36	20972	0.12244898	0.048192771
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GO:BP	GO:190481 excitatory	5.65E-09	TRUE	"The aggre	41	1172	16	20972	0.013651877	0.390243902
GO:BP	GO:008601 cardiac mu	5.67E-09	TRUE	"An action	59	1172	19	20972	0.016211604	0.322033898
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GO:BP	GO:190171 response t	6.08E-09	TRUE	"Any proce	1632	682	102	20972	0.149560117	0.0625
GO:BP	GO:003541 protein loc	6.41E-09	TRUE	"Any proce	87	1172	23	20972	0.019624573	0.264367816
GO:BP	GO:005191 regulation	6.7E-09	TRUE	"Any proce	163	682	25	20972	0.036656891	0.153374233
GO:BP	GO:002241 cell cycle p	6.75E-09	TRUE	"The cellul	1290	294	49	20972	0.166666667	0.037984496
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GO:BP	GO:009701 dendritic s	6.81E-09	TRUE	"A process	80	1172	22	20972	0.018771331	0.275
GO:BP	GO:014111 intracellul	7.16E-09	TRUE	"An intrac	1866	682	112	20972	0.164222874	0.060021436
GO:BP	GO:009951 maintenanc	7.23E-09	TRUE	"A process	31	1172	14	20972	0.011945392	0.451612903

GO:BP	GO:00335! multicellul	7.25E-09	TRUE	"Any proce	95	1172	24	20972	0.020477816	0.252631579
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GO:BP	GO:00329! secretion k	1.04E-08	TRUE	"The contr	835	682	64	20972	0.093841642	0.076646707
GO:BP	GO:00192! neuronal a	1.06E-08	TRUE	"An action	37	1172	15	20972	0.012798635	0.405405405
GO:BP	GO:00303! regulation	1.07E-08	TRUE	"Any proce	944	1172	100	20972	0.085324232	0.105932203
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GO:BP	GO:00708! cellular res	1.46E-08	TRUE	"Any proce	2199	172	49	20972	0.284883721	0.022282856
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GO:BP	GO:00301! neuron dif	2.77E-08	TRUE	"The proce	1444	172	38	20972	0.220930233	0.026315789

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GO:BP	GO:00098: tissue deve	3.44E-08	TRUE	"The proce	2014	1172	175	20972	0.149317406	0.086891758
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GO:BP	GO:00615: axon devel	3.77E-08	TRUE	"The progr	513	294	28	20972	0.095238095	0.054580897
GO:BP	GO:00512: chromosom	3.9E-08	TRUE	"A process	575	1172	69	20972	0.05887372	0.12
GO:BP	GO:00071: neuron cel	3.9E-08	TRUE	"The attac	16	1172	10	20972	0.008532423	0.625
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GO:BP	GO:00074: sensory or	4.27E-08	TRUE	"The proce	625	1172	73	20972	0.062286689	0.1168
GO:BP	GO:00434: regulation	4.48E-08	TRUE	"Any proce	638	1172	74	20972	0.063139932	0.115987461
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GO:BP	GO:00303: positive re	4.62E-08	TRUE	"Any proce	155	1172	30	20972	0.02559727	0.193548387
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GO:BP	GO:00435' negative re	6.1E-08	TRUE	"Any proce	166	1172	31	20972	0.026450512	0.186746988
GO:BP	GO:00109' positive re	6.24E-08	TRUE	"Any proce	148	1172	29	20972	0.024744027	0.195945946
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GO:BP	GO:00352' tube morp	6.9E-08	TRUE	"The proce	877	682	64	20972	0.093841642	0.072976055
GO:BP	GO:19017' cellular res	6.97E-08	TRUE	"Any proce	1194	682	79	20972	0.115835777	0.066164154
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GO:BP	GO:00510' positive re	7.27E-08	TRUE	"Any proce	1305	682	84	20972	0.123167155	0.064367816
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GO:BP	GO:00971' postsynapti	8.28E-08	TRUE	"The cluste	10	1172	8	20972	0.006825939	0.8
GO:BP	GO:19018' regulation	8.3E-08	TRUE	"Any proce	258	682	30	20972	0.04398827	0.11627907
GO:BP	GO:00508' regulation	8.42E-08	TRUE	"Any proce	305	294	21	20972	0.071428571	0.068852459
GO:BP	GO:00970' synaptic ve	8.47E-08	TRUE	"The proce	17	1172	10	20972	0.008532423	0.588235294
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GO:BP	GO:000721	small GTPa	1.02E-07	TRUE	"An intrac	493	1172	61	20972	0.052047782	0.123732252
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GO:BP	GO:00217 limbic syst	1.61E-07	TRUE	"The progr	119	1172	25	20972	0.021331058	0.210084034
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GO:BP	GO:00507 regulation	1.81E-07	TRUE	"Any proce	350	294	22	20972	0.074829932	0.062857143
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GO:BP	GO:00511 regulation	2.13E-07	TRUE	"Any proce	2427	172	49	20972	0.284883721	0.020189534
GO:BP	GO:00082 positive re	2.23E-07	TRUE	"Any proce	923	1172	94	20972	0.080204778	0.10184182
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GO:BP	GO:00400 locomotio	2.29E-07	TRUE	"Self-prop	1236	1172	117	20972	0.099829352	0.094660194
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GO:BP	GO:00995 synaptic m	2.6E-07	TRUE	"The attac	34	682	11	20972	0.016129032	0.323529412
GO:BP	GO:00215 telencepha	2.63E-07	TRUE	"The proce	272	682	30	20972	0.04398827	0.110294118
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GO:BP	GO:00507 regulation	2.69E-07	TRUE	"Any proce	350	227	19	20972	0.083700441	0.054285714
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GO:BP	GO:00107 positive re	2.78E-07	TRUE	"Any proce	430	227	21	20972	0.092511013	0.048837209
GO:BP	GO:00448 cell cycle C	2.85E-07	TRUE	"The cell c	239	294	18	20972	0.06122449	0.075313808
GO:BP	GO:00450 regulated c	2.87E-07	TRUE	"A process	227	1172	36	20972	0.030716724	0.158590308
GO:BP	GO:00000 G1/S trans	2.9E-07	TRUE	"The mitot	212	294	17	20972	0.057823129	0.080188679
GO:BP	GO:00076 learning or	2.9E-07	TRUE	"The acqui	268	294	19	20972	0.06462585	0.070895522
GO:BP	GO:00447 mitotic cel	2.91E-07	TRUE	"The cell c	426	294	24	20972	0.081632653	0.056338028
GO:BP	GO:00072 cell-cell sig	2.91E-07	TRUE	"Any proce	1316	294	46	20972	0.156462585	0.034954407
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GO:BP	GO:00974: dendrite e	3E-07	TRUE	"Long dista	34	1172	13	20972	0.01109215	0.382352941
GO:BP	GO:00315: heterochro	3.08E-07	TRUE	"An epiger	162	294	15	20972	0.051020408	0.092592593
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GO:BP	GO:00971 receptor lc	5.93E-07	TRUE	"Any proce	77	1172	19	20972	0.016211604	0.246753247
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GO:BP	GO:00426 muscle cel	5.98E-07	TRUE	"The proce	388	1172	50	20972	0.042662116	0.128865979
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GO:BP	GO:00352 ionotropic	6.54E-07	TRUE	"The serie	30	227	7	20972	0.030837004	0.233333333
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GO:BP	GO:00481	regulation	6.66E-07	TRUE	"A process	221	682	26	20972	0.038123167	0.117647059
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GO:BP	GO:00019	startle res	8.44E-07	TRUE	"An action	30	682	10	20972	0.014662757	0.333333333
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GO:BP	GO:00097: response t	1.28E-06	TRUE	"Any proce	1467	682	87	20972	0.127565982	0.059304703
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GO:BP	GO:00510: positive re	1.42E-06	TRUE	"Any proce	312	682	31	20972	0.045454545	0.099358974
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GO:BP	GO:006501: regulation	2.41E-06	TRUE	"Any proce	1070	1172	101	20972	0.086177474	0.094392523
GO:BP	GO:004241: dopamine	2.42E-06	TRUE	"The chem	42	682	11	20972	0.016129032	0.261904762
GO:BP	GO:000191: postsynapt	2.43E-06	TRUE	"A process	40	1172	13	20972	0.01109215	0.325
GO:BP	GO:009891: excitatory	2.44E-06	TRUE	"Synaptic t	12	227	5	20972	0.022026432	0.416666667
GO:BP	GO:000301: vascular pr	2.51E-06	TRUE	"A circulat	270	682	28	20972	0.041055718	0.103703704
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GO:BP	GO:000991: positive re	2.58E-06	TRUE	"Any proce	1569	227	41	20972	0.18061674	0.026131294
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GO:BP	GO:00977! blood vess	3.87E-06	TRUE	"Any proce	152	682	20	20972	0.029325513	0.131578947
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GO:BP	GO:00447! mitotic cel	4.06E-06	TRUE	"The cell c	426	1172	51	20972	0.043515358	0.11971831
GO:BP	GO:00717! membrane	4.09E-06	TRUE	"The aggre	63	1172	16	20972	0.013651877	0.253968254
GO:BP	GO:00610! membrane	4.11E-06	TRUE	"A process	783	1172	79	20972	0.067406143	0.100893997
GO:BP	GO:00083! associative	4.13E-06	TRUE	"Learning l	87	682	15	20972	0.021994135	0.172413793
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GO:BP	GO:00456! negative re	4.34E-06	TRUE	"Any proce	71	1172	17	20972	0.014505119	0.23943662
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GO:BP	GO:00514! regulation	4.5E-06	TRUE	"Any proce	56	1172	15	20972	0.012798635	0.267857143
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GO:BP	GO:00603! head morp	4.71E-06	TRUE	"The proce	40	227	7	20972	0.030837004	0.175
GO:BP	GO:00519! regulation	4.78E-06	TRUE	"Any proce	163	227	12	20972	0.052863436	0.073619632
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GO:BP	GO:00002! microtubu	4.87E-06	TRUE	"A process	681	1172	71	20972	0.060580205	0.104258443
GO:BP	GO:00197! calcium-m	5.01E-06	TRUE	"Any intrac	234	1172	34	20972	0.029010239	0.145299145
GO:BP	GO:00068! sodium ior	5.1E-06	TRUE	"The direc	245	1172	35	20972	0.029863481	0.142857143
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GO:BP	GO:00518! regulation	5.62E-06	TRUE	"Any proce	268	1172	37	20972	0.031569966	0.138059701
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GO:BP	GO:00068! exocytosis	6E-06	TRUE	"A process	352	682	32	20972	0.046920821	0.090909091
GO:BP	GO:00020! regulation	6.05E-06	TRUE	"Any proce	102	682	16	20972	0.023460411	0.156862745
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GO:BP	GO:00488: cell projec	4.09E-05	TRUE	"The proce	669	172	20	20972	0.11627907	0.029895366
GO:BP	GO:00030: vascular pr	4.15E-05	TRUE	"A circulat	270	1172	35	20972	0.029863481	0.12962963
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GO:BP	GO:19901: neuron prc	4.17E-05	TRUE	"Long dista	169	227	11	20972	0.04845815	0.065088757
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GO:BP	GO:00218: forebrain r	4.29E-05	TRUE	"The proce	36	682	9	20972	0.013196481	0.25
GO:BP	GO:00459: positive re	4.34E-05	TRUE	"Any proce	242	227	13	20972	0.057268722	0.053719008
GO:BP	GO:00216: nerve deve	4.35E-05	TRUE	"The proce	93	1172	18	20972	0.015358362	0.193548387
GO:BP	GO:20001: regulation	4.35E-05	TRUE	"Any proce	93	1172	18	20972	0.015358362	0.193548387
GO:BP	GO:00482: vesicle doc	4.36E-05	TRUE	"The initial	59	1172	14	20972	0.011945392	0.237288136
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GO:BP	GO:00062: nucleotide	4.58E-05	TRUE	"A DNA rej	83	294	9	20972	0.030612245	0.108433735

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GO:BP	GO:00300: lymphocyt	5.25E-05	TRUE	"The proce	427	294	20	20972	0.068027211	0.046838407
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GO:BP	GO:00080: neuron rec	5.36E-05	TRUE	"The proce	47	682	10	20972	0.014662757	0.212765957
GO:BP	GO:00516: establishm	5.37E-05	TRUE	"Any proce	1938	172	37	20972	0.215116279	0.019091847
GO:BP	GO:00026: regulation	5.42E-05	TRUE	"Any proce	572	1172	59	20972	0.050341297	0.103146853
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GO:BP	GO:00458: negative re	5.42E-05	TRUE	"An epiger	195	1172	28	20972	0.023890785	0.143589744
GO:BP	GO:19038: positive re	5.52E-05	TRUE	"Any proce	433	682	34	20972	0.049853372	0.07852194
GO:BP	GO:00065: catecholar	5.52E-05	TRUE	"The chem	58	682	11	20972	0.016129032	0.189655172
GO:BP	GO:00097: catechol-c	5.52E-05	TRUE	"The chem	58	682	11	20972	0.016129032	0.189655172
GO:BP	GO:00718: potassium	5.6E-05	TRUE	"A process	199	682	21	20972	0.030791789	0.105527638
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GO:BP	GO:000701: actin filament assembly	5.77E-05	TRUE	"A process	457	1172	50	20972	0.042662116	0.10940919
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GO:BP	GO:003221: secretory	5.92E-05	TRUE	"Any process	25	1172	9	20972	0.007679181	0.36
GO:BP	GO:009871: inorganic	5.96E-05	TRUE	"Any process	445	1172	49	20972	0.041808874	0.11011236
GO:BP	GO:000631: double-strand	5.98E-05	TRUE	"The repair	322	1172	39	20972	0.033276451	0.121118012
GO:BP	GO:006091: positive re	6.01E-05	TRUE	"Any process	39	227	6	20972	0.026431718	0.153846154
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GO:BP	GO:200011: positive re	6.34E-05	TRUE	"Any process	53	1172	13	20972	0.01109215	0.245283019
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GO:BP	GO:000151: regulation	6.41E-05	TRUE	"Any process	399	172	15	20972	0.087209302	0.037593985
GO:BP	GO:009901: postsynap	6.44E-05	TRUE	"The aggreg	83	682	13	20972	0.019061584	0.156626506
GO:BP	GO:000761: circadian r	6.53E-05	TRUE	"Any biolo	201	682	21	20972	0.030791789	0.104477612
GO:BP	GO:006131: neural pre	6.57E-05	TRUE	"The multi	164	294	12	20972	0.040816327	0.073170732
GO:BP	GO:000831: associative	6.58E-05	TRUE	"Learning l	87	294	9	20972	0.030612245	0.103448276
GO:BP	GO:004801: regulation	6.68E-05	TRUE	"Any process	111	294	10	20972	0.034013605	0.09009009
GO:BP	GO:003281: cellular res	6.68E-05	TRUE	"Any process	677	294	26	20972	0.088435374	0.038404727
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GO:BP	GO:003331: protein loc	6.75E-05	TRUE	"A process	887	227	26	20972	0.114537445	0.029312289
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GO:BP	GO:00224' membrane	7.03E-05	TRUE	"The initial	88	294	9	20972	0.030612245	0.102272727
GO:BP	GO:00454' myoblast c	7.03E-05	TRUE	"The proce	112	294	10	20972	0.034013605	0.089285714
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GO:BP	GO:00015' blood vess	7.1E-05	TRUE	"The proce	727	682	48	20972	0.070381232	0.066024759
GO:BP	GO:00098' negative re	7.14E-05	TRUE	"Any proce	363	294	18	20972	0.06122449	0.049586777
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GO:BP	GO:00512' positive re	7.26E-05	TRUE	"Any proce	863	1172	80	20972	0.068259386	0.092699884
GO:BP	GO:00315' cell-substr	7.29E-05	TRUE	"The attac	349	1172	41	20972	0.034982935	0.11747851
GO:BP	GO:00604' epithelium	7.29E-05	TRUE	"The proce	1234	1172	106	20972	0.090443686	0.085899514
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GO:BP	GO:00160' cell growth	7.54E-05	TRUE	"The proce	478	294	21	20972	0.071428571	0.043933054
GO:BP	GO:00101' body morp	7.65E-05	TRUE	"The proce	48	294	7	20972	0.023809524	0.145833333
GO:BP	GO:00519' catecholar	7.66E-05	TRUE	"The direc	60	682	11	20972	0.016129032	0.183333333
GO:BP	GO:00075' muscle org	7.69E-05	TRUE	"The proce	362	1172	42	20972	0.035836177	0.116022099
GO:BP	GO:00015' action pot	7.75E-05	TRUE	"A process	170	172	10	20972	0.058139535	0.058823529
GO:BP	GO:00362' response t	7.76E-05	TRUE	"Any proce	290	1172	36	20972	0.030716724	0.124137931
GO:BP	GO:00069' apoptotic	7.82E-05	TRUE	"A prograr	1898	294	51	20972	0.173469388	0.02687039
GO:BP	GO:00217' hippocam	7.92E-05	TRUE	"The progr	88	1172	17	20972	0.014505119	0.193181818
GO:BP	GO:00224' membrane	7.92E-05	TRUE	"The initial	88	1172	17	20972	0.014505119	0.193181818
GO:BP	GO:00457' positive re	7.93E-05	TRUE	"Any proce	140	294	11	20972	0.037414966	0.078571429
GO:BP	GO:00971' GABAergic	7.94E-05	TRUE	"The proce	20	1172	8	20972	0.006825939	0.4
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GO:BP	GO:00486' cell morph	8.3E-05	TRUE	"The proce	584	172	18	20972	0.104651163	0.030821918
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GO:BP	GO:00215' pallium de	8.33E-05	TRUE	"The proce	188	682	20	20972	0.029325513	0.106382979
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GO:BP	GO:005151: positive re	8.35E-05	TRUE	"Any proce	26	1172	9	20972	0.007679181	0.346153846
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GO:BP	GO:009871: maintenanc	9.09E-05	TRUE	"Any proce	185	227	11	20972	0.04845815	0.059459459
GO:BP	GO:001581: acidic amir	9.09E-05	TRUE	"The direc	80	1172	16	20972	0.013651877	0.2
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GO:BP	GO:190581: regulation	9.7E-05	TRUE	"Any proce	22	682	7	20972	0.01026393	0.318181818
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GO:BP	GO:007051: regulation	9.77E-05	TRUE	"Any proce	169	1172	25	20972	0.021331058	0.147928994
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GO:BP	GO:004251: myelinatio	9.86E-05	TRUE	"The proce	152	227	10	20972	0.044052863	0.065789474
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GO:BP	GO:200001: regulation	9.91E-05	TRUE	"Any proce	70	294	8	20972	0.027210884	0.114285714
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GO:BP	GO:190491: positive re	0.000102	TRUE	"Any proce	282	1172	35	20972	0.029863481	0.124113475
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GO:BP	GO:00507	positive re	0.000103	TRUE	"Any proce	72	1172	15	20972	0.012798635	0.208333333
GO:BP	GO:00099	hormone t	0.000103	TRUE	"The direc	330	1172	39	20972	0.033276451	0.118181818
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GO:BP	GO:00481	behavioral	0.000104	TRUE	"Any proce	15	682	6	20972	0.008797654	0.4
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GO:BP	GO:00072	ensheathm	0.000109	TRUE	"The proce	154	227	10	20972	0.044052863	0.064935065
GO:BP	GO:00083	axon ensh	0.000109	TRUE	"Any proce	154	227	10	20972	0.044052863	0.064935065
GO:BP	GO:00017	in utero er	0.000109	TRUE	"The proce	405	1172	45	20972	0.038395904	0.111111111
GO:BP	GO:00160	cell growth	0.000109	TRUE	"The proce	478	172	16	20972	0.093023256	0.033472803
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GO:BP	GO:008601 cardiac muscle tissue morphogenesis	0.000427	TRUE	"An action"	83	227	7	20972	0.030837004	0.084337349
GO:BP	GO:200071 regulation of gene expression	0.000427	TRUE	"Any process"	147	227	9	20972	0.039647577	0.06122449
GO:BP	GO:004001 regulation of gene expression	0.000427	TRUE	"Any process"	590	682	39	20972	0.057184751	0.066101695
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GO:BP	GO:009911 presynaptic transmission	0.000429	TRUE	"Any process"	38	682	8	20972	0.011730205	0.210526316
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GO:BP	GO:199001 response to hypoxia	0.000432	TRUE	"A process"	47	1172	11	20972	0.009385666	0.234042553
GO:BP	GO:004511 establishment of cell-cell contact	0.000432	TRUE	"Any cellular process"	47	1172	11	20972	0.009385666	0.234042553

GO:BP	GO:19017!response t	0.000433	TRUE	"Any proce	1632	227	36	20972	0.158590308	0.022058824
GO:BP	GO:00488!inner ear c	0.000434	TRUE	"The proce	209	1172	27	20972	0.023037543	0.129186603
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GO:BP	GO:00074!synapse as	0.000447	TRUE	"The aggre	266	172	11	20972	0.063953488	0.041353383
GO:BP	GO:00064!N-terminal	0.00045	TRUE	"The acety	14	294	4	20972	0.013605442	0.285714286
GO:BP	GO:00601!prepulse ir	0.00045	TRUE	"The proce	14	294	4	20972	0.013605442	0.285714286
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GO:BP	GO:00613!neural pre	0.000453	TRUE	"The multi	164	682	17	20972	0.024926686	0.103658537
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GO:BP	GO:00071!cell-matrix	0.000458	TRUE	"The bindi	233	1172	29	20972	0.024744027	0.124463519
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GO:BP	GO:00610!positive re	0.000459	TRUE	"Any proce	19	1172	7	20972	0.005972696	0.368421053
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GO:BP	GO:00100!response t	0.000471	TRUE	"Any proce	320	172	12	20972	0.069767442	0.0375
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GO:BP	GO:00425: fear respo	0.000501	TRUE	"The respc	50	682	9	20972	0.013196481	0.18
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GO:BP	GO:00030: regionaliza	0.00052	TRUE	"The patte	422	1172	44	20972	0.037542662	0.104265403
GO:BP	GO:00382: TORC1 sig	0.000522	TRUE	"A series o	113	1172	18	20972	0.015358362	0.159292035
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GO:BP	GO:00216: cerebellar	0.000565	TRUE	"The proce	26	1172	8	20972	0.006825939	0.307692308
GO:BP	GO:00480: neurotrop	0.000565	TRUE	"The serie	26	1172	8	20972	0.006825939	0.307692308
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GO:BP	GO:00487: calcium io	0.000581	TRUE	"The relea	15	294	4	20972	0.013605442	0.266666667
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GO:BP	GO:00343: protein loc	0.000594	TRUE	"A process	66	1172	13	20972	0.01109215	0.196969697
GO:BP	GO:00063: chromatin	0.000596	TRUE	"A dynami	1057	172	23	20972	0.13372093	0.021759697
GO:BP	GO:00305: intracellu	0.0006	TRUE	"The serie	399	1172	42	20972	0.035836177	0.105263158
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GO:BP	GO:00230: negative re	0.000601	TRUE	"Any proce	1449	172	28	20972	0.162790698	0.019323671
GO:BP	GO:00988: modulatio	0.000602	TRUE	"Any proce	47	294	6	20972	0.020408163	0.127659574
GO:BP	GO:00027: peptide se	0.000602	TRUE	"The contr	256	682	22	20972	0.032258065	0.0859375
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GO:BP	GO:01060: neuron pr	0.000636	TRUE	"A process	91	682	12	20972	0.017595308	0.131868132
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GO:BP	GO:00300! metal ion t	0.000749	TRUE	"The direc	860	227	23	20972	0.101321586	0.026744186
GO:BP	GO:00400! positive re	0.000749	TRUE	"Any proce	586	682	38	20972	0.055718475	0.064846416
GO:BP	GO:19002! regulation	0.00075	TRUE	"Any proce	42	1172	10	20972	0.008532423	0.238095238
GO:BP	GO:01500! regulation	0.00075	TRUE	"Any proce	42	1172	10	20972	0.008532423	0.238095238
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GO:BP	GO:00600! cardiac mu	0.000771	TRUE	"Muscle co	139	682	15	20972	0.021994135	0.107913669
GO:BP	GO:00352! organ grov	0.000779	TRUE	"The incre	160	1172	22	20972	0.018771331	0.1375
GO:BP	GO:00009! cytokines	0.00078	TRUE	"The divisi	194	1172	25	20972	0.021331058	0.128865979
GO:BP	GO:01411! positive re	0.000781	TRUE	"An epiger	59	1172	12	20972	0.010238908	0.203389831
GO:BP	GO:00330! regulation	0.000786	TRUE	"Any proce	241	227	11	20972	0.04845815	0.045643154
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GO:BP	GO:00322! negative re	0.000788	TRUE	"Any proce	31	682	7	20972	0.01026393	0.225806452
GO:BP	GO:00468! hormone s	0.000793	TRUE	"The regul	319	682	25	20972	0.036656891	0.078369906
GO:BP	GO:00716! protein loc	0.0008	TRUE	"Any proce	366	1172	39	20972	0.033276451	0.106557377
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GO:BP	GO:00070! metaphase	0.0008	TRUE	"The cell c	97	294	8	20972	0.027210884	0.082474227
GO:BP	GO:00512! protein ho	0.000804	TRUE	"The proce	206	1172	26	20972	0.0221843	0.126213592
GO:BP	GO:00486! positive re	0.000806	TRUE	"Any proce	156	682	16	20972	0.023460411	0.102564103
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GO:BP	GO:00217! cerebral cc	0.000816	TRUE	"The order	50	294	6	20972	0.020408163	0.12
GO:BP	GO:00085! visual learn	0.000816	TRUE	"Any proce	50	294	6	20972	0.020408163	0.12
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GO:BP	GO:00158! organic hy	0.000817	TRUE	"The direc	266	1172	31	20972	0.026450512	0.116541353
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GO:BP	GO:00609! dendritic s	0.000832	TRUE	"The proce	94	682	12	20972	0.017595308	0.127659574
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GO:BP	GO:00158! peptide tra	0.000839	TRUE	"The direc	282	682	23	20972	0.03372434	0.081560284
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GO:BP	GO:19021! regulation	0.000857	TRUE	"Any proce	329	1172	36	20972	0.030716724	0.109422492
GO:BP	GO:00009! nuclear-tra	0.000863	TRUE	"The chem	126	294	9	20972	0.030612245	0.071428571
GO:BP	GO:00320! negative re	0.000866	TRUE	"Any proce	73	294	7	20972	0.023809524	0.095890411
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GO:BP	GO:00063! transcripti	0.000869	TRUE	"The exter	118	1172	18	20972	0.015358362	0.152542373
GO:BP	GO:00714! cellular res	0.00087	TRUE	"Any proce	1238	227	29	20972	0.127753304	0.023424879
GO:BP	GO:00080! retrograde	0.000874	TRUE	"The direc	21	1172	7	20972	0.005972696	0.333333333
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GO:BP	GO:00063! nucleosom	0.000874	TRUE	"The contr	21	1172	7	20972	0.005972696	0.333333333
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GO:BP	GO:00487! calcium io	0.000874	TRUE	"The relea	15	1172	6	20972	0.005119454	0.4
GO:BP	GO:00481! regulation	0.000874	TRUE	"A process	15	1172	6	20972	0.005119454	0.4
GO:BP	GO:00301! regulation	0.000881	TRUE	"Any proce	292	1172	33	20972	0.028156997	0.113013699
GO:BP	GO:00447! cell cycle p	0.000888	TRUE	"The cell c	534	227	17	20972	0.074889868	0.031835206
GO:BP	GO:00905! sensory or	0.000888	TRUE	"Morphog	294	294	14	20972	0.047619048	0.047619048
GO:BP	GO:00315! protein-co	0.0009	TRUE	"A localiza	88	1172	15	20972	0.012798635	0.170454545
GO:BP	GO:00335! multicellul	0.000901	TRUE	"Any proce	95	227	7	20972	0.030837004	0.073684211
GO:BP	GO:00093! response t	0.000901	TRUE	"Any proce	432	227	15	20972	0.066079295	0.034722222
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GO:BP	GO:00161!	vesicle-me	0.000904	TRUE	"A cellular	1572	172	29	20972	0.168604651	0.018447837
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GO:BP	GO:00459!	positive re	0.000912	TRUE	"Any proce	242	172	10	20972	0.058139535	0.041322314
GO:BP	GO:00072!	neurotran	0.000912	TRUE	"The regul	149	172	8	20972	0.046511628	0.053691275
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GO:BP	GO:00511!	meiotic sis	0.000913	TRUE	"The cell c	17	294	4	20972	0.013605442	0.235294118
GO:BP	GO:20001!	positive re	0.000915	TRUE	"Any proce	571	682	37	20972	0.054252199	0.064798599
GO:BP	GO:00072!	phospholiq	0.000921	TRUE	"A G prote	140	1172	20	20972	0.017064846	0.142857143
GO:BP	GO:00432!	regulation	0.000928	TRUE	"Any proce	81	682	11	20972	0.016129032	0.135802469
GO:BP	GO:00071!	adenylate	0.000928	TRUE	"A G prote	81	682	11	20972	0.016129032	0.135802469
GO:BP	GO:00902!	regulation	0.000934	TRUE	"Any proce	318	1172	35	20972	0.029863481	0.110062893
GO:BP	GO:00011!	response t	0.000943	TRUE	"Any proce	129	227	8	20972	0.035242291	0.062015504
GO:BP	GO:00019!	synaptic tr	0.000944	TRUE	"The vesic	28	1172	8	20972	0.006825939	0.285714286
GO:BP	GO:20008!	regulation	0.000944	TRUE	"Any proce	28	1172	8	20972	0.006825939	0.285714286
GO:BP	GO:00301!	regulation	0.000951	TRUE	"Any proce	331	1172	36	20972	0.030716724	0.108761329
GO:BP	GO:00421!	regulation	0.000955	TRUE	"Any proce	1639	227	35	20972	0.154185022	0.021354484
GO:BP	GO:00486!	neuron fat	0.000957	TRUE	"The proce	10	1172	5	20972	0.004266212	0.5
GO:BP	GO:00709!	RISC comp	0.000957	TRUE	"The proce	10	1172	5	20972	0.004266212	0.5
GO:BP	GO:00506!	epithelial c	0.000957	TRUE	"The multi	435	1172	44	20972	0.037542662	0.101149425
GO:BP	GO:00447!	metaphase	0.000965	TRUE	"The cell c	100	294	8	20972	0.027210884	0.08
GO:BP	GO:00190!	reproducti	0.000966	TRUE	"The speci	43	682	8	20972	0.011730205	0.186046512
GO:BP	GO:00423!	vasodilatic	0.000973	TRUE	"An increa	55	682	9	20972	0.013196481	0.163636364
GO:BP	GO:00514!	negative re	0.000973	TRUE	"Any proce	159	682	16	20972	0.023460411	0.100628931
GO:BP	GO:00512!	regulation	0.000976	TRUE	"Any proce	1574	227	34	20972	0.149779736	0.021601017
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GO:BP	GO:20012!	regulation	0.000994	TRUE	"Any proce	52	294	6	20972	0.020408163	0.115384615
GO:BP	GO:00511!	nuclear tra	0.001005	TRUE	"The direc	332	1172	36	20972	0.030716724	0.108433735
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GO:BP	GO:20007!	positive re	0.001011	TRUE	"Any proce	97	227	7	20972	0.030837004	0.072164948
GO:BP	GO:00434!	regulation	0.001015	TRUE	"Any proce	186	1172	24	20972	0.020477816	0.129032258

GO:BP	GO:003501	establishment of	0.001016	TRUE	"Any cellular process"	52	1172	11	20972	0.009385666	0.211538462
GO:BP	GO:003271	regulation of	0.001016	TRUE	"Any process"	99	1172	16	20972	0.013651877	0.161616162
GO:BP	GO:006121	establishment of	0.001016	TRUE	"Any cellular process"	52	1172	11	20972	0.009385666	0.211538462
GO:BP	GO:004001	negative regulation of	0.001016	TRUE	"Any process"	325	682	25	20972	0.036656891	0.076923077
GO:BP	GO:000711	cell adhesion	0.001026	TRUE	"The attachment of cells"	1512	227	33	20972	0.145374449	0.021825397
GO:BP	GO:002261	regulation of	0.00103	TRUE	"Any process"	822	227	22	20972	0.0969163	0.02676399
GO:BP	GO:005071	RNA destination	0.00103	TRUE	"Any process"	101	294	8	20972	0.027210884	0.079207921
GO:BP	GO:190161	cellular response to	0.001031	TRUE	"Any process"	676	294	23	20972	0.078231293	0.034023669
GO:BP	GO:009901	vesicle-mediated	0.001032	TRUE	"Any vesicle transport"	246	172	10	20972	0.058139535	0.040650407
GO:BP	GO:006031	face morphology	0.001034	TRUE	"The process of morphogenesis"	33	294	5	20972	0.017006803	0.151515152
GO:BP	GO:005121	regulation of	0.001034	TRUE	"Any process"	1574	294	41	20972	0.139455782	0.026048285
GO:BP	GO:006021	long-term	0.001034	TRUE	"A process"	33	294	5	20972	0.017006803	0.151515152
GO:BP	GO:009051	sensory response	0.001037	TRUE	"Morphogenesis"	294	227	12	20972	0.052863436	0.040816327
GO:BP	GO:190291	regulation of	0.001047	TRUE	"Any process"	377	294	16	20972	0.054421769	0.042440318
GO:BP	GO:001061	negative regulation of	0.001055	TRUE	"Any process"	1449	227	32	20972	0.140969163	0.022084196
GO:BP	GO:002301	negative regulation of	0.001055	TRUE	"Any process"	1449	227	32	20972	0.140969163	0.022084196
GO:BP	GO:004001	positive regulation of	0.001056	TRUE	"Any process"	586	1172	55	20972	0.046928328	0.093856655
GO:BP	GO:007121	cellular response to	0.001059	TRUE	"Any process"	333	1172	36	20972	0.030716724	0.108108108
GO:BP	GO:010401	cellular response to	0.001059	TRUE	"Any process"	333	1172	36	20972	0.030716724	0.108108108
GO:BP	GO:000301	heart morphogenesis	0.001064	TRUE	"The development of the heart"	263	294	13	20972	0.044217687	0.049429658
GO:BP	GO:190481	regulation of	0.001074	TRUE	"Any process"	23	682	6	20972	0.008797654	0.260869565
GO:BP	GO:001581	norepinephrine	0.001074	TRUE	"The direction of movement"	23	682	6	20972	0.008797654	0.260869565
GO:BP	GO:000931	amine metabolism	0.001074	TRUE	"The chemical reaction"	97	682	12	20972	0.017595308	0.12371134
GO:BP	GO:000971	embryo development	0.001074	TRUE	"The process of development"	687	682	42	20972	0.061583578	0.061135371
GO:BP	GO:000971	response to	0.001075	TRUE	"Any process"	222	1172	27	20972	0.023037543	0.121621622
GO:BP	GO:005081	negative regulation of	0.001077	TRUE	"Any process"	53	294	6	20972	0.020408163	0.113207547
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GO:BP	GO:007031	regulation of	0.00108	TRUE	"A cell cycle"	36	1172	9	20972	0.007679181	0.25
GO:BP	GO:003111	regulation of	0.00108	TRUE	"Any process"	36	1172	9	20972	0.007679181	0.25
GO:BP	GO:190541	regulation of	0.00108	TRUE	"Any process"	169	227	9	20972	0.039647577	0.053254438
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GO:BP	GO:002151	cerebellum	0.001086	TRUE	"The process of development"	44	1172	10	20972	0.008532423	0.227272727

GO:BP	GO:00091: purine nuc	0.001086	TRUE	"The chem	44	1172	10	20972	0.008532423	0.227272727
GO:BP	GO:00351: limb morp	0.001092	TRUE	"The proce	153	1172	21	20972	0.017918089	0.137254902
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GO:BP	GO:00082: cell popula	0.001099	TRUE	"The multi	1961	294	48	20972	0.163265306	0.024477307
GO:BP	GO:00515: response t	0.001104	TRUE	"Any proce	131	294	9	20972	0.030612245	0.06870229
GO:BP	GO:00718: potassium	0.001104	TRUE	"A process	199	172	9	20972	0.052325581	0.045226131
GO:BP	GO:00323: intracellu	0.001104	TRUE	"The direc	47	172	5	20972	0.029069767	0.106382979
GO:BP	GO:00071: cell adhesi	0.001104	TRUE	"The attac	1512	172	28	20972	0.162790698	0.018518519
GO:BP	GO:00148: regulation	0.001105	TRUE	"Any proce	18	294	4	20972	0.013605442	0.222222222
GO:BP	GO:00459: positive re	0.001105	TRUE	"Any proce	18	294	4	20972	0.013605442	0.222222222
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GO:BP	GO:00091: purine nuc	0.001109	TRUE	"The chem	44	682	8	20972	0.011730205	0.181818182
GO:BP	GO:00504: catecholar	0.001109	TRUE	"The regul	44	682	8	20972	0.011730205	0.181818182
GO:BP	GO:00100: glial cell di	0.00111	TRUE	"The proce	249	172	10	20972	0.058139535	0.040160643
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GO:BP	GO:00974: dendrite e	0.001155	TRUE	"Long dista	34	294	5	20972	0.017006803	0.147058824
GO:BP	GO:00164: cell migrat	0.001155	TRUE	"The contr	1532	294	40	20972	0.136054422	0.026109661
GO:BP	GO:00109: regulation	0.001161	TRUE	"Any proce	342	294	15	20972	0.051020408	0.043859649
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GO:BP	GO:00328: regulation	0.001168	TRUE	"Any proce	272	1172	31	20972	0.026450512	0.113970588
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GO:BP	GO:00071: leukocyte	0.001175	TRUE	"The attac	413	1172	42	20972	0.035836177	0.101694915
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GO:BP	GO:00362: protein mc	0.001187	TRUE	"The coval	2360	227	45	20972	0.198237885	0.019067797
GO:BP	GO:00217: developme	0.001187	TRUE	"A develop	346	227	13	20972	0.057268722	0.037572254
GO:BP	GO:00860: membrane	0.001188	TRUE	"The proce	53	1172	11	20972	0.009385666	0.20754717
GO:BP	GO:00443: cell-cell ad	0.001188	TRUE	"The attac	53	1172	11	20972	0.009385666	0.20754717
GO:BP	GO:00216: cerebellar	0.001188	TRUE	"The proce	53	1172	11	20972	0.009385666	0.20754717
GO:BP	GO:19026: regulation	0.001191	TRUE	"Any proce	29	1172	8	20972	0.006825939	0.275862069
GO:BP	GO:00220: telenceph	0.001191	TRUE	"The order	29	1172	8	20972	0.006825939	0.275862069
GO:BP	GO:00218: cerebral cc	0.001191	TRUE	"The radia	29	1172	8	20972	0.006825939	0.275862069

GO:BP	GO:00357 endothelia	0.001191	TRUE	"The direc	29	1172	8	20972	0.006825939	0.275862069
GO:BP	GO:00990 presynapsi	0.001192	TRUE	"The aggre	44	227	5	20972	0.022026432	0.113636364
GO:BP	GO:00487 oligodendr	0.001194	TRUE	"The proce	104	294	8	20972	0.027210884	0.076923077
GO:BP	GO:00432 response t	0.001195	TRUE	"Any proce	15	682	5	20972	0.007331378	0.333333333
GO:BP	GO:00140 response t	0.001195	TRUE	"Any proce	15	682	5	20972	0.007331378	0.333333333
GO:BP	GO:00072 G protein-i	0.001195	TRUE	"A G prote	15	682	5	20972	0.007331378	0.333333333
GO:BP	GO:20000 regulation	0.001203	TRUE	"Any proce	70	227	6	20972	0.026431718	0.085714286
GO:BP	GO:00508 regulation	0.001208	TRUE	"Any proce	305	172	11	20972	0.063953488	0.036065574
GO:BP	GO:00702 actin-medi	0.001208	TRUE	"The actin	115	172	7	20972	0.040697674	0.060869565
GO:BP	GO:00328 cellular res	0.001208	TRUE	"Any proce	252	172	10	20972	0.058139535	0.03968254
GO:BP	GO:00075 skeletal m	0.001214	TRUE	"The devel	177	1172	23	20972	0.019624573	0.129943503
GO:BP	GO:00430 chordate e	0.001218	TRUE	"The proce	669	682	41	20972	0.060117302	0.061285501
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GO:BP	GO:00314: positive re	0.005452	TRUE	"Any proce	450	1172	42	20972	0.035836177	0.093333333
GO:BP	GO:19016: response t	0.005452	TRUE	"A respons	205	172	8	20972	0.046511628	0.03902439
GO:BP	GO:00016: behavioral	0.00548	TRUE	"An acute	44	682	7	20972	0.01026393	0.159090909
GO:BP	GO:00096: response t	0.005497	TRUE	"Any proce	1108	172	21	20972	0.122093023	0.018953069
GO:BP	GO:00217: limbic syst	0.005503	TRUE	"The progr	119	682	12	20972	0.017595308	0.100840336
GO:BP	GO:20001: negative re	0.005515	TRUE	"Any proce	308	682	22	20972	0.032258065	0.071428571
GO:BP	GO:00063: transcripti	0.005523	TRUE	"The synth	76	294	6	20972	0.020408163	0.078947368
GO:BP	GO:00506: epithelial c	0.005543	TRUE	"The multi	435	682	28	20972	0.041055718	0.064367816

GO:BP	GO:00704: response t	0.005547	TRUE	"Any proce	317	227	11	20972	0.04845815	0.034700315
GO:BP	GO:19033: positive re	0.005547	TRUE	"Any proce	136	227	7	20972	0.030837004	0.051470588
GO:BP	GO:00485: camera-ty	0.005547	TRUE	"The proce	136	227	7	20972	0.030837004	0.051470588
GO:BP	GO:00068: calcium ior	0.005549	TRUE	"The direc	421	227	13	20972	0.057268722	0.03087886
GO:BP	GO:00605: muscle tiss	0.005549	TRUE	"The progr	421	227	13	20972	0.057268722	0.03087886
GO:BP	GO:00070: actin filam	0.005571	TRUE	"A process	457	682	29	20972	0.042521994	0.06345733
GO:BP	GO:00512: protein ho	0.005579	TRUE	"The proce	206	172	8	20972	0.046511628	0.038834951
GO:BP	GO:00602: positive re	0.005582	TRUE	"Any proce	13	294	3	20972	0.010204082	0.230769231
GO:BP	GO:00608: random in	0.005582	TRUE	"Compens	13	294	3	20972	0.010204082	0.230769231
GO:BP	GO:00458: positive re	0.005603	TRUE	"Any proce	154	682	14	20972	0.020527859	0.090909091
GO:BP	GO:00425: myelinatio	0.005606	TRUE	"The proce	152	1172	19	20972	0.016211604	0.125
GO:BP	GO:00163: morphoge	0.005606	TRUE	"The proce	152	1172	19	20972	0.016211604	0.125
GO:BP	GO:00720: kidney epi	0.005606	TRUE	"The proce	152	1172	19	20972	0.016211604	0.125
GO:BP	GO:00354: protein loc	0.005617	TRUE	"Any proce	87	682	10	20972	0.014662757	0.114942529
GO:BP	GO:00430: regulation	0.005628	TRUE	"Any proce	188	1172	22	20972	0.018771331	0.117021277
GO:BP	GO:19034: regulation	0.005651	TRUE	"Any proce	99	227	6	20972	0.026431718	0.060606061
GO:BP	GO:00030: regionaliza	0.005651	TRUE	"The patte	422	227	13	20972	0.057268722	0.030805687
GO:BP	GO:00068: sodium ior	0.005687	TRUE	"The direc	245	294	11	20972	0.037414966	0.044897959
GO:BP	GO:00708: response t	0.005687	TRUE	"Any proce	734	294	22	20972	0.074829932	0.029972752
GO:BP	GO:19024: L-alpha-arr	0.005691	TRUE	"The direc	72	682	9	20972	0.013196481	0.125
GO:BP	GO:00217: cerebral cc	0.005702	TRUE	"The migr	38	227	4	20972	0.017621145	0.105263158
GO:BP	GO:00356: exploratio	0.005702	TRUE	"The speci	38	227	4	20972	0.017621145	0.105263158
GO:BP	GO:00218: forebrain c	0.005709	TRUE	"The order	66	227	5	20972	0.022026432	0.075757576
GO:BP	GO:00343: protein loc	0.005709	TRUE	"A process	66	227	5	20972	0.022026432	0.075757576
GO:BP	GO:00611: morphoge	0.005717	TRUE	"The proce	191	682	16	20972	0.023460411	0.083769634
GO:BP	GO:00711: protein loc	0.005731	TRUE	"Any proce	51	294	5	20972	0.017006803	0.098039216
GO:BP	GO:00713: cellular res	0.005744	TRUE	"Any proce	210	682	17	20972	0.024926686	0.080952381
GO:BP	GO:00094: response t	0.005745	TRUE	"Any proce	319	227	11	20972	0.04845815	0.034482759
GO:BP	GO:00083: regulation	0.005785	TRUE	"Any proce	137	682	13	20972	0.019061584	0.094890511
GO:BP	GO:00459: positive re	0.0058	TRUE	"Any proce	46	1172	9	20972	0.007679181	0.195652174
GO:BP	GO:19043: positive re	0.005813	TRUE	"Any proce	75	1172	12	20972	0.010238908	0.16
GO:BP	GO:00070: regulation	0.005815	TRUE	"Any proce	17	227	3	20972	0.013215859	0.176470588

GO:BP	GO:00218: layer form	0.005815	TRUE	"The detach	17	227	3	20972	0.013215859	0.176470588
GO:BP	GO:19018: positive re	0.005815	TRUE	"Any proce	106	294	7	20972	0.023809524	0.066037736
GO:BP	GO:00712: cellular res	0.005815	TRUE	"Any proce	77	294	6	20972	0.020408163	0.077922078
GO:BP	GO:00324: positive re	0.005815	TRUE	"Any proce	77	294	6	20972	0.020408163	0.077922078
GO:BP	GO:00435: regulation	0.005815	TRUE	"Any proce	77	294	6	20972	0.020408163	0.077922078
GO:BP	GO:00345: protein loc	0.005815	TRUE	"Any proce	106	294	7	20972	0.023809524	0.066037736
GO:BP	GO:00026: positive re	0.005828	TRUE	"Any proce	1111	1172	87	20972	0.074232082	0.078307831
GO:BP	GO:00069: vesicle tar	0.005828	TRUE	"The proce	65	1172	11	20972	0.009385666	0.169230769
GO:BP	GO:19030: regulation	0.005843	TRUE	"Any proce	58	682	8	20972	0.011730205	0.137931034
GO:BP	GO:00481: regulation	0.005843	TRUE	"A process	58	682	8	20972	0.011730205	0.137931034
GO:BP	GO:00216: cerebellar	0.005844	TRUE	"The proce	29	1172	7	20972	0.005972696	0.24137931
GO:BP	GO:00480: platelet-de	0.005871	TRUE	"The serie	107	1172	15	20972	0.012798635	0.140186916
GO:BP	GO:00330: regulation	0.005871	TRUE	"Any proce	107	1172	15	20972	0.012798635	0.140186916
GO:BP	GO:00447: metaphase	0.005871	TRUE	"The cell c	100	227	6	20972	0.026431718	0.06
GO:BP	GO:00030: developm	0.005873	TRUE	"A develop	1024	227	23	20972	0.101321586	0.022460938
GO:BP	GO:00509: regulation	0.005882	TRUE	"Any proce	201	1172	23	20972	0.019624573	0.114427861
GO:BP	GO:19037: regulation	0.005922	TRUE	"Any proce	425	227	13	20972	0.057268722	0.030588235
GO:BP	GO:00098: positive re	0.005928	TRUE	"Any proce	536	227	15	20972	0.066079295	0.027985075
GO:BP	GO:00512: positive re	0.005953	TRUE	"Any proce	331	682	23	20972	0.03372434	0.069486405
GO:BP	GO:00420: gliogenesis	0.005953	TRUE	"The proce	331	682	23	20972	0.03372434	0.069486405
GO:BP	GO:00429: regulation	0.005962	TRUE	"Any proce	1426	227	29	20972	0.127753304	0.020336606
GO:BP	GO:00068: intracellul	0.005982	TRUE	"The direc	653	227	17	20972	0.074889868	0.026033691
GO:BP	GO:00430: negative re	0.005995	TRUE	"Any proce	887	294	25	20972	0.085034014	0.028184893
GO:BP	GO:00487: cardiac mu	0.006029	TRUE	"The proce	225	227	9	20972	0.039647577	0.04
GO:BP	GO:00102: vascular tr	0.006039	TRUE	"The direc	88	682	10	20972	0.014662757	0.113636364
GO:BP	GO:01501: transport	0.006039	TRUE	"The direc	88	682	10	20972	0.014662757	0.113636364
GO:BP	GO:00434: response t	0.006047	TRUE	"Any proce	457	294	16	20972	0.054421769	0.035010941
GO:BP	GO:00108: regulation	0.006047	TRUE	"Any proce	30	294	4	20972	0.013605442	0.133333333
GO:BP	GO:00033: type B pan	0.006047	TRUE	"The proce	30	294	4	20972	0.013605442	0.133333333
GO:BP	GO:00352: tube devel	0.006047	TRUE	"The proce	1094	294	29	20972	0.098639456	0.026508227
GO:BP	GO:00160: synaptic ve	0.006066	TRUE	"Fusion of	101	227	6	20972	0.026431718	0.059405941
GO:BP	GO:00600: cardiac mu	0.006066	TRUE	"Muscle co	139	227	7	20972	0.030837004	0.050359712

GO:BP	GO:000181: neural tube morphogenesis	0.006066	TRUE	"The formation of the neural tube from the neural plate." (GO:000181)	101	227	6	20972	0.026431718	0.059405941
GO:BP	GO:001024: response to stress	0.006074	TRUE	"Any process that is a response to stress." (GO:001024)	17	172	3	20972	0.01744186	0.176470588
GO:BP	GO:002241: positive regulation of gene expression	0.006077	TRUE	"Any process that is a positive regulation of gene expression." (GO:002241)	328	294	13	20972	0.044217687	0.039634146
GO:BP	GO:000301: circulatory system process	0.006077	TRUE	"An organ system process." (GO:000301)	595	294	19	20972	0.06462585	0.031932773
GO:BP	GO:000221: behavioral process	0.006091	TRUE	"A behavior." (GO:000221)	45	682	7	20972	0.01026393	0.155555556
GO:BP	GO:000761: rhythmic behavior	0.006091	TRUE	"The specification of a rhythmic behavior." (GO:000761)	45	682	7	20972	0.01026393	0.155555556
GO:BP	GO:005091: negative regulation of gene expression	0.006091	TRUE	"The direct negative regulation of gene expression." (GO:005091)	45	682	7	20972	0.01026393	0.155555556
GO:BP	GO:003571: sodium ion transport	0.006109	TRUE	"A process that involves the transport of sodium ions." (GO:003571)	181	227	8	20972	0.035242291	0.044198895
GO:BP	GO:000691: striated muscle process	0.006109	TRUE	"A process that involves the contraction of striated muscle." (GO:000691)	181	227	8	20972	0.035242291	0.044198895
GO:BP	GO:005081: regulation of gene expression	0.006119	TRUE	"Any process that is a regulation of gene expression." (GO:005081)	374	682	25	20972	0.036656891	0.06684492
GO:BP	GO:004591: negative regulation of gene expression	0.006126	TRUE	"Any process that is a negative regulation of gene expression." (GO:004591)	52	294	5	20972	0.017006803	0.096153846
GO:BP	GO:007131: cellular response to stress	0.006127	TRUE	"Any process that is a cellular response to stress." (GO:007131)	210	294	10	20972	0.034013605	0.047619048
GO:BP	GO:005071: regulation of gene expression	0.006149	TRUE	"Any process that is a regulation of gene expression." (GO:005071)	73	682	9	20972	0.013196481	0.123287671
GO:BP	GO:004211: negative regulation of gene expression	0.006153	TRUE	"Any process that is a negative regulation of gene expression." (GO:004211)	130	1172	17	20972	0.014505119	0.130769231
GO:BP	GO:001811: peptidyl-aminopeptidase activity	0.006184	TRUE	"The alteration of a peptide by the action of a peptidyl-aminopeptidase." (GO:001811)	495	1172	45	20972	0.038395904	0.090909091
GO:BP	GO:003001: actin cytoskeleton organization	0.006204	TRUE	"A process that involves the organization of the actin cytoskeleton." (GO:003001)	716	227	18	20972	0.079295154	0.025139665
GO:BP	GO:001481: muscle hypertrophy	0.006208	TRUE	"The muscle hypertrophy." (GO:001481)	86	1172	13	20972	0.01109215	0.151162791
GO:BP	GO:000171: cell activation	0.006218	TRUE	"A multicellular process that involves the activation of a cell." (GO:000171)	1123	682	57	20972	0.083577713	0.050756901
GO:BP	GO:004271: circadian rhythm	0.006218	TRUE	"The cycle of a circadian rhythm." (GO:004271)	22	682	5	20972	0.007331378	0.227272727
GO:BP	GO:190021: positive regulation of gene expression	0.006218	TRUE	"Any process that is a positive regulation of gene expression." (GO:190021)	22	682	5	20972	0.007331378	0.227272727
GO:BP	GO:003221: dense core vesicle transport	0.006218	TRUE	"Any process that is a dense core vesicle transport." (GO:003221)	22	682	5	20972	0.007331378	0.227272727
GO:BP	GO:006001: reflex	0.006218	TRUE	"An automatic response to a stimulus." (GO:006001)	22	682	5	20972	0.007331378	0.227272727
GO:BP	GO:004211: regulation of gene expression	0.006236	TRUE	"Any process that is a regulation of gene expression." (GO:004211)	1639	294	39	20972	0.132653061	0.023794997
GO:BP	GO:000161: eye development	0.006295	TRUE	"The process of eye development." (GO:000161)	418	682	27	20972	0.039589443	0.064593301
GO:BP	GO:006031: regulation of gene expression	0.006296	TRUE	"Any process that is a regulation of gene expression." (GO:006031)	517	172	13	20972	0.075581395	0.025145068
GO:BP	GO:009881: nuclear chromosome organization	0.006315	TRUE	"The process of nuclear chromosome organization." (GO:009881)	324	227	11	20972	0.04845815	0.033950617
GO:BP	GO:002151: cerebellum development	0.006331	TRUE	"The process of cerebellum development." (GO:002151)	102	227	6	20972	0.026431718	0.058823529
GO:BP	GO:009711: intrinsic apoptosis	0.006386	TRUE	"The series of events that lead to the activation of the intrinsic pathway of apoptosis." (GO:009711)	318	1172	32	20972	0.027303754	0.100628931
GO:BP	GO:003571: platelet degranulation	0.006386	TRUE	"The series of events that lead to the activation of the platelet degranulation." (GO:003571)	56	1172	10	20972	0.008532423	0.178571429
GO:BP	GO:009011: epithelium morphogenesis	0.006386	TRUE	"The process of epithelium morphogenesis." (GO:009011)	97	1172	14	20972	0.011945392	0.144329897
GO:BP	GO:004821: vesicle docking	0.006397	TRUE	"The initial docking of a vesicle." (GO:004821)	59	682	8	20972	0.011730205	0.13559322
GO:BP	GO:004571: regulation of gene expression	0.006397	TRUE	"Any process that is a regulation of gene expression." (GO:004571)	292	682	21	20972	0.030791789	0.071917808

GO:BP	GO:00480: vascular er	0.006418	TRUE	"The serie	105	682	11	20972	0.016129032	0.104761905
GO:BP	GO:00072: ensheathm	0.00645	TRUE	"The proce	154	1172	19	20972	0.016211604	0.123376623
GO:BP	GO:00083: axon ensh	0.00645	TRUE	"Any proce	154	1172	19	20972	0.016211604	0.123376623
GO:BP	GO:00024: antigen pri	0.006474	TRUE	"The proce	13	682	4	20972	0.005865103	0.307692308
GO:BP	GO:00602: positive re	0.006474	TRUE	"Any proce	13	682	4	20972	0.005865103	0.307692308
GO:BP	GO:00482: norepinep	0.006474	TRUE	"The regul	13	682	4	20972	0.005865103	0.307692308
GO:BP	GO:19029: regulation	0.006489	TRUE	"Any proce	377	227	12	20972	0.052863436	0.031830239
GO:BP	GO:00400: regulation	0.006544	TRUE	"Any proce	66	1172	11	20972	0.009385666	0.166666667
GO:BP	GO:00063: DNA-temp	0.006565	TRUE	"The initial	212	294	10	20972	0.034013605	0.047169811
GO:BP	GO:00469: intracellul	0.006567	TRUE	"The direc	1375	172	24	20972	0.139534884	0.017454545
GO:BP	GO:00705: calcium ioi	0.006571	TRUE	"A process	326	172	10	20972	0.058139535	0.030674847
GO:BP	GO:00324: protein mc	0.006571	TRUE	"A protein	810	172	17	20972	0.098837209	0.020987654
GO:BP	GO:00602: regulation	0.006589	TRUE	"Any proce	15	1172	5	20972	0.004266212	0.333333333
GO:BP	GO:20010: positive re	0.006589	TRUE	"Any proce	15	1172	5	20972	0.004266212	0.333333333
GO:BP	GO:19049: interneuro	0.006589	TRUE	"The order	15	1172	5	20972	0.004266212	0.333333333
GO:BP	GO:20003: regulation	0.006589	TRUE	"Any proce	15	1172	5	20972	0.004266212	0.333333333
GO:BP	GO:00550: neuroblast	0.006589	TRUE	"The proce	15	1172	5	20972	0.004266212	0.333333333
GO:BP	GO:00072: G protein-i	0.006589	TRUE	"A G prote	15	1172	5	20972	0.004266212	0.333333333
GO:BP	GO:00705: calcium ioi	0.006604	TRUE	"A process	326	227	11	20972	0.04845815	0.033742331
GO:BP	GO:00469: secretion	0.00662	TRUE	"The contr	971	227	22	20972	0.0969163	0.022657055
GO:BP	GO:00351: forelimb r	0.006626	TRUE	"The proce	40	227	4	20972	0.017621145	0.1
GO:BP	GO:19029: regulation	0.006628	TRUE	"Any proce	377	682	25	20972	0.036656891	0.066312997
GO:BP	GO:00860: atrial cardi	0.006633	TRUE	"The proce	18	227	3	20972	0.013215859	0.166666667
GO:BP	GO:00860: atrial cardi	0.006633	TRUE	"Any proce	18	227	3	20972	0.013215859	0.166666667
GO:BP	GO:00860: atrial cardi	0.006633	TRUE	"An action	18	227	3	20972	0.013215859	0.166666667
GO:BP	GO:00216: cerebellar	0.006642	TRUE	"The proce	53	294	5	20972	0.017006803	0.094339623
GO:BP	GO:00074: brain deve	0.00666	TRUE	"The proce	736	172	16	20972	0.093023256	0.02173913
GO:BP	GO:00434: steroid ho	0.006671	TRUE	"The serie	140	682	13	20972	0.019061584	0.092857143
GO:BP	GO:00350: response t	0.006671	TRUE	"Any proce	46	682	7	20972	0.01026393	0.152173913
GO:BP	GO:00454: photorece	0.006671	TRUE	"Any proce	46	682	7	20972	0.01026393	0.152173913
GO:BP	GO:00305: nuclear rex	0.006671	TRUE	"A nuclear	140	682	13	20972	0.019061584	0.092857143
GO:BP	GO:00955: acetylcholi	0.006671	TRUE	"The serie	46	682	7	20972	0.01026393	0.152173913

GO:BP	GO:00434: regulation	0.006687	TRUE	"Any proce	176	294	9	20972	0.030612245	0.051136364
GO:BP	GO:00600: reflex	0.006717	TRUE	"An autom	22	1172	6	20972	0.005119454	0.272727273
GO:BP	GO:00427: circadian s	0.006717	TRUE	"The cycle	22	1172	6	20972	0.005119454	0.272727273
GO:BP	GO:00016: eye develc	0.006734	TRUE	"The proce	418	294	15	20972	0.051020408	0.035885167
GO:BP	GO:00713: cellular res	0.006736	TRUE	"Any proce	600	682	35	20972	0.051319648	0.058333333
GO:BP	GO:01405: chromatin	0.006747	TRUE	"A chroma	14	294	3	20972	0.010204082	0.214285714
GO:BP	GO:00650: protein-DN	0.006747	TRUE	"The aggre	213	294	10	20972	0.034013605	0.046948357
GO:BP	GO:00611: morphoge	0.006768	TRUE	"The proce	191	1172	22	20972	0.018771331	0.115183246
GO:BP	GO:00311: microtubu	0.006801	TRUE	"Assembly	143	1172	18	20972	0.015358362	0.125874126
GO:BP	GO:00455: negative re	0.006823	TRUE	"Any proce	665	172	15	20972	0.087209302	0.022556391
GO:BP	GO:00197: cellular ho	0.006823	TRUE	"Any proce	738	172	16	20972	0.093023256	0.021680217
GO:BP	GO:00430: regulation	0.006823	TRUE	"Any proce	1467	172	25	20972	0.145348837	0.017041581
GO:BP	GO:00197: calcium-m	0.00683	TRUE	"Any intrac	234	682	18	20972	0.026392962	0.076923077
GO:BP	GO:00508: protein sta	0.006844	TRUE	"Any proce	230	227	9	20972	0.039647577	0.039130435
GO:BP	GO:00016: response t	0.006855	TRUE	"Any proce	278	227	10	20972	0.044052863	0.035971223
GO:BP	GO:00224: positive re	0.006855	TRUE	"Any proce	328	227	11	20972	0.04845815	0.033536585
GO:BP	GO:20012: positive re	0.006857	TRUE	"Any proce	18	172	3	20972	0.01744186	0.166666667
GO:BP	GO:00602: regulation	0.006857	TRUE	"Any proce	816	172	17	20972	0.098837209	0.020833333
GO:BP	GO:00434: response t	0.006857	TRUE	"Any proce	457	172	12	20972	0.069767442	0.026258206
GO:BP	GO:00300: actin filam	0.006857	TRUE	"Any cellul	816	172	17	20972	0.098837209	0.020833333
GO:BP	GO:00488: cell motilit	0.006883	TRUE	"Any proce	1794	227	34	20972	0.149779736	0.018952062
GO:BP	GO:00075: skeletal m	0.006912	TRUE	"The devel	177	294	9	20972	0.030612245	0.050847458
GO:BP	GO:19013: regulation	0.006927	TRUE	"Any proce	60	682	8	20972	0.011730205	0.133333333
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GO:BP	GO:00971: neuronal s	0.008772	TRUE	"Any proce	24	682	5	20972	0.007331378	0.208333333
GO:BP	GO:00024: antigen pri	0.008772	TRUE	"The proce	24	682	5	20972	0.007331378	0.208333333
GO:BP	GO:19005: regulation	0.008793	TRUE	"Any proce	85	294	6	20972	0.020408163	0.070588235
GO:BP	GO:00902: positive re	0.008799	TRUE	"Any proce	110	682	11	20972	0.016129032	0.1
GO:BP	GO:00106: negative re	0.00881	TRUE	"Any proce	1678	294	39	20972	0.132653061	0.023241955
GO:BP	GO:00519: regulation	0.00881	TRUE	"Any proce	223	294	10	20972	0.034013605	0.044843049
GO:BP	GO:19013: negative re	0.008832	TRUE	"Any proce	112	1172	15	20972	0.012798635	0.133928571
GO:BP	GO:00027: positive re	0.008832	TRUE	"Any proce	112	1172	15	20972	0.012798635	0.133928571
GO:BP	GO:00096: response t	0.00885	TRUE	"Any proce	2427	682	106	20972	0.15542522	0.043675319
GO:BP	GO:00902: regulation	0.008863	TRUE	"Any proce	226	172	8	20972	0.046511628	0.03539823
GO:BP	GO:19028: microtubu	0.008874	TRUE	"Any micro	171	1172	20	20972	0.017064846	0.116959064
GO:BP	GO:00351: forelimb r	0.008915	TRUE	"The proce	40	1172	8	20972	0.006825939	0.2
GO:BP	GO:19031: regulation	0.008915	TRUE	"Any proce	40	1172	8	20972	0.006825939	0.2
GO:BP	GO:00350: cardiocyte	0.008917	TRUE	"The proce	147	1172	18	20972	0.015358362	0.12244898
GO:BP	GO:00605: respiratory	0.008917	TRUE	"The progr	221	1172	24	20972	0.020477816	0.108597285
GO:BP	GO:00510: positive re	0.008975	TRUE	"Any proce	818	294	23	20972	0.078231293	0.028117359
GO:BP	GO:00486: animal org	0.009012	TRUE	"The proce	69	1172	11	20972	0.009385666	0.15942029
GO:BP	GO:00480: regulation	0.009014	TRUE	"Any proce	111	227	6	20972	0.026431718	0.054054054
GO:BP	GO:01200: positive re	0.009014	TRUE	"Any proce	111	227	6	20972	0.026431718	0.054054054
GO:BP	GO:00488: cell motilit	0.009022	TRUE	"Any proce	1794	294	41	20972	0.139455782	0.022853958
GO:BP	GO:00485: developm	0.009081	TRUE	"The grow	221	682	17	20972	0.024926686	0.076923077
GO:BP	GO:00323: regulation	0.009095	TRUE	"Any proce	260	1172	27	20972	0.023037543	0.103846154
GO:BP	GO:01400: organelle l	0.009105	TRUE	"The proce	78	682	9	20972	0.013196481	0.115384615
GO:BP	GO:00424: photorece	0.009132	TRUE	"Developm	58	294	5	20972	0.017006803	0.086206897
GO:BP	GO:00715: response t	0.009174	TRUE	"Any proce	264	294	11	20972	0.037414966	0.041666667
GO:BP	GO:00485: embryonic	0.009174	TRUE	"Morphog	305	294	12	20972	0.040816327	0.039344262
GO:BP	GO:00488: regulation	0.009175	TRUE	"Any proce	63	682	8	20972	0.011730205	0.126984127
GO:BP	GO:00148: muscle hy	0.009211	TRUE	"The musc	86	294	6	20972	0.020408163	0.069767442
GO:BP	GO:00061: regulation	0.009211	TRUE	"Any proce	86	294	6	20972	0.020408163	0.069767442
GO:BP	GO:20007: regulation	0.00922	TRUE	"Any proce	16	294	3	20972	0.010204082	0.1875
GO:BP	GO:19000: regulation	0.00922	TRUE	"Any proce	16	294	3	20972	0.010204082	0.1875
GO:BP	GO:00454: locomotor	0.00922	TRUE	"The rhyth	16	294	3	20972	0.010204082	0.1875

GO:BP	GO:000821: positive re	0.00922	TRUE	"Any proce	923	294	25	20972	0.085034014	0.02708559
GO:BP	GO:014061: membran	0.009237	TRUE	"The aggre	436	294	15	20972	0.051020408	0.03440367
GO:BP	GO:003301: regulation	0.009237	TRUE	"Any proce	1167	172	21	20972	0.122093023	0.017994859
GO:BP	GO:190431: regulation	0.009252	TRUE	"Any proce	152	227	7	20972	0.030837004	0.046052632
GO:BP	GO:004591: negative re	0.009303	TRUE	"Any cellul	1510	172	25	20972	0.145348837	0.016556291
GO:BP	GO:004871: astrocyte c	0.009311	TRUE	"The proce	83	172	5	20972	0.029069767	0.060240964
GO:BP	GO:005501: potassium	0.009354	TRUE	"Any proce	36	682	6	20972	0.008797654	0.166666667
GO:BP	GO:004271: regulation	0.009394	TRUE	"Any proce	112	227	6	20972	0.026431718	0.053571429
GO:BP	GO:002201: metencepl	0.009394	TRUE	"The proce	112	227	6	20972	0.026431718	0.053571429
GO:BP	GO:004401: regulation	0.009399	TRUE	"Any proce	1006	172	19	20972	0.110465116	0.01888668
GO:BP	GO:004571: positive re	0.009399	TRUE	"Any proce	229	172	8	20972	0.046511628	0.034934498
GO:BP	GO:005171: regulation	0.009399	TRUE	"Any proce	1088	172	20	20972	0.11627907	0.018382353
GO:BP	GO:004291: regulation	0.009399	TRUE	"Any proce	1426	172	24	20972	0.139534884	0.016830295
GO:BP	GO:005111: striated m	0.009401	TRUE	"The proce	287	172	9	20972	0.052325581	0.031358885
GO:BP	GO:008011: regulation	0.009409	TRUE	"Any proce	1410	227	28	20972	0.123348018	0.019858156
GO:BP	GO:000991: negative re	0.00943	TRUE	"Any proce	1341	172	23	20972	0.13372093	0.01715138
GO:BP	GO:001061: positive re	0.009431	TRUE	"Any proce	499	682	30	20972	0.04398827	0.06012024
GO:BP	GO:000711: negative re	0.009459	TRUE	"Any proce	287	1172	29	20972	0.024744027	0.101045296
GO:BP	GO:004571: positive re	0.009468	TRUE	"Any proce	349	294	13	20972	0.044217687	0.037249284
GO:BP	GO:000221: behavioral	0.009491	TRUE	"A behavic	45	227	4	20972	0.017621145	0.088888889
GO:BP	GO:003511: appendage	0.009509	TRUE	"The proce	153	227	7	20972	0.030837004	0.045751634
GO:BP	GO:003511: limb morp	0.009509	TRUE	"The proce	153	227	7	20972	0.030837004	0.045751634
GO:BP	GO:005121: negative re	0.009509	TRUE	"Any proce	569	227	15	20972	0.066079295	0.026362039
GO:BP	GO:004851: negative re	0.009513	TRUE	"Any proce	1693	172	27	20972	0.156976744	0.015948021
GO:BP	GO:000861: intrinsic a	0.009537	TRUE	"The serie	10	1172	4	20972	0.003412969	0.4
GO:BP	GO:009891: retrograde	0.009537	TRUE	"Cell-cell s	10	1172	4	20972	0.003412969	0.4
GO:BP	GO:004651: retinal rod	0.009537	TRUE	"Developm	10	1172	4	20972	0.003412969	0.4
GO:BP	GO:005191: amino acic	0.009537	TRUE	"The uptak	10	1172	4	20972	0.003412969	0.4
GO:BP	GO:004871: branching	0.009537	TRUE	"The proce	10	1172	4	20972	0.003412969	0.4
GO:BP	GO:000631: nucleosom	0.009539	TRUE	"The aggre	118	294	7	20972	0.023809524	0.059322034
GO:BP	GO:000691: response t	0.009556	TRUE	"Any proce	80	1172	12	20972	0.010238908	0.15
GO:BP	GO:000911: nucleoside	0.009556	TRUE	"The chem	80	1172	12	20972	0.010238908	0.15

GO:BP	GO:004241: mechanoreception	0.009556	TRUE	"The process of detecting mechanical stimuli"	80	1172	12	20972	0.010238908	0.15
GO:BP	GO:001481: muscle cell contraction	0.009556	TRUE	"The order of muscle contraction"	80	1172	12	20972	0.010238908	0.15
GO:BP	GO:007081: positive regulation of gene expression	0.009573	TRUE	"Any process that positively regulates gene expression"	21	172	3	20972	0.01744186	0.142857143
GO:BP	GO:010601: positive regulation of gene expression	0.009573	TRUE	"Any process that positively regulates gene expression"	21	172	3	20972	0.01744186	0.142857143
GO:BP	GO:004661: response to stress	0.009613	TRUE	"Any process that is a response to stress"	35	294	4	20972	0.013605442	0.114285714
GO:BP	GO:003461: cellular response to stress	0.009613	TRUE	"Any process that is a cellular response to stress"	87	294	6	20972	0.020408163	0.068965517
GO:BP	GO:001481: release of neurotransmitter	0.009613	TRUE	"The process of releasing neurotransmitters"	35	294	4	20972	0.013605442	0.114285714
GO:BP	GO:003311: regulation of gene expression	0.009613	TRUE	"Any process that regulates gene expression"	35	294	4	20972	0.013605442	0.114285714
GO:BP	GO:005111: negative regulation of gene expression	0.009622	TRUE	"Any process that negatively regulates gene expression"	59	294	5	20972	0.017006803	0.084745763
GO:BP	GO:003821: TORC2 signaling	0.009647	TRUE	"A series of events in TORC2 signaling"	21	227	3	20972	0.013215859	0.142857143
GO:BP	GO:008601: membrane transport	0.009647	TRUE	"The process of transporting substances across a membrane"	21	227	3	20972	0.013215859	0.142857143
GO:BP	GO:000681: sodium ion transport	0.009712	TRUE	"The direct transport of sodium ions"	245	227	9	20972	0.039647577	0.036734694
GO:BP	GO:002191: cerebral cortex development	0.00973	TRUE	"The program of development of the cerebral cortex"	128	172	6	20972	0.034883721	0.046875
GO:BP	GO:004501: regulated gene expression	0.009742	TRUE	"A process of regulated gene expression"	227	294	10	20972	0.034013605	0.044052863
GO:BP	GO:004341: regulation of gene expression	0.009777	TRUE	"Any process that regulates gene expression"	189	294	9	20972	0.030612245	0.047619048
GO:BP	GO:190421: negative regulation of gene expression	0.009805	TRUE	"Any process that negatively regulates gene expression"	50	1172	9	20972	0.007679181	0.18
GO:BP	GO:003241: positive regulation of gene expression	0.009805	TRUE	"Any process that positively regulates gene expression"	50	1172	9	20972	0.007679181	0.18
GO:BP	GO:003631: vascular endothelial cell growth	0.009805	TRUE	"The series of events in vascular endothelial cell growth"	50	1172	9	20972	0.007679181	0.18
GO:BP	GO:001091: negative regulation of gene expression	0.009845	TRUE	"Any process that negatively regulates gene expression"	125	1172	16	20972	0.013651877	0.128
GO:BP	GO:005161: establishment of gene expression	0.009904	TRUE	"The direct establishment of gene expression"	32	1172	7	20972	0.005972696	0.21875
GO:BP	GO:000201: regulation of gene expression	0.009924	TRUE	"Any process that regulates gene expression"	70	1172	11	20972	0.009385666	0.157142857
GO:BP	GO:000161: osteoblast differentiation	0.009924	TRUE	"The process of osteoblast differentiation"	236	1172	25	20972	0.021331058	0.105932203
GO:BP	GO:003201: positive regulation of gene expression	0.009939	TRUE	"Any process that positively regulates gene expression"	77	227	5	20972	0.022026432	0.064935065
GO:BP	GO:007121: cellular response to stress	0.009939	TRUE	"Any process that is a cellular response to stress"	77	227	5	20972	0.022026432	0.064935065
GO:BP	GO:001631: phosphorylation	0.009987	TRUE	"The process of phosphorylation"	1077	227	23	20972	0.101321586	0.021355617
GO:BP	GO:000271: positive regulation of gene expression	0.010073	TRUE	"Any process that positively regulates gene expression"	112	682	11	20972	0.016129032	0.098214286
GO:BP	GO:000681: exocytosis	0.010077	TRUE	"A process of exocytosis"	352	294	13	20972	0.044217687	0.036931818
GO:BP	GO:001491: myotube development	0.010097	TRUE	"The process of myotube development"	129	172	6	20972	0.034883721	0.046511628
GO:BP	GO:005501: muscle cell contraction	0.010109	TRUE	"The process of muscle cell contraction"	190	294	9	20972	0.030612245	0.047368421
GO:BP	GO:004851: embryonic development	0.010153	TRUE	"Morphogenesis of an embryo"	305	682	21	20972	0.030791789	0.068852459
GO:BP	GO:009711: neuronal signaling	0.010201	TRUE	"Any process that involves neuronal signaling"	24	1172	6	20972	0.005119454	0.25
GO:BP	GO:009961: regulation of gene expression	0.010201	TRUE	"Any process that regulates gene expression"	24	1172	6	20972	0.005119454	0.25

GO:BP	GO:00714: protein loc	0.010242	TRUE	"Any proce	41	1172	8	20972	0.006825939	0.195121951
GO:BP	GO:01407: constitutiv	0.01027	TRUE	"The comp	60	294	5	20972	0.017006803	0.083333333
GO:BP	GO:00072: glutamate	0.01027	TRUE	"The serie	60	294	5	20972	0.017006803	0.083333333
GO:BP	GO:00325: regulation	0.01027	TRUE	"A process	353	294	13	20972	0.044217687	0.036827195
GO:BP	GO:00085: visual learn	0.010378	TRUE	"Any proce	50	682	7	20972	0.01026393	0.14
GO:BP	GO:00481: regulation	0.010391	TRUE	"A process	25	682	5	20972	0.007331378	0.2
GO:BP	GO:19034: regulation	0.010391	TRUE	"Any proce	25	682	5	20972	0.007331378	0.2
GO:BP	GO:00322: telomere c	0.010402	TRUE	"A process	191	294	9	20972	0.030612245	0.047120419
GO:BP	GO:00605: skeletal m	0.010402	TRUE	"The progr	191	294	9	20972	0.030612245	0.047120419
GO:BP	GO:00431: proteasom	0.010445	TRUE	"The chem	468	1172	42	20972	0.035836177	0.08974359
GO:BP	GO:00434: phosphatic	0.01046	TRUE	"An intrac	306	682	21	20972	0.030791789	0.068627451
GO:BP	GO:00316: cellular res	0.010498	TRUE	"Any proce	250	1172	26	20972	0.0221843	0.104
GO:BP	GO:00432: anoikis	0.010509	TRUE	"Apoptosis	36	294	4	20972	0.013605442	0.111111111
GO:BP	GO:00901: tissue mig	0.010523	TRUE	"The proce	103	1172	14	20972	0.011945392	0.13592233
GO:BP	GO:00347: negative re	0.010523	TRUE	"Any proce	103	1172	14	20972	0.011945392	0.13592233
GO:BP	GO:00024: antigen pri	0.010546	TRUE	"The proce	15	682	4	20972	0.005865103	0.266666667
GO:BP	GO:00019: positive re	0.010546	TRUE	"Any proce	37	682	6	20972	0.008797654	0.162162162
GO:BP	GO:00986: postsynap	0.010546	TRUE	"The aggre	37	682	6	20972	0.008797654	0.162162162
GO:BP	GO:00995: modificati	0.010546	TRUE	"Any proce	37	682	6	20972	0.008797654	0.162162162
GO:BP	GO:00026: regulation	0.010546	TRUE	"Any proce	572	682	33	20972	0.048387097	0.057692308
GO:BP	GO:00602: regulation	0.010546	TRUE	"Any proce	15	682	4	20972	0.005865103	0.266666667
GO:BP	GO:20010: positive re	0.010546	TRUE	"Any proce	15	682	4	20972	0.005865103	0.266666667
GO:BP	GO:00702: actin-medi	0.010553	TRUE	"The actin	115	227	6	20972	0.026431718	0.052173913
GO:BP	GO:00424: mechanori	0.010556	TRUE	"The proce	80	682	9	20972	0.013196481	0.1125
GO:BP	GO:00158: acidic amir	0.010556	TRUE	"The direc	80	682	9	20972	0.013196481	0.1125
GO:BP	GO:00315: cell-substr	0.010563	TRUE	"The attac	349	682	23	20972	0.03372434	0.065902579
GO:BP	GO:00347: positive re	0.01061	TRUE	"Any proce	126	1172	16	20972	0.013651877	0.126984127
GO:BP	GO:00435: regulation	0.01061	TRUE	"Any proce	92	1172	13	20972	0.01109215	0.141304348
GO:BP	GO:01200: regulation	0.01063	TRUE	"Any proce	201	227	8	20972	0.035242291	0.039800995
GO:BP	GO:00070: regulation	0.010694	TRUE	"Any proce	17	294	3	20972	0.010204082	0.176470588
GO:BP	GO:00020: chondrocy	0.010694	TRUE	"The proce	121	294	7	20972	0.023809524	0.05785124
GO:BP	GO:00076: memory	0.010694	TRUE	"The activi	121	294	7	20972	0.023809524	0.05785124

GO:BP	GO:00970!	synaptic ve	0.010694	TRUE	"The proce	17	294	3	20972	0.010204082	0.176470588
GO:BP	GO:00610!	regulation	0.010694	TRUE	"Any proce	192	294	9	20972	0.030612245	0.046875
GO:BP	GO:00068!	calcium io	0.010711	TRUE	"The direc	421	172	11	20972	0.063953488	0.026128266
GO:BP	GO:00100!	glial cell di	0.010752	TRUE	"The proce	249	227	9	20972	0.039647577	0.036144578
GO:BP	GO:00311!	cytoplasm	0.010894	TRUE	"A process	65	682	8	20972	0.011730205	0.123076923
GO:BP	GO:00488!	genitalia d	0.010944	TRUE	"The proce	47	227	4	20972	0.017621145	0.085106383
GO:BP	GO:00430!	positive re	0.010985	TRUE	"Any proce	512	1172	45	20972	0.038395904	0.087890625
GO:BP	GO:00715!	response t	0.010996	TRUE	"Any proce	264	1172	27	20972	0.023037543	0.102272727
GO:BP	GO:00070!	cytoskelet	0.010998	TRUE	"A process	1534	172	25	20972	0.145348837	0.016297262
GO:BP	GO:00328!	regulation	0.011031	TRUE	"Any proce	71	1172	11	20972	0.009385666	0.154929577
GO:BP	GO:00611!	mammary	0.011031	TRUE	"The proce	71	1172	11	20972	0.009385666	0.154929577
GO:BP	GO:19021!	regulation	0.011031	TRUE	"Any proce	238	1172	25	20972	0.021331058	0.105042017
GO:BP	GO:00456!	regulation	0.011031	TRUE	"Any proce	71	1172	11	20972	0.009385666	0.154929577
GO:BP	GO:00325!	regulation	0.011065	TRUE	"A process	353	227	11	20972	0.04845815	0.031161473
GO:BP	GO:00466!	lymphocyt	0.011075	TRUE	"A change	811	682	43	20972	0.063049853	0.053020962
GO:BP	GO:00604!	regulation	0.011086	TRUE	"Any proce	203	227	8	20972	0.035242291	0.039408867
GO:BP	GO:00071!	negative re	0.0111	TRUE	"Any proce	287	682	20	20972	0.029325513	0.069686411
GO:BP	GO:00456!	positive re	0.01112	TRUE	"Any proce	90	294	6	20972	0.020408163	0.066666667
GO:BP	GO:00514!	regulation	0.01112	TRUE	"Any proce	90	294	6	20972	0.020408163	0.066666667
GO:BP	GO:00104!	proteasom	0.011156	TRUE	"The chem	599	1172	51	20972	0.043515358	0.085141903
GO:BP	GO:00711!	protein loc	0.011156	TRUE	"Any proce	51	1172	9	20972	0.007679181	0.176470588
GO:BP	GO:00614!	connective	0.011244	TRUE	"The progr	291	1172	29	20972	0.024744027	0.099656357
GO:BP	GO:00069!	apoptotic	0.011268	TRUE	"A program	1898	172	29	20972	0.168604651	0.015279241
GO:BP	GO:00603!	head deve	0.011268	TRUE	"The biolo	787	172	16	20972	0.093023256	0.020330368
GO:BP	GO:00070!	regulation	0.01127	TRUE	"Any proce	17	1172	5	20972	0.004266212	0.294117647
GO:BP	GO:20006!	regulation	0.01127	TRUE	"Any proce	17	1172	5	20972	0.004266212	0.294117647
GO:BP	GO:00511!	meiotic sis	0.01127	TRUE	"The cell c	17	1172	5	20972	0.004266212	0.294117647
GO:BP	GO:00108!	negative re	0.01127	TRUE	"Any proce	17	1172	5	20972	0.004266212	0.294117647
GO:BP	GO:00512!	regulation	0.011274	TRUE	"Any proce	506	682	30	20972	0.04398827	0.059288538
GO:BP	GO:00316!	regulation	0.011274	TRUE	"Any proce	114	682	11	20972	0.016129032	0.096491228
GO:BP	GO:00480!	hepatocyt	0.0113	TRUE	"The serie	61	1172	10	20972	0.008532423	0.163934426
GO:BP	GO:19051!	response t	0.011394	TRUE	"Any proce	51	682	7	20972	0.01026393	0.137254902

GO:BP	GO:00986! postsynap	0.011395	TRUE	"The aggre	37	294	4	20972	0.013605442	0.108108108
GO:BP	GO:00456! regulation	0.011395	TRUE	"Any proce	37	294	4	20972	0.013605442	0.108108108
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GO:BP	GO:00508: positive re	0.014023	TRUE	"Any proce	380	682	24	20972	0.035190616	0.063157895
GO:BP	GO:00108: regulation	0.014023	TRUE	"Any proce	212	682	16	20972	0.023460411	0.075471698
GO:BP	GO:00015: skeletal sy	0.014023	TRUE	"The proce	538	682	31	20972	0.045454545	0.057620818
GO:BP	GO:00451: developm	0.014023	TRUE	"The proce	232	682	17	20972	0.024926686	0.073275862
GO:BP	GO:00905: sensory or	0.014023	TRUE	"Morphog	294	682	20	20972	0.029325513	0.068027211
GO:BP	GO:01400: mitotic nu	0.014054	TRUE	"A mitotic	283	1172	28	20972	0.023890785	0.098939929
GO:BP	GO:00987: L-glutamat	0.014132	TRUE	"The direc	18	1172	5	20972	0.004266212	0.277777778
GO:BP	GO:00381: angiotensi	0.014132	TRUE	"A G prote	18	1172	5	20972	0.004266212	0.277777778
GO:BP	GO:00148: regulation	0.014132	TRUE	"Any proce	18	1172	5	20972	0.004266212	0.277777778
GO:BP	GO:00158: gamma-an	0.014132	TRUE	"The direc	18	1172	5	20972	0.004266212	0.277777778
GO:BP	GO:00430: negative re	0.014132	TRUE	"Any proce	18	1172	5	20972	0.004266212	0.277777778
GO:BP	GO:00215: cell migrat	0.014132	TRUE	"The order	18	1172	5	20972	0.004266212	0.277777778
GO:BP	GO:00198: antigen pri	0.014144	TRUE	"The proce	118	682	11	20972	0.016129032	0.093220339
GO:BP	GO:00310: actomyosi	0.014162	TRUE	"A process	202	294	9	20972	0.030612245	0.044554455
GO:BP	GO:00105: positive re	0.014191	TRUE	"Any proce	107	1172	14	20972	0.011945392	0.130841121
GO:BP	GO:00007: recombina	0.014274	TRUE	"A DNA re	192	1172	21	20972	0.017918089	0.109375
GO:BP	GO:00513: cell divisio	0.014335	TRUE	"The proce	656	682	36	20972	0.052785924	0.054878049
GO:BP	GO:00480: insulin-like	0.014411	TRUE	"The serie	101	682	10	20972	0.014662757	0.099009901
GO:BP	GO:01201: positive re	0.014411	TRUE	"Any proce	101	682	10	20972	0.014662757	0.099009901
GO:BP	GO:19035: regulation	0.014415	TRUE	"Any proce	556	294	17	20972	0.057823129	0.03057554

GO:BP	GO:005071	negative re	0.014415	TRUE	"Any proce	129	294	7	20972	0.023809524	0.054263566
GO:BP	GO:007151	cellular res	0.014419	TRUE	"Any proce	257	1172	26	20972	0.0221843	0.101167315
GO:BP	GO:001061	regulation	0.014432	TRUE	"Any proce	55	172	4	20972	0.023255814	0.072727273
GO:BP	GO:003811	neurotropi	0.014441	TRUE	"The serie	40	294	4	20972	0.013605442	0.1
GO:BP	GO:000631	transcripti	0.014441	TRUE	"The synth	66	294	5	20972	0.017006803	0.075757576
GO:BP	GO:003511	forelimb m	0.014441	TRUE	"The proce	40	294	4	20972	0.013605442	0.1
GO:BP	GO:001601	synaptic ve	0.014441	TRUE	"A process	25	172	3	20972	0.01744186	0.12
GO:BP	GO:001901	viral trans	0.014441	TRUE	"The proce	25	172	3	20972	0.01744186	0.12
GO:BP	GO:003591	response t	0.014441	TRUE	"Any proce	25	172	3	20972	0.01744186	0.12
GO:BP	GO:007081	inclusion b	0.014441	TRUE	"The aggre	25	172	3	20972	0.01744186	0.12
GO:BP	GO:199081	adaptive tl	0.014477	TRUE	"The regul	167	227	7	20972	0.030837004	0.041916168
GO:BP	GO:000821	insulin rec	0.014499	TRUE	"The serie	174	682	14	20972	0.020527859	0.08045977
GO:BP	GO:190251	negative re	0.014522	TRUE	"Any proce	681	682	37	20972	0.054252199	0.054331865
GO:BP	GO:000171	cell activat	0.014571	TRUE	"A multice	1123	294	28	20972	0.095238095	0.024933215
GO:BP	GO:005101	negative re	0.014602	TRUE	"Any proce	404	682	25	20972	0.036656891	0.061881188
GO:BP	GO:000961	response t	0.014612	TRUE	"Any proce	214	227	8	20972	0.035242291	0.037383178
GO:BP	GO:190201	regulation	0.014636	TRUE	"Any proce	96	1172	13	20972	0.01109215	0.135416667
GO:BP	GO:000311	outflow tr	0.014732	TRUE	"The proce	26	1172	6	20972	0.005119454	0.230769231
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GO:BP	GO:004551	positive re	0.014807	TRUE	"Any proce	119	682	11	20972	0.016129032	0.092436975
GO:BP	GO:005101	mRNA trar	0.014809	TRUE	"The direc	131	1172	16	20972	0.013651877	0.122137405
GO:BP	GO:003161	protein de	0.014815	TRUE	"Any proce	52	227	4	20972	0.017621145	0.076923077
GO:BP	GO:003021	platelet fo	0.014815	TRUE	"The proce	25	227	3	20972	0.013215859	0.12
GO:BP	GO:004651	negative re	0.014815	TRUE	"Any proce	25	227	3	20972	0.013215859	0.12
GO:BP	GO:003031	embryonic	0.014829	TRUE	"The proce	125	227	6	20972	0.026431718	0.048
GO:BP	GO:003511	embryonic	0.014829	TRUE	"The proce	125	227	6	20972	0.026431718	0.048
GO:BP	GO:000721	JNK cascac	0.014838	TRUE	"A MAPK c	194	172	7	20972	0.040697674	0.036082474
GO:BP	GO:000701	cell cycle	0.014853	TRUE	"The progr	1668	172	26	20972	0.151162791	0.01558753

GO:BP	GO:005131 cell division	0.014858	TRUE	"The process of cell division"	656	294	19	20972	0.06462585	0.028963415
GO:BP	GO:007131 cellular response	0.014858	TRUE	"Any process that is a response to an external stimulus"	706	294	20	20972	0.068027211	0.028328612
GO:BP	GO:190291 regulation of cell cycle	0.014915	TRUE	"Any process that is a response to an external stimulus"	377	172	10	20972	0.058139535	0.026525199
GO:BP	GO:007061 leukocyte chemotaxis	0.014936	TRUE	"The expansion of a cell population"	339	682	22	20972	0.032258065	0.064896755
GO:BP	GO:004351 negative regulation of cell cycle	0.014967	TRUE	"Any process that is a response to an external stimulus"	166	294	8	20972	0.027210884	0.048192771
GO:BP	GO:000021 polysaccharide catabolic process	0.015012	TRUE	"The chemical breakdown of a substance"	56	172	4	20972	0.023255814	0.071428571
GO:BP	GO:006131 regulation of cell cycle	0.015096	TRUE	"Any process that is a response to an external stimulus"	97	294	6	20972	0.020408163	0.06185567
GO:BP	GO:001051 negative regulation of cell cycle	0.015096	TRUE	"Any process that is a response to an external stimulus"	97	294	6	20972	0.020408163	0.06185567
GO:BP	GO:003351 carbohydrate catabolic process	0.015122	TRUE	"A homeostatic process that maintains the internal environment of an organism"	258	1172	26	20972	0.0221843	0.100775194
GO:BP	GO:004351 regulation of cell cycle	0.015247	TRUE	"Any process that is a response to an external stimulus"	353	1172	33	20972	0.028156997	0.093484419
GO:BP	GO:005081 positive regulation of cell cycle	0.015273	TRUE	"Any process that is a response to an external stimulus"	255	682	18	20972	0.026392962	0.070588235
GO:BP	GO:003291 circadian rhythm	0.01528	TRUE	"Any process that is a response to an external stimulus"	67	294	5	20972	0.017006803	0.074626866
GO:BP	GO:001921 regulation of cell cycle	0.015347	TRUE	"Any process that is a response to an external stimulus"	64	1172	10	20972	0.008532423	0.15625
GO:BP	GO:003591 aorta development	0.015347	TRUE	"The progression of a cell through the cell cycle"	64	1172	10	20972	0.008532423	0.15625
GO:BP	GO:004591 positive regulation of cell cycle	0.015397	TRUE	"Any process that is a response to an external stimulus"	395	1172	36	20972	0.030716724	0.091139241
GO:BP	GO:001051 positive regulation of cell cycle	0.015397	TRUE	"Any process that is a response to an external stimulus"	395	1172	36	20972	0.030716724	0.091139241
GO:BP	GO:007031 regulation of cell cycle	0.015398	TRUE	"Any process that is a response to an external stimulus"	276	682	19	20972	0.027859238	0.06884058
GO:BP	GO:005101 regulation of cell cycle	0.015399	TRUE	"Any process that is a response to an external stimulus"	485	227	13	20972	0.057268722	0.026804124
GO:BP	GO:000021 protein catabolic process	0.015493	TRUE	"Addition of a substance to a mixture"	266	227	9	20972	0.039647577	0.033834586
GO:BP	GO:007121 cellular response	0.015581	TRUE	"Any process that is a response to an external stimulus"	87	227	5	20972	0.022026432	0.057471264
GO:BP	GO:005071 regulation of cell cycle	0.015671	TRUE	"Any process that is a response to an external stimulus"	138	682	12	20972	0.017595308	0.086956522
GO:BP	GO:002161 cerebellar development	0.015723	TRUE	"The process of a cell dividing into two daughter cells"	53	227	4	20972	0.017621145	0.075471698
GO:BP	GO:190271 positive regulation of cell cycle	0.015728	TRUE	"Any process that is a response to an external stimulus"	35	1172	7	20972	0.005972696	0.2
GO:BP	GO:000761 adult locomotion	0.015728	TRUE	"The behavior of an organism in response to an external stimulus"	35	1172	7	20972	0.005972696	0.2
GO:BP	GO:004861 neuron fate determination	0.015728	TRUE	"The process of a cell dividing into two daughter cells"	35	1172	7	20972	0.005972696	0.2
GO:BP	GO:003451 protein localization	0.01578	TRUE	"A process that involves the movement of a substance from one location to another"	319	227	10	20972	0.044052863	0.031347962
GO:BP	GO:007091 organelle organization	0.015792	TRUE	"The aggregation of a substance"	1057	227	22	20972	0.0969163	0.020813623
GO:BP	GO:007131 cellular response	0.015813	TRUE	"Any process that is a response to an external stimulus"	144	1172	17	20972	0.014505119	0.118055556
GO:BP	GO:000601 glycolytic process	0.015829	TRUE	"The chemical breakdown of a substance"	98	294	6	20972	0.020408163	0.06122449
GO:BP	GO:190301 regulation of cell cycle	0.015829	TRUE	"Any process that is a response to an external stimulus"	375	294	13	20972	0.044217687	0.034666667
GO:BP	GO:009001 positive regulation of cell cycle	0.015832	TRUE	"Any process that is a response to an external stimulus"	259	1172	26	20972	0.0221843	0.1003861
GO:BP	GO:000701 metaphase	0.015846	TRUE	"The cell cycle phase in which the chromosomes are aligned at the metaphase plate"	97	1172	13	20972	0.01109215	0.134020619

GO:BP	GO:00611! mRNA des	0.015846	TRUE	"Any proce	97	1172	13	20972	0.01109215	0.134020619
GO:BP	GO:00326! type I inter	0.015847	TRUE	"The appe	127	227	6	20972	0.026431718	0.047244094
GO:BP	GO:00324! regulation	0.015847	TRUE	"Any proce	127	227	6	20972	0.026431718	0.047244094
GO:BP	GO:19029! mitotic DN	0.01586	TRUE	"Any nucle	17	682	4	20972	0.005865103	0.235294118
GO:BP	GO:00300! intracellul	0.01586	TRUE	"A homeos	17	682	4	20972	0.005865103	0.235294118
GO:BP	GO:00019! positive re	0.015874	TRUE	"Any proce	20	294	3	20972	0.010204082	0.15
GO:BP	GO:00083! asymmetri	0.015874	TRUE	"The asym	20	294	3	20972	0.010204082	0.15
GO:BP	GO:00715! genomic ir	0.015874	TRUE	"The estab	20	294	3	20972	0.010204082	0.15
GO:BP	GO:00971! GABAergic	0.015874	TRUE	"The proce	20	294	3	20972	0.010204082	0.15
GO:BP	GO:00514! negative re	0.01589	TRUE	"Any proce	28	682	5	20972	0.007331378	0.178571429
GO:BP	GO:00312! positive re	0.01589	TRUE	"Any proce	28	682	5	20972	0.007331378	0.178571429
GO:BP	GO:00459! positive re	0.01589	TRUE	"Any proce	28	682	5	20972	0.007331378	0.178571429
GO:BP	GO:00313! regulation	0.015923	TRUE	"Any proce	194	1172	21	20972	0.017918089	0.108247423
GO:BP	GO:00070! cell-cell j	0.015924	TRUE	"The aggre	157	682	13	20972	0.019061584	0.082802548
GO:BP	GO:00860! cell comm	0.01598	TRUE	"Any proce	57	172	4	20972	0.023255814	0.070175439
GO:BP	GO:00429! amyloid pr	0.015984	TRUE	"The chem	75	1172	11	20972	0.009385666	0.146666667
GO:BP	GO:00100! response t	0.016023	TRUE	"Any proce	320	227	10	20972	0.044052863	0.03125
GO:BP	GO:00518! regulation	0.016077	TRUE	"Any proce	268	227	9	20972	0.039647577	0.03358209
GO:BP	GO:00481! regulation	0.016143	TRUE	"Any proce	88	227	5	20972	0.022026432	0.056818182
GO:BP	GO:00315! protein-co	0.016143	TRUE	"A localiza	88	227	5	20972	0.022026432	0.056818182
GO:BP	GO:00313! N-terminal	0.016167	TRUE	"The alter	26	227	3	20972	0.013215859	0.115384615
GO:BP	GO:00600! uterus dev	0.016167	TRUE	"The reproc	26	227	3	20972	0.013215859	0.115384615
GO:BP	GO:00216! cerebellar	0.016167	TRUE	"The proce	26	227	3	20972	0.013215859	0.115384615
GO:BP	GO:00215! dentate gy	0.016167	TRUE	"The proce	26	227	3	20972	0.013215859	0.115384615
GO:BP	GO:00432! regulation	0.016276	TRUE	"Any proce	422	294	14	20972	0.047619048	0.033175355
GO:BP	GO:00610! membrane	0.01633	TRUE	"A process	783	682	41	20972	0.060117302	0.052362708
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GO:BP	GO:00621! cellular res	0.016369	TRUE	"Any proce	300	1172	29	20972	0.024744027	0.096666667
GO:BP	GO:00974! supramole	0.016389	TRUE	"A process	798	227	18	20972	0.079295154	0.022556391
GO:BP	GO:00512! positive re	0.016427	TRUE	"Any proce	863	227	19	20972	0.083700441	0.022016222
GO:BP	GO:00508! regulation	0.016493	TRUE	"Any proce	386	682	24	20972	0.035190616	0.062176166
GO:BP	GO:00600! synapse m	0.016493	TRUE	"The proce	41	682	6	20972	0.008797654	0.146341463

GO:BP	GO:00107: negative re	0.016614	TRUE	"Any proce	260	1172	26	20972	0.0221843	0.1
GO:BP	GO:00706: protein me	0.016656	TRUE	"A protein	146	172	6	20972	0.034883721	0.04109589
GO:BP	GO:00481: regulation	0.016656	TRUE	"A process	58	172	4	20972	0.023255814	0.068965517
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GO:BP	GO:00313: regulation	0.016656	TRUE	"Any proce	748	172	15	20972	0.087209302	0.020053476
GO:BP	GO:00504: amyloid-be	0.016656	TRUE	"The chem	58	172	4	20972	0.023255814	0.068965517
GO:BP	GO:00362: response t	0.016678	TRUE	"Any proce	290	294	11	20972	0.037414966	0.037931034
GO:BP	GO:00159: long-chain	0.016747	TRUE	"The direc	65	1172	10	20972	0.008532423	0.153846154
GO:BP	GO:00182: peptidyl-ly	0.016747	TRUE	"The modi	133	1172	16	20972	0.013651877	0.120300752
GO:BP	GO:00106: regulation	0.016747	TRUE	"Any proce	65	1172	10	20972	0.008532423	0.153846154
GO:BP	GO:00311: regulation	0.016747	TRUE	"Any proce	65	1172	10	20972	0.008532423	0.153846154
GO:BP	GO:00001: MAPK casc	0.016754	TRUE	"An intrac	749	172	15	20972	0.087209302	0.020026702
GO:BP	GO:00218: forebrain g	0.016788	TRUE	"The proce	89	227	5	20972	0.022026432	0.056179775
GO:BP	GO:00712: cellular res	0.016816	TRUE	"Any proce	173	227	7	20972	0.030837004	0.040462428
GO:BP	GO:00507: negative re	0.016816	TRUE	"Any proce	129	227	6	20972	0.026431718	0.046511628
GO:BP	GO:00512: protein tet	0.016827	TRUE	"The formi	99	172	5	20972	0.029069767	0.050505051
GO:BP	GO:19018: regulation	0.016827	TRUE	"Any proce	258	172	8	20972	0.046511628	0.031007752
GO:BP	GO:00017: cell fate de	0.01687	TRUE	"The cellul	45	1172	8	20972	0.006825939	0.177777778
GO:BP	GO:00605: neuroepith	0.01687	TRUE	"The proce	45	1172	8	20972	0.006825939	0.177777778
GO:BP	GO:00342: amyloid-be	0.01687	TRUE	"The genei	45	1172	8	20972	0.006825939	0.177777778
GO:BP	GO:00484: autonomic	0.01687	TRUE	"The proce	45	1172	8	20972	0.006825939	0.177777778
GO:BP	GO:00323: icosanoid s	0.01687	TRUE	"The contr	45	1172	8	20972	0.006825939	0.177777778
GO:BP	GO:00171: regulation	0.01688	TRUE	"Any proce	42	294	4	20972	0.013605442	0.095238095
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GO:BP	GO:00015: ossificatio	0.01689	TRUE	"The formi	424	294	14	20972	0.047619048	0.033018868
GO:BP	GO:00602: anatomica	0.016996	TRUE	"A homeos	271	227	9	20972	0.039647577	0.033210332
GO:BP	GO:00018: tissue hor	0.016996	TRUE	"A homeos	271	227	9	20972	0.039647577	0.033210332
GO:BP	GO:00703: ERK1 and f	0.017049	TRUE	"A MAPK c	300	682	20	20972	0.029325513	0.066666667
GO:BP	GO:00713: cellular res	0.017056	TRUE	"Any proce	600	172	13	20972	0.075581395	0.021666667
GO:BP	GO:19040: regulation	0.017082	TRUE	"Any proce	258	682	18	20972	0.026392962	0.069767442
GO:BP	GO:00061: regulation	0.0171	TRUE	"Any proce	55	1172	9	20972	0.007679181	0.163636364
GO:BP	GO:00381: macrophag	0.0171	TRUE	"The serie	55	1172	9	20972	0.007679181	0.163636364

GO:BP	GO:20000! negative re	0.0171	TRUE	"Any proce	55	1172	9	20972	0.007679181	0.163636364
GO:BP	GO:00331! regulation	0.0171	TRUE	"Any proce	55	1172	9	20972	0.007679181	0.163636364
GO:BP	GO:00308! regulation	0.0171	TRUE	"Any proce	55	1172	9	20972	0.007679181	0.163636364
GO:BP	GO:00342! regulation	0.017102	TRUE	"Any proce	87	1172	12	20972	0.010238908	0.137931034
GO:BP	GO:00712! cellular res	0.017102	TRUE	"Any proce	87	1172	12	20972	0.010238908	0.137931034
GO:BP	GO:00480! ephrin rec	0.017102	TRUE	"The serie	87	1172	12	20972	0.010238908	0.137931034
GO:BP	GO:00460! ADP catab	0.017136	TRUE	"The chem	100	294	6	20972	0.020408163	0.06
GO:BP	GO:00456! regulation	0.017136	TRUE	"Any proce	134	294	7	20972	0.023809524	0.052238806
GO:BP	GO:00162! telomere c	0.017162	TRUE	"A process	27	1172	6	20972	0.005119454	0.222222222
GO:BP	GO:00105! positive re	0.017162	TRUE	"Any proce	27	1172	6	20972	0.005119454	0.222222222
GO:BP	GO:00216! cerebellar	0.017232	TRUE	"The proce	19	1172	5	20972	0.004266212	0.263157895
GO:BP	GO:00076! chemosen	0.017232	TRUE	"Behavior	19	1172	5	20972	0.004266212	0.263157895
GO:BP	GO:00018! trophector	0.017232	TRUE	"The proce	19	1172	5	20972	0.004266212	0.263157895
GO:BP	GO:00096! response t	0.017232	TRUE	"Any proce	19	1172	5	20972	0.004266212	0.263157895
GO:BP	GO:00427! regulation	0.017232	TRUE	"Any proce	19	1172	5	20972	0.004266212	0.263157895
GO:BP	GO:00068! dicarboxyli	0.017264	TRUE	"The direc	104	682	10	20972	0.014662757	0.096153846
GO:BP	GO:00426! muscle cel	0.017269	TRUE	"The proce	388	172	10	20972	0.058139535	0.025773196
GO:BP	GO:00455! negative re	0.017272	TRUE	"Any proce	665	682	36	20972	0.052785924	0.054135338
GO:BP	GO:00171! negative re	0.0174	TRUE	"Any proce	130	227	6	20972	0.026431718	0.046153846
GO:BP	GO:00486! neuron fat	0.01741	TRUE	"The proce	71	682	8	20972	0.011730205	0.112676056
GO:BP	GO:00484! synaptic ve	0.01741	TRUE	"A vesicle-	71	682	8	20972	0.011730205	0.112676056
GO:BP	GO:00611! mammary	0.01741	TRUE	"The proce	71	682	8	20972	0.011730205	0.112676056
GO:BP	GO:00510! actin filam	0.017534	TRUE	"The asser	159	682	13	20972	0.019061584	0.081761006
GO:BP	GO:00300! establishm	0.017534	TRUE	"The speci	159	682	13	20972	0.019061584	0.081761006
GO:BP	GO:00720! nephron d	0.017534	TRUE	"The proce	159	682	13	20972	0.019061584	0.081761006
GO:BP	GO:00508! positive re	0.017569	TRUE	"Any proce	380	227	11	20972	0.04845815	0.028947368
GO:BP	GO:00328! cellular res	0.017569	TRUE	"Any proce	677	227	16	20972	0.070484581	0.023633678
GO:BP	GO:00355! protein K2	0.017598	TRUE	"A protein	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00355! G protein-i	0.017598	TRUE	"A G prote	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00706! centromer	0.017598	TRUE	"The cell c	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00312! auditory b	0.017598	TRUE	"The beha	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:19026! regulation	0.017598	TRUE	"Any proce	12	1172	4	20972	0.003412969	0.333333333

GO:BP	GO:19029!	regulation	0.017598	TRUE	"Any proce	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00336!	negative re	0.017598	TRUE	"Any proce	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00019!	G protein-i	0.017598	TRUE	"The serie	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00105!	positive re	0.017598	TRUE	"Any proce	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00702!	establishm	0.017598	TRUE	"The direc	12	1172	4	20972	0.003412969	0.333333333
GO:BP	GO:00019!	negative re	0.0176	TRUE	"Any proce	36	1172	7	20972	0.005972696	0.194444444
GO:BP	GO:19907!	response t	0.0176	TRUE	"Any proce	36	1172	7	20972	0.005972696	0.194444444
GO:BP	GO:00181!	peptidyl-a	0.017642	TRUE	"The altera	495	227	13	20972	0.057268722	0.026262626
GO:BP	GO:00105!	positive re	0.017643	TRUE	"Any proce	27	227	3	20972	0.013215859	0.111111111
GO:BP	GO:00363!	platelet m	0.017643	TRUE	"Generatic	27	227	3	20972	0.013215859	0.111111111
GO:BP	GO:00903!	positive re	0.017643	TRUE	"Any proce	27	227	3	20972	0.013215859	0.111111111
GO:BP	GO:00019!	response t	0.017643	TRUE	"Any proce	27	227	3	20972	0.013215859	0.111111111
GO:BP	GO:00650!	regulation	0.017643	TRUE	"Any proce	1070	227	22	20972	0.0969163	0.020560748
GO:BP	GO:00514!	neuron ap	0.017749	TRUE	"Any apop	293	294	11	20972	0.037414966	0.037542662
GO:BP	GO:00519!	regulation	0.017796	TRUE	"Any proce	223	227	8	20972	0.035242291	0.035874439
GO:BP	GO:00069!	regulation	0.01787	TRUE	"Any proce	56	682	7	20972	0.01026393	0.125
GO:BP	GO:00382!	TORC2 sig	0.017886	TRUE	"A series o	21	294	3	20972	0.010204082	0.142857143
GO:BP	GO:00363!	sodium ior	0.017886	TRUE	"The direc	21	294	3	20972	0.010204082	0.142857143
GO:BP	GO:19038!	positive re	0.017886	TRUE	"Any proce	21	294	3	20972	0.010204082	0.142857143
GO:BP	GO:00108!	regulation	0.017886	TRUE	"Any proce	21	294	3	20972	0.010204082	0.142857143
GO:BP	GO:19051!	positive re	0.01799	TRUE	"Any proce	43	294	4	20972	0.013605442	0.093023256
GO:BP	GO:00511!	positive re	0.01799	TRUE	"Any proce	43	294	4	20972	0.013605442	0.093023256
GO:BP	GO:00424!	eye photo	0.01799	TRUE	"Developm	43	294	4	20972	0.013605442	0.093023256
GO:BP	GO:19034!	regulation	0.018004	TRUE	"Any proce	99	1172	13	20972	0.01109215	0.131313131
GO:BP	GO:00316!	response t	0.018024	TRUE	"Any proce	475	294	15	20972	0.051020408	0.031578947
GO:BP	GO:19033!	regulation	0.018035	TRUE	"Any proce	236	1172	24	20972	0.020477816	0.101694915
GO:BP	GO:00329!	regulation	0.018081	TRUE	"Any proce	382	227	11	20972	0.04845815	0.028795812
GO:BP	GO:00610!	regulation	0.018091	TRUE	"Any proce	42	682	6	20972	0.008797654	0.142857143
GO:BP	GO:01500!	regulation	0.018091	TRUE	"Any proce	42	682	6	20972	0.008797654	0.142857143
GO:BP	GO:00506!	regulation	0.018091	TRUE	"Any proce	239	682	17	20972	0.024926686	0.071129707
GO:BP	GO:19002!	regulation	0.018091	TRUE	"Any proce	42	682	6	20972	0.008797654	0.142857143
GO:BP	GO:00320!	positive re	0.018091	TRUE	"Any proce	88	682	9	20972	0.013196481	0.102272727

GO:BP	GO:00329: mononucleotide	0.018091	TRUE	"The expansion of the genome"	302	682	20	20972	0.029325513	0.066225166
GO:BP	GO:00328: positive regulation of gene expression	0.018091	TRUE	"Any process that results in the production of a functional gene product"	42	682	6	20972	0.008797654	0.142857143
GO:BP	GO:00381: ERBB signaling pathway	0.018091	TRUE	"The series of events starting with the binding of a ligand to a receptor tyrosine kinase, leading to the activation of a signaling pathway"	179	682	14	20972	0.020527859	0.078212291
GO:BP	GO:00219: central nervous system development	0.018091	TRUE	"Generative processes that lead to the formation of the central nervous system"	29	682	5	20972	0.007331378	0.172413793
GO:BP	GO:00315: protein complex assembly	0.018091	TRUE	"A localization process that involves the assembly of a protein complex"	88	682	9	20972	0.013196481	0.102272727
GO:BP	GO:00224: membrane organization	0.018091	TRUE	"The initial organization of the membrane"	88	682	9	20972	0.013196481	0.102272727
GO:BP	GO:00020: positive regulation of gene expression	0.018091	TRUE	"Any process that results in the production of a functional gene product"	29	682	5	20972	0.007331378	0.172413793
GO:BP	GO:00019: negative regulation of gene expression	0.018091	TRUE	"Any process that results in the production of a functional gene product"	29	682	5	20972	0.007331378	0.172413793
GO:BP	GO:00075: body fluid homeostasis	0.018091	TRUE	"The contraction of the body fluid"	88	682	9	20972	0.013196481	0.102272727
GO:BP	GO:00085: female gamete development	0.018091	TRUE	"The process of female gamete development"	105	682	10	20972	0.014662757	0.095238095
GO:BP	GO:00093: 2'-deoxyribonucleic acid metabolism	0.018091	TRUE	"The chemical reaction that involves the metabolism of 2'-deoxyribonucleic acid"	42	682	6	20972	0.008797654	0.142857143
GO:BP	GO:00328: cellular response to stress	0.018103	TRUE	"Any process that results in the production of a functional gene product"	252	294	10	20972	0.034013605	0.03968254
GO:BP	GO:00511: positive regulation of gene expression	0.018184	TRUE	"Any process that results in the production of a functional gene product"	66	1172	10	20972	0.008532423	0.151515152
GO:BP	GO:00149: smooth muscle cell development	0.018184	TRUE	"The order of the smooth muscle cell development"	66	1172	10	20972	0.008532423	0.151515152
GO:BP	GO:00026: regulation of gene expression	0.018235	TRUE	"Any process that results in the production of a functional gene product"	572	294	17	20972	0.057823129	0.02972028
GO:BP	GO:00352: endocrine system development	0.018289	TRUE	"Progression of the endocrine system development"	136	294	7	20972	0.023809524	0.051470588
GO:BP	GO:00519: negative regulation of gene expression	0.018289	TRUE	"Any process that results in the production of a functional gene product"	136	294	7	20972	0.023809524	0.051470588
GO:BP	GO:00080: regulation of gene expression	0.018326	TRUE	"Any process that results in the production of a functional gene product"	204	172	7	20972	0.040697674	0.034313725
GO:BP	GO:00488: cell motility	0.018342	TRUE	"Any process that results in the production of a functional gene product"	1794	172	27	20972	0.156976744	0.015050167
GO:BP	GO:00030: heart morphogenesis	0.018393	TRUE	"The development of the heart"	263	1172	26	20972	0.0221843	0.098859316
GO:BP	GO:00488: multicellular organismal development	0.018445	TRUE	"Any process that results in the production of a functional gene product"	839	682	43	20972	0.063049853	0.05125149
GO:BP	GO:00063: mitochondrial protein import	0.018496	TRUE	"The synthesis of mitochondrial protein"	28	172	3	20972	0.01744186	0.107142857
GO:BP	GO:00481: regulation of gene expression	0.018501	TRUE	"Any process that results in the production of a functional gene product"	88	1172	12	20972	0.010238908	0.136363636
GO:BP	GO:00320: positive regulation of gene expression	0.018501	TRUE	"Any process that results in the production of a functional gene product"	88	1172	12	20972	0.010238908	0.136363636
GO:BP	GO:00193: pyridine nucleotide metabolism	0.018537	TRUE	"The chemical reaction that involves the metabolism of pyridine nucleotide"	102	294	6	20972	0.020408163	0.058823529
GO:BP	GO:00511: import into nucleus	0.018587	TRUE	"The direction of the import into the nucleus"	172	1172	19	20972	0.016211604	0.110465116
GO:BP	GO:19021: regulation of gene expression	0.018695	TRUE	"Any process that results in the production of a functional gene product"	329	227	10	20972	0.044052863	0.030395137
GO:BP	GO:19030: positive regulation of gene expression	0.018707	TRUE	"Any process that results in the production of a functional gene product"	276	227	9	20972	0.039647577	0.032608696
GO:BP	GO:00459: regulation of gene expression	0.018743	TRUE	"Any process that results in the production of a functional gene product"	92	227	5	20972	0.022026432	0.054347826
GO:BP	GO:00466: lymphocyte development	0.018743	TRUE	"A change in the lymphocyte development"	811	227	18	20972	0.079295154	0.022194821
GO:BP	GO:00610: hepaticobiliary system development	0.018803	TRUE	"The progression of the hepaticobiliary system development"	123	1172	15	20972	0.012798635	0.12195122
GO:BP	GO:00513: regulation of gene expression	0.018849	TRUE	"Any process that results in the production of a functional gene product"	277	1172	27	20972	0.023037543	0.097472924

GO:BP	GO:00352: tube morphogenesis	0.018883	TRUE	"The process of forming a tube or tube-like structure"	877	227	19	20972	0.083700441	0.021664766
GO:BP	GO:00316: regulation of cell growth	0.018896	TRUE	"Any process that results in the growth of a cell"	385	227	11	20972	0.04845815	0.028571429
GO:BP	GO:00506: regulation of gene expression	0.018896	TRUE	"Any process that results in the expression of a gene"	133	227	6	20972	0.026431718	0.045112782
GO:BP	GO:00987: L-glutamate transport	0.018942	TRUE	"The direct transport of L-glutamate"	18	682	4	20972	0.005865103	0.222222222
GO:BP	GO:00381: angiotensin transport	0.018942	TRUE	"A G protein-coupled receptor-mediated transport of angiotensin"	18	682	4	20972	0.005865103	0.222222222
GO:BP	GO:00430: negative regulation of gene expression	0.018942	TRUE	"Any process that results in the downregulation of gene expression"	18	682	4	20972	0.005865103	0.222222222
GO:BP	GO:19904: embryonic development	0.018942	TRUE	"The process of developing from an embryo"	18	682	4	20972	0.005865103	0.222222222
GO:BP	GO:00069: regulation of cell cycle	0.018957	TRUE	"Any process that results in the regulation of the cell cycle"	56	1172	9	20972	0.007679181	0.160714286
GO:BP	GO:00350: response to stress	0.018957	TRUE	"Any process that results in the response to stress"	46	1172	8	20972	0.006825939	0.173913043
GO:BP	GO:00423: regulation of cell growth	0.018957	TRUE	"Any process that results in the growth of a cell"	560	1172	47	20972	0.040102389	0.083928571
GO:BP	GO:00346: nucleobase metabolic process	0.018985	TRUE	"The chemical reaction of nucleobases"	526	294	16	20972	0.054421769	0.030418251
GO:BP	GO:00300: actin filament-based process	0.018985	TRUE	"Movement of actin filaments"	137	294	7	20972	0.023809524	0.051094891
GO:BP	GO:00506: RNA transport	0.019178	TRUE	"The direct transport of RNA"	160	1172	18	20972	0.015358362	0.1125
GO:BP	GO:00506: nucleic acid transport	0.019178	TRUE	"The direct transport of nucleic acids"	160	1172	18	20972	0.015358362	0.1125
GO:BP	GO:00508: regulation of gene expression	0.019217	TRUE	"Any process that results in the expression of a gene"	386	227	11	20972	0.04845815	0.028497409
GO:BP	GO:00343: adherens junction assembly	0.019221	TRUE	"A process of forming a junction between cells"	57	682	7	20972	0.01026393	0.122807018
GO:BP	GO:00519: synaptic transmission	0.019221	TRUE	"The vesicle-mediated transport of neurotransmitters"	57	682	7	20972	0.01026393	0.122807018
GO:BP	GO:00860: cell communication	0.019221	TRUE	"Any process that results in the communication between cells"	57	682	7	20972	0.01026393	0.122807018
GO:BP	GO:00076: visual behavior	0.019221	TRUE	"The behavior of an organism in response to visual stimuli"	57	682	7	20972	0.01026393	0.122807018
GO:BP	GO:19000: positive regulation of gene expression	0.019252	TRUE	"Any process that results in the upregulation of gene expression"	44	294	4	20972	0.013605442	0.090909091
GO:BP	GO:00091: purine nucleotide metabolic process	0.019252	TRUE	"The chemical reaction of purine nucleotides"	103	294	6	20972	0.020408163	0.058252427
GO:BP	GO:00165: protein ubiquitination	0.019252	TRUE	"The process of marking proteins for degradation"	726	294	20	20972	0.068027211	0.027548209
GO:BP	GO:00074: sensory perception	0.019252	TRUE	"The process of receiving and responding to sensory stimuli"	625	294	18	20972	0.06122449	0.0288
GO:BP	GO:00215: cerebellum development	0.019252	TRUE	"The process of developing the cerebellum"	44	294	4	20972	0.013605442	0.090909091
GO:BP	GO:00091: purine ribonucleotide metabolic process	0.019252	TRUE	"The chemical reaction of purine ribonucleotides"	103	294	6	20972	0.020408163	0.058252427
GO:BP	GO:00096: response to stimulus	0.019337	TRUE	"Any process that results in the response to a stimulus"	214	294	9	20972	0.030612245	0.042056075
GO:BP	GO:01100: regulation of cell growth	0.019345	TRUE	"Any process that results in the growth of a cell"	262	682	18	20972	0.026392962	0.06870229
GO:BP	GO:00450: regulated gene expression	0.019421	TRUE	"A process of regulating gene expression"	227	227	8	20972	0.035242291	0.035242291
GO:BP	GO:00344: substrate transport	0.019458	TRUE	"The morphological process of transporting a substrate"	100	1172	13	20972	0.01109215	0.13
GO:BP	GO:00447: metaphase chromosome segregation	0.019458	TRUE	"The cell cycle process of separating chromosomes"	100	1172	13	20972	0.01109215	0.13
GO:BP	GO:00216: nerve development	0.019474	TRUE	"The process of developing a nerve"	93	227	5	20972	0.022026432	0.053763441
GO:BP	GO:00457: positive regulation of gene expression	0.019474	TRUE	"Any process that results in the upregulation of gene expression"	134	227	6	20972	0.026431718	0.044776119

GO:BP	GO:000631: DNA-temp	0.019474	TRUE	"The exter	134	227	6	20972	0.026431718	0.044776119
GO:BP	GO:003571: sodium ion	0.019559	TRUE	"A process	181	682	14	20972	0.020527859	0.077348066
GO:BP	GO:000631: transcripti	0.019688	TRUE	"A transcri	173	1172	19	20972	0.016211604	0.10982659
GO:BP	GO:004821: mRNA stat	0.019714	TRUE	"Preventio	72	294	5	20972	0.017006803	0.069444444
GO:BP	GO:006151: actin filam	0.019747	TRUE	"A process	162	682	13	20972	0.019061584	0.080246914
GO:BP	GO:004661: positive re	0.019762	TRUE	"Any proce	73	682	8	20972	0.011730205	0.109589041
GO:BP	GO:199081: cellular res	0.019779	TRUE	"Any proce	22	294	3	20972	0.010204082	0.136363636
GO:BP	GO:200101: regulation	0.019779	TRUE	"Any proce	22	294	3	20972	0.010204082	0.136363636
GO:BP	GO:000681: intracellul	0.019779	TRUE	"A homeos	22	294	3	20972	0.010204082	0.136363636
GO:BP	GO:000221: stimulator	0.019779	TRUE	"The serie	22	294	3	20972	0.010204082	0.136363636
GO:BP	GO:006061: regulation	0.019779	TRUE	"Any proce	529	294	16	20972	0.054421769	0.030245747
GO:BP	GO:199081: response t	0.019779	TRUE	"Any proce	22	294	3	20972	0.010204082	0.136363636
GO:BP	GO:003291: secretion t	0.019858	TRUE	"The contr	43	682	6	20972	0.008797654	0.139534884
GO:BP	GO:005091: negative re	0.019858	TRUE	"Any proce	43	682	6	20972	0.008797654	0.139534884
GO:BP	GO:000921: deoxyribo	0.019858	TRUE	"The chem	43	682	6	20972	0.008797654	0.139534884
GO:BP	GO:006051: epithelial t	0.020032	TRUE	"The proce	333	227	10	20972	0.044052863	0.03003003
GO:BP	GO:190331: positive re	0.020035	TRUE	"Any proce	136	1172	16	20972	0.013651877	0.117647059
GO:BP	GO:003521: endocrine	0.020035	TRUE	"Progressi	136	1172	16	20972	0.013651877	0.117647059
GO:BP	GO:004571: regulation	0.020068	TRUE	"Any proce	292	1172	28	20972	0.023890785	0.095890411
GO:BP	GO:007051: protein K6	0.020092	TRUE	"A protein	67	1172	10	20972	0.008532423	0.149253731
GO:BP	GO:009751: lamellipod	0.020113	TRUE	"A process	89	1172	12	20972	0.010238908	0.134831461
GO:BP	GO:004561: positive re	0.020146	TRUE	"Any proce	135	227	6	20972	0.026431718	0.044444444
GO:BP	GO:001091: negative re	0.020171	TRUE	"Any proce	125	682	11	20972	0.016129032	0.088
GO:BP	GO:005101: regulation	0.02019	TRUE	"Any proce	485	682	28	20972	0.041055718	0.057731959
GO:BP	GO:007151: cellular res	0.020192	TRUE	"Any proce	257	294	10	20972	0.034013605	0.038910506
GO:BP	GO:008601: membrane	0.020198	TRUE	"The proce	30	682	5	20972	0.007331378	0.166666667
GO:BP	GO:005041: arachidon	0.020198	TRUE	"The contr	30	682	5	20972	0.007331378	0.166666667
GO:BP	GO:190561: regulation	0.020198	TRUE	"Any proce	30	682	5	20972	0.007331378	0.166666667
GO:BP	GO:003101: actomyosi	0.020198	TRUE	"A process	202	682	15	20972	0.021994135	0.074257426
GO:BP	GO:190431: cellular res	0.020198	TRUE	"Any proce	30	682	5	20972	0.007331378	0.166666667
GO:BP	GO:190391: arachidon	0.020198	TRUE	"The direc	30	682	5	20972	0.007331378	0.166666667
GO:BP	GO:002241: positive re	0.020198	TRUE	"Any proce	328	682	21	20972	0.030791789	0.06402439

GO:BP	GO:004871 oligodendr	0.020205	TRUE	"The proce	104	172	5	20972	0.029069767	0.048076923
GO:BP	GO:000901 macromoli	0.020244	TRUE	"The chem	1444	172	23	20972	0.13372093	0.015927978
GO:BP	GO:001091 regulation	0.020244	TRUE	"Any proce	29	172	3	20972	0.01744186	0.103448276
GO:BP	GO:000591 regulation	0.020244	TRUE	"Any proce	29	172	3	20972	0.01744186	0.103448276
GO:BP	GO:003121 positive re	0.020263	TRUE	"Any proce	28	1172	6	20972	0.005119454	0.214285714
GO:BP	GO:003511 embryonic	0.020263	TRUE	"The proce	28	1172	6	20972	0.005119454	0.214285714
GO:BP	GO:000761 locomotor	0.020272	TRUE	"The speci	216	294	9	20972	0.030612245	0.041666667
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GO:BP	GO:00024' antigen pri	0.029521	TRUE	"The proce	14	1172	4	20972	0.003412969	0.285714286

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GO:BP	GO:00215: facial nerv	0.030205	TRUE	"The proce	11	682	3	20972	0.004398827	0.272727273
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GO:BP	GO:002161	cranial nerve development	0.030205	TRUE	"The process of cranial nerve development"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:004241	catecholamine metabolic process	0.030205	TRUE	"The chemical processes of catecholamine metabolism"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:003551	purinergic transmission	0.030205	TRUE	"The series of events in purinergic transmission"	34	682	5	20972	0.007331378	0.147058824
GO:BP	GO:003591	aortic development	0.030205	TRUE	"The process of aortic development"	64	682	7	20972	0.01026393	0.109375
GO:BP	GO:000191	endothelial cell proliferation	0.030205	TRUE	"The multi-step process of endothelial cell proliferation"	154	682	12	20972	0.017595308	0.077922078
GO:BP	GO:003231	positive regulation of cell proliferation	0.030205	TRUE	"Any process that positively regulates cell proliferation"	34	682	5	20972	0.007331378	0.147058824
GO:BP	GO:003191	locomotion	0.030205	TRUE	"Self-propulsion of an organism"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:003231	positive regulation of cell proliferation	0.030205	TRUE	"Any process that positively regulates cell proliferation"	34	682	5	20972	0.007331378	0.147058824
GO:BP	GO:003221	purine nucleotide metabolism	0.030205	TRUE	"Any process involving the metabolism of purine nucleotides"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:004201	olfactory behavior	0.030205	TRUE	"The behavioral response to olfactory stimuli"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:003261	interleukin production	0.030205	TRUE	"The appearance and production of interleukins"	64	682	7	20972	0.01026393	0.109375
GO:BP	GO:003261	regulation of cell proliferation	0.030205	TRUE	"Any process that regulates cell proliferation"	64	682	7	20972	0.01026393	0.109375
GO:BP	GO:000711	leukocyte chemotaxis	0.030205	TRUE	"The attachment and movement of leukocytes towards a chemical stimulus"	413	682	24	20972	0.035190616	0.05811138
GO:BP	GO:003041	peristalsis	0.030205	TRUE	"A wavelike contraction of smooth muscle"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:009741	dendrite extension	0.030205	TRUE	"Long distance transport of materials in a dendrite"	34	682	5	20972	0.007331378	0.147058824
GO:BP	GO:004361	engulfment	0.030205	TRUE	"The removal of particles or cells by phagocytosis or pinocytosis"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:004671	GDP metal ion binding	0.030205	TRUE	"The chemical binding of GDP to metal ions"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:001031	membrane organization	0.030205	TRUE	"The infolding and organization of membranes"	64	682	7	20972	0.01026393	0.109375
GO:BP	GO:000911	deoxyribonucleic acid metabolic process	0.030205	TRUE	"The chemical processes of deoxyribonucleic acid metabolism"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:002171	olfactory behavior	0.030205	TRUE	"The process of olfactory behavior"	34	682	5	20972	0.007331378	0.147058824
GO:BP	GO:200121	regulation of cell proliferation	0.030205	TRUE	"Any process that regulates cell proliferation"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:190521	negative regulation of cell proliferation	0.030205	TRUE	"Any process that negatively regulates cell proliferation"	11	682	3	20972	0.004398827	0.272727273
GO:BP	GO:010611	cold-induced response	0.030244	TRUE	"The process of response to cold stimuli"	151	227	6	20972	0.026431718	0.039735099
GO:BP	GO:004351	ear development	0.030274	TRUE	"The process of ear development"	236	682	16	20972	0.023460411	0.06779661
GO:BP	GO:001501	protein transport	0.030359	TRUE	"The directed movement of proteins across or through membranes"	1422	172	22	20972	0.127906977	0.015471167
GO:BP	GO:190001	regulation of cell proliferation	0.030382	TRUE	"Any process that regulates cell proliferation"	73	172	4	20972	0.023255814	0.054794521
GO:BP	GO:000741	neuroblast differentiation	0.030382	TRUE	"The expansion and differentiation of neuroblasts"	73	172	4	20972	0.023255814	0.054794521
GO:BP	GO:007061	protein metabolism	0.030383	TRUE	"A protein's chemical and physical changes"	927	294	23	20972	0.078231293	0.024811219
GO:BP	GO:004241	catecholamine metabolic process	0.030456	TRUE	"The chemical processes of catecholamine metabolism"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:004811	astrocyte differentiation	0.030456	TRUE	"A change in the state of an astrocyte"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:000911	purine nucleotide metabolism	0.030456	TRUE	"The chemical processes of purine nucleotide metabolism"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:000971	catecholamine metabolic process	0.030456	TRUE	"The chemical processes of catecholamine metabolism"	22	1172	5	20972	0.004266212	0.227272727

GO:BP	GO:00484: sympathetic	0.030456	TRUE	"The process of sympathetic nervous system activity"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:00701: establishment of	0.030456	TRUE	"The direct establishment of a biological process"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:20010: regulation of	0.030456	TRUE	"Any process of regulation"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:00519: positive regulation of	0.030456	TRUE	"Any process of positive regulation"	22	1172	5	20972	0.004266212	0.227272727
GO:BP	GO:00075: muscle organ	0.030467	TRUE	"The process of muscle organ function"	362	172	9	20972	0.052325581	0.024861878
GO:BP	GO:00510: positive regulation of	0.030467	TRUE	"Any process of positive regulation"	818	172	15	20972	0.087209302	0.018337408
GO:BP	GO:00481: fibroblast junction	0.030526	TRUE	"The multi-cellular fibroblast junction"	107	227	5	20972	0.022026432	0.046728972
GO:BP	GO:00030: heart process	0.030526	TRUE	"A circulatory heart process"	251	227	8	20972	0.035242291	0.03187251
GO:BP	GO:00485: embryonic	0.030526	TRUE	"Morphogenesis of embryonic tissue"	305	227	9	20972	0.039647577	0.029508197
GO:BP	GO:00507: negative regulation of	0.030542	TRUE	"Any process of negative regulation"	52	294	4	20972	0.013605442	0.076923077
GO:BP	GO:00016: response to	0.030686	TRUE	"Any process of response to stimulus"	278	294	10	20972	0.034013605	0.035971223
GO:BP	GO:00021: cytoplasmic	0.030758	TRUE	"The chemical process of cytoplasmic function"	200	227	7	20972	0.030837004	0.035
GO:BP	GO:00901: regulation of	0.030759	TRUE	"Any process of regulation"	61	1172	9	20972	0.007679181	0.147540984
GO:BP	GO:19033: positive regulation of	0.030759	TRUE	"Any process of positive regulation"	131	1172	15	20972	0.012798635	0.114503817
GO:BP	GO:00518: regulation of	0.030759	TRUE	"Any process of regulation"	61	1172	9	20972	0.007679181	0.147540984
GO:BP	GO:00457: negative regulation of	0.030759	TRUE	"Any process of negative regulation"	61	1172	9	20972	0.007679181	0.147540984
GO:BP	GO:00360: cellular response	0.030759	TRUE	"Any process of cellular response"	61	1172	9	20972	0.007679181	0.147540984
GO:BP	GO:00717: nitrogen catabolism	0.030887	TRUE	"The direct process of nitrogen catabolism"	798	227	17	20972	0.074889868	0.021303258
GO:BP	GO:19048: cranial skeletal	0.030889	TRUE	"The process of cranial skeletal development"	72	1172	10	20972	0.008532423	0.138888889
GO:BP	GO:19024: L-alpha-amino acid	0.030889	TRUE	"The direct process of L-alpha-amino acid metabolism"	72	1172	10	20972	0.008532423	0.138888889
GO:BP	GO:00098: negative regulation of	0.030922	TRUE	"Any process of negative regulation"	363	172	9	20972	0.052325581	0.024793388
GO:BP	GO:00421: regulation of	0.030961	TRUE	"Any process of regulation"	182	1172	19	20972	0.016211604	0.104395604
GO:BP	GO:00328: cellular response	0.030983	TRUE	"Any process of cellular response"	252	227	8	20972	0.035242291	0.031746032
GO:BP	GO:00481: negative regulation of	0.030983	TRUE	"Any process of negative regulation"	35	227	3	20972	0.013215859	0.085714286
GO:BP	GO:00331: regulation of	0.030983	TRUE	"Any process of regulation"	35	227	3	20972	0.013215859	0.085714286
GO:BP	GO:00468: filopodium	0.030983	TRUE	"The assertion of filopodium function"	68	227	4	20972	0.017621145	0.058823529
GO:BP	GO:00148: release of	0.030983	TRUE	"The process of release of a substance"	35	227	3	20972	0.013215859	0.085714286
GO:BP	GO:19027: positive regulation of	0.030983	TRUE	"Any process of positive regulation"	35	227	3	20972	0.013215859	0.085714286
GO:BP	GO:00457: positive regulation of	0.030983	TRUE	"Any process of positive regulation"	35	227	3	20972	0.013215859	0.085714286
GO:BP	GO:00508: regulation of	0.031017	TRUE	"Any process of regulation"	374	1172	33	20972	0.028156997	0.088235294
GO:BP	GO:00082: insulin receptor	0.031136	TRUE	"The series of insulin receptor activation"	174	172	6	20972	0.034883721	0.034482759
GO:BP	GO:19046: D-glucose	0.031136	TRUE	"The process of D-glucose metabolism"	95	1172	12	20972	0.010238908	0.126315789

GO:BP	GO:00456: negative re	0.031136	TRUE	"Any proce	95	1172	12	20972	0.010238908	0.126315789
GO:BP	GO:00400: regulation	0.031166	TRUE	"Any proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00988: postsynap	0.031166	TRUE	"A recepto	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00432: positive re	0.031166	TRUE	"Any proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:20003: negative re	0.031166	TRUE	"Any proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00607: regulation	0.031166	TRUE	"Any proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00995: neurotran	0.031166	TRUE	"A recepto	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00072: neuromusi	0.031166	TRUE	"The proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:01402: postsynap	0.031166	TRUE	"A vesicle-	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:19038: positive re	0.031166	TRUE	"Any proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00305: positive re	0.031166	TRUE	"Any proce	31	1172	6	20972	0.005119454	0.193548387
GO:BP	GO:00487: astrocyte c	0.031195	TRUE	"The proce	83	294	5	20972	0.017006803	0.060240964
GO:BP	GO:19028: negative re	0.031195	TRUE	"Any proce	27	294	3	20972	0.010204082	0.111111111
GO:BP	GO:00990: postsynap	0.031195	TRUE	"The aggre	83	294	5	20972	0.017006803	0.060240964
GO:BP	GO:00105: regulation	0.031195	TRUE	"Any proce	83	294	5	20972	0.017006803	0.060240964
GO:BP	GO:00106: positive re	0.031195	TRUE	"Any proce	27	294	3	20972	0.010204082	0.111111111
GO:BP	GO:01410: positive re	0.031195	TRUE	"Any proce	27	294	3	20972	0.010204082	0.111111111
GO:BP	GO:00363: platelet m	0.031195	TRUE	"Generatic	27	294	3	20972	0.010204082	0.111111111
GO:BP	GO:00000: mitotic sis	0.031195	TRUE	"The cell c	194	294	8	20972	0.027210884	0.041237113
GO:BP	GO:00550: cardiac cel	0.031195	TRUE	"The proce	83	294	5	20972	0.017006803	0.060240964
GO:BP	GO:00064: RNA locali	0.031253	TRUE	"A process	201	227	7	20972	0.030837004	0.034825871
GO:BP	GO:00481: fibroblast j	0.03127	TRUE	"The multi	107	1172	13	20972	0.01109215	0.121495327
GO:BP	GO:00226: gland morj	0.03127	TRUE	"The proce	107	1172	13	20972	0.01109215	0.121495327
GO:BP	GO:00435: ear develo	0.031317	TRUE	"The proce	236	294	9	20972	0.030612245	0.038135593
GO:BP	GO:00105: regulation	0.031322	TRUE	"Any proce	364	172	9	20972	0.052325581	0.024725275
GO:BP	GO:00313: positive re	0.031328	TRUE	"Any proce	108	227	5	20972	0.022026432	0.046296296
GO:BP	GO:00451: cell fate cc	0.031347	TRUE	"The cellul	297	172	8	20972	0.046511628	0.026936027
GO:BP	GO:00076: memory	0.031387	TRUE	"The activi	121	172	5	20972	0.029069767	0.041322314
GO:BP	GO:00074: axon guida	0.031387	TRUE	"The chem	234	172	7	20972	0.040697674	0.02991453
GO:BP	GO:00086: carbohydr	0.031387	TRUE	"The direc	121	172	5	20972	0.029069767	0.041322314
GO:BP	GO:00163: dendrite d	0.031387	TRUE	"The proce	234	172	7	20972	0.040697674	0.02991453
GO:BP	GO:00432: regulation	0.031559	TRUE	"Any proce	422	227	11	20972	0.04845815	0.026066351

GO:BP	GO:005101	positive re	0.031576	TRUE	"Any proce	280	682	18	20972	0.026392962	0.064285714
GO:BP	GO:001051	regulation	0.031689	TRUE	"Any proce	364	227	10	20972	0.044052863	0.027472527
GO:BP	GO:001821	peptidyl-tr	0.031807	TRUE	"The modi	49	682	6	20972	0.008797654	0.12244898
GO:BP	GO:005191	negative re	0.031807	TRUE	"Any proce	136	682	11	20972	0.016129032	0.080882353
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GO:BP	GO:007151	semaphori	0.031807	TRUE	"The serie	49	682	6	20972	0.008797654	0.12244898
GO:BP	GO:003281	response t	0.03193	TRUE	"Any proce	308	227	9	20972	0.039647577	0.029220779
GO:BP	GO:006051	developm	0.031967	TRUE	"The incre	235	172	7	20972	0.040697674	0.029787234
GO:BP	GO:009741	neuron pr	0.031967	TRUE	"The proce	235	172	7	20972	0.040697674	0.029787234
GO:BP	GO:007141	cellular res	0.032012	TRUE	"Any proce	118	294	6	20972	0.020408163	0.050847458
GO:BP	GO:004251	glucose ho	0.032105	TRUE	"Any proce	254	227	8	20972	0.035242291	0.031496063
GO:BP	GO:000701	cell-cell j	0.032147	TRUE	"The aggre	157	1172	17	20972	0.014505119	0.108280255
GO:BP	GO:001401	glial cell pr	0.032163	TRUE	"The multi	53	294	4	20972	0.013605442	0.075471698
GO:BP	GO:003411	cellular res	0.032163	TRUE	"Any proce	53	294	4	20972	0.013605442	0.075471698
GO:BP	GO:000711	enzyme-lir	0.032357	TRUE	"The serie	988	294	24	20972	0.081632653	0.024291498
GO:BP	GO:004311	receptor n	0.032382	TRUE	"The chem	75	172	4	20972	0.023255814	0.053333333
GO:BP	GO:007231	signal tran	0.032466	TRUE	"An intrac	183	1172	19	20972	0.016211604	0.103825137
GO:BP	GO:004851	thymus de	0.032466	TRUE	"The proce	51	1172	8	20972	0.006825939	0.156862745
GO:BP	GO:004211	T cell activ	0.032509	TRUE	"The chang	581	682	31	20972	0.045454545	0.053356282
GO:BP	GO:007011	establishm	0.032509	TRUE	"The direc	22	682	4	20972	0.005865103	0.181818182
GO:BP	GO:000191	positive re	0.032509	TRUE	"Any proce	65	682	7	20972	0.01026393	0.107692308
GO:BP	GO:000691	vesicle tar	0.032509	TRUE	"The proce	65	682	7	20972	0.01026393	0.107692308
GO:BP	GO:001061	regulation	0.032509	TRUE	"Any proce	65	682	7	20972	0.01026393	0.107692308
GO:BP	GO:000241	antigen pri	0.032509	TRUE	"The proce	22	682	4	20972	0.005865103	0.181818182
GO:BP	GO:005191	positive re	0.032509	TRUE	"Any proce	22	682	4	20972	0.005865103	0.181818182
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GO:BP	GO:000911	purine nuc	0.032509	TRUE	"The chem	22	682	4	20972	0.005865103	0.181818182
GO:BP	GO:009731	response t	0.032509	TRUE	"Any proce	238	682	16	20972	0.023460411	0.067226891
GO:BP	GO:003221	methyлатic	0.032518	TRUE	"The proce	238	294	9	20972	0.030612245	0.037815126
GO:BP	GO:190181	positive re	0.032519	TRUE	"Any proce	132	1172	15	20972	0.012798635	0.113636364
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GO:BP	GO:007081	regulation	0.032741	TRUE	"Any proce	37	172	3	20972	0.01744186	0.081081081

GO:BP	GO:00091: purine ribc	0.032771	TRUE	"The chem	119	294	6	20972	0.020408163	0.050420168
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GO:BP	GO:00608: regulation	0.032807	TRUE	"Any proce	255	227	8	20972	0.035242291	0.031372549
GO:BP	GO:00364: synaptic ve	0.032981	TRUE	"The traffi	82	682	8	20972	0.011730205	0.097560976
GO:BP	GO:00600: canonical \	0.033035	TRUE	"A type of	310	227	9	20972	0.039647577	0.029032258
GO:BP	GO:00107: positive re	0.033035	TRUE	"Any proce	36	227	3	20972	0.013215859	0.083333333
GO:BP	GO:00433: CD4-positi	0.033035	TRUE	"The proce	36	227	3	20972	0.013215859	0.083333333
GO:BP	GO:00219: olfactory k	0.033035	TRUE	"The progr	36	227	3	20972	0.013215859	0.083333333
GO:BP	GO:00311: regulation	0.033035	TRUE	"Any proce	36	227	3	20972	0.013215859	0.083333333
GO:BP	GO:00346: response t	0.033035	TRUE	"Any proce	36	227	3	20972	0.013215859	0.083333333
GO:BP	GO:00076: adult walk	0.03306	TRUE	"The beha	35	682	5	20972	0.007331378	0.142857143
GO:BP	GO:00327: positive re	0.03306	TRUE	"Any proce	35	682	5	20972	0.007331378	0.142857143
GO:BP	GO:00423: positive re	0.033074	TRUE	"Any proce	41	1172	7	20972	0.005972696	0.170731707
GO:BP	GO:00906: walking be	0.033074	TRUE	"The beha	41	1172	7	20972	0.005972696	0.170731707
GO:BP	GO:00605: ventral spi	0.033084	TRUE	"The proce	11	172	2	20972	0.011627907	0.181818182
GO:BP	GO:00148: response t	0.033084	TRUE	"Any proce	11	172	2	20972	0.011627907	0.181818182
GO:BP	GO:00709: CRD-medi	0.033084	TRUE	"An mRNA	11	172	2	20972	0.011627907	0.181818182
GO:BP	GO:00605: cell fate cc	0.033084	TRUE	"The comr	11	172	2	20972	0.011627907	0.181818182
GO:BP	GO:00018: liver devel	0.033261	TRUE	"The proce	120	1172	14	20972	0.011945392	0.116666667
GO:BP	GO:00902: regulation	0.033298	TRUE	"Any proce	204	227	7	20972	0.030837004	0.034313725
GO:BP	GO:00512: negative re	0.033309	TRUE	"Any proce	569	294	16	20972	0.054421769	0.028119508
GO:BP	GO:01400: mitotic nu	0.033554	TRUE	"A mitotic	283	294	10	20972	0.034013605	0.035335689
GO:BP	GO:00466: positive re	0.033572	TRUE	"Any proce	73	1172	10	20972	0.008532423	0.136986301
GO:BP	GO:00973: response t	0.033579	TRUE	"Any proce	238	172	7	20972	0.040697674	0.029411765
GO:BP	GO:00457: positive re	0.033597	TRUE	"Any proce	54	294	4	20972	0.013605442	0.074074074
GO:BP	GO:00063: mitochondr	0.033597	TRUE	"The synth	28	294	3	20972	0.010204082	0.107142857
GO:BP	GO:00987: sodium ior	0.033597	TRUE	"The direc	28	294	3	20972	0.010204082	0.107142857
GO:BP	GO:00323: regulation	0.033597	TRUE	"Any proce	54	294	4	20972	0.013605442	0.074074074
GO:BP	GO:00147: positive re	0.033597	TRUE	"Any proce	28	294	3	20972	0.010204082	0.107142857
GO:BP	GO:00344: substrate	0.03365	TRUE	"The morp	100	682	9	20972	0.013196481	0.09
GO:BP	GO:00456: regulation	0.03365	TRUE	"Any proce	218	682	15	20972	0.021994135	0.068807339
GO:BP	GO:01100: regulation	0.03365	TRUE	"Any proce	100	682	9	20972	0.013196481	0.09

GO:BP	GO:00063: DNA recon	0.033664	TRUE	"Any proce	348	1172	31	20972	0.026450512	0.08908046
GO:BP	GO:00427: embryonic	0.033668	TRUE	"The proce	62	1172	9	20972	0.007679181	0.14516129
GO:BP	GO:00700: neuron cel	0.033668	TRUE	"The cellul	62	1172	9	20972	0.007679181	0.14516129
GO:BP	GO:00090: macromol	0.033707	TRUE	"The chem	1444	294	32	20972	0.108843537	0.022160665
GO:BP	GO:00601: regulation	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:19001: negative re	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00715: cellular res	0.03378	TRUE	"Any proce	257	227	8	20972	0.035242291	0.031128405
GO:BP	GO:00724: mesenchy	0.03378	TRUE	"The proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00971: apoptotic :	0.03378	TRUE	"The serie:	612	227	14	20972	0.061674009	0.022875817
GO:BP	GO:20002: positive re	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00105: positive re	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:19039: regulation	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00312: auditory b	0.03378	TRUE	"The beha:	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:19033: regulation	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00706: centromer	0.03378	TRUE	"The cell c	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00726: protein loc	0.03378	TRUE	"A process	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00216: lateral ven	0.03378	TRUE	"The proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00351: genitalia r	0.03378	TRUE	"The proce	12	227	2	20972	0.008810573	0.166666667
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GO:BP	GO:19026: regulation	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00336: negative re	0.03378	TRUE	"Any proce	12	227	2	20972	0.008810573	0.166666667
GO:BP	GO:00335: carbohydr	0.034119	TRUE	"A homeos	258	227	8	20972	0.035242291	0.031007752
GO:BP	GO:00026: positive re	0.034536	TRUE	"Any proce	1111	682	52	20972	0.076246334	0.04680468
GO:BP	GO:00305: androgen i	0.034539	TRUE	"A nuclear	50	682	6	20972	0.008797654	0.12
GO:BP	GO:00363: vascular er	0.034539	TRUE	"The serie:	50	682	6	20972	0.008797654	0.12
GO:BP	GO:00096: response t	0.034669	TRUE	"Any proce	240	682	16	20972	0.023460411	0.066666667
GO:BP	GO:00069: vesicle doc	0.034675	TRUE	"The initial	37	227	3	20972	0.013215859	0.081081081
GO:BP	GO:00219: central nei	0.034675	TRUE	"Generatic	37	227	3	20972	0.013215859	0.081081081
GO:BP	GO:00105: regulation	0.034675	TRUE	"Any proce	37	227	3	20972	0.013215859	0.081081081
GO:BP	GO:19035: release of	0.034675	TRUE	"The direc	37	227	3	20972	0.013215859	0.081081081
GO:BP	GO:00216: cerebellar	0.034675	TRUE	"The proce	37	227	3	20972	0.013215859	0.081081081
GO:BP	GO:00486: neuron fat	0.034675	TRUE	"The proce	71	227	4	20972	0.017621145	0.056338028

GO:BP	GO:00991: presynaptic	0.034709	TRUE	"Any process"	38	172	3	20972	0.01744186	0.078947368
GO:BP	GO:00972: amyloid-beta	0.034709	TRUE	"The process"	38	172	3	20972	0.01744186	0.078947368
GO:BP	GO:00435: regulation	0.034709	TRUE	"Any process"	77	172	4	20972	0.023255814	0.051948052
GO:BP	GO:00156: organic cation	0.034754	TRUE	"The direction"	119	682	10	20972	0.014662757	0.084033613
GO:BP	GO:00400: regulation	0.034782	TRUE	"Any process"	66	682	7	20972	0.01026393	0.106060606
GO:BP	GO:00989: axonal transport	0.034782	TRUE	"The direction"	66	682	7	20972	0.01026393	0.106060606
GO:BP	GO:00218: forebrain c	0.034782	TRUE	"The order"	66	682	7	20972	0.01026393	0.106060606
GO:BP	GO:00027: regulation	0.034956	TRUE	"Any process"	207	227	7	20972	0.030837004	0.033816425
GO:BP	GO:00068: monoatom	0.035077	TRUE	"The direction"	1052	294	25	20972	0.085034014	0.023764259
GO:BP	GO:00510: actin filament	0.035132	TRUE	"The assertion"	159	294	7	20972	0.023809524	0.044025157
GO:BP	GO:00030: circulatory	0.035195	TRUE	"An organ"	595	172	12	20972	0.069767442	0.020168067
GO:BP	GO:00423: positive re	0.035202	TRUE	"Any process"	335	1172	30	20972	0.02559727	0.089552239
GO:BP	GO:00002: nuclear div	0.035249	TRUE	"The division"	450	1172	38	20972	0.032423208	0.084444444
GO:BP	GO:00432: negative re	0.035332	TRUE	"Any process"	85	1172	11	20972	0.009385666	0.129411765
GO:BP	GO:00506: positive re	0.035533	TRUE	"Any process"	146	1172	16	20972	0.013651877	0.109589041
GO:BP	GO:00075: response to	0.035533	TRUE	"Any process"	146	1172	16	20972	0.013651877	0.109589041
GO:BP	GO:19909: response to	0.035613	TRUE	"Any process"	55	294	4	20972	0.013605442	0.072727273
GO:BP	GO:00106: regulation	0.035613	TRUE	"Any process"	55	294	4	20972	0.013605442	0.072727273
GO:BP	GO:00303: respiratory	0.035665	TRUE	"The process"	198	1172	20	20972	0.017064846	0.101010101
GO:BP	GO:00486: regulation	0.035711	TRUE	"Any process"	306	172	8	20972	0.046511628	0.026143791
GO:BP	GO:00357: sodium ion	0.035711	TRUE	"A process"	181	172	6	20972	0.034883721	0.033149171
GO:BP	GO:00303: regulation	0.035755	TRUE	"Any process"	944	294	23	20972	0.078231293	0.024364407
GO:BP	GO:00002: nuclear-tra	0.03577	TRUE	"Shortenin	32	1172	6	20972	0.005119454	0.1875
GO:BP	GO:00018: blastocyst	0.03577	TRUE	"An increa	23	1172	5	20972	0.004266212	0.217391304
GO:BP	GO:00507: negative re	0.03577	TRUE	"Any process"	52	1172	8	20972	0.006825939	0.153846154
GO:BP	GO:00350: negative re	0.03577	TRUE	"Any process"	23	1172	5	20972	0.004266212	0.217391304
GO:BP	GO:00458: positive re	0.03577	TRUE	"Any process"	23	1172	5	20972	0.004266212	0.217391304
GO:BP	GO:00305: BMP signa	0.03577	TRUE	"The series"	159	1172	17	20972	0.014505119	0.106918239
GO:BP	GO:00509: detection	0.03577	TRUE	"The series"	23	1172	5	20972	0.004266212	0.217391304
GO:BP	GO:00611: establishm	0.03577	TRUE	"The speci	23	1172	5	20972	0.004266212	0.217391304
GO:BP	GO:00973: cellular res	0.03577	TRUE	"Any process"	97	1172	12	20972	0.010238908	0.12371134
GO:BP	GO:19018: positive re	0.03577	TRUE	"Any process"	23	1172	5	20972	0.004266212	0.217391304

GO:BP	GO:00897: amino acid	0.03577	TRUE	"The direc	52	1172	8	20972	0.006825939	0.153846154
GO:BP	GO:00090: dosage cor	0.03577	TRUE	"Compens	23	1172	5	20972	0.004266212	0.217391304
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GO:BP	GO:00400: positive re	0.03577	TRUE	"Any proce	32	1172	6	20972	0.005119454	0.1875
GO:BP	GO:00486: positive re	0.03577	TRUE	"Any proce	23	1172	5	20972	0.004266212	0.217391304
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GO:BP	GO:00108: lipid localiz	0.03577	TRUE	"Any proce	451	1172	38	20972	0.032423208	0.084257206
GO:BP	GO:00149: negative re	0.03577	TRUE	"Any proce	23	1172	5	20972	0.004266212	0.217391304
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GO:BP	GO:00331: regulation	0.035863	TRUE	"Any proce	74	1172	10	20972	0.008532423	0.135135135
GO:BP	GO:00705: platelet ag	0.035863	TRUE	"The adhe	74	1172	10	20972	0.008532423	0.135135135
GO:BP	GO:00224: regulation	0.035949	TRUE	"Any proce	493	227	12	20972	0.052863436	0.024340771
GO:BP	GO:00063: DNA-temp	0.035989	TRUE	"The initial	212	1172	21	20972	0.017918089	0.099056604
GO:BP	GO:00072: Rho protei	0.035989	TRUE	"An intrac	134	1172	15	20972	0.012798635	0.111940299
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GO:BP	GO:19030: positive re	0.036116	TRUE	"Any proce	63	1172	9	20972	0.007679181	0.142857143
GO:BP	GO:00709: left/right a	0.036116	TRUE	"The estab	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00721: epithelial c	0.036116	TRUE	"The proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00431: purine-cor	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00513: regulation	0.036116	TRUE	"Any proce	408	1172	35	20972	0.029863481	0.085784314
GO:BP	GO:00309: positive re	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00322: positive re	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00061: GMP biosy	0.036116	TRUE	"The chem	15	1172	4	20972	0.003412969	0.266666667
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GO:BP	GO:00183: peptidyl-ly	0.036116	TRUE	"The acety	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00988: modificati	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00986: regulation	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00607: negative re	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:01500: negative re	0.036116	TRUE	"Any proce	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00024: antigen pri	0.036116	TRUE	"The proce	15	1172	4	20972	0.003412969	0.266666667

GO:BP	GO:00024: T cell medi	0.036116	TRUE	"An immu	15	1172	4	20972	0.003412969	0.266666667
GO:BP	GO:00105: positive re	0.036153	TRUE	"Any proce	160	294	7	20972	0.023809524	0.04375
GO:BP	GO:00328: positive re	0.036238	TRUE	"Any proce	42	1172	7	20972	0.005972696	0.166666667
GO:BP	GO:00218: cerebral cc	0.036395	TRUE	"The radia	29	294	3	20972	0.010204082	0.103448276
GO:BP	GO:00140: glutamate	0.036395	TRUE	"The contr	29	294	3	20972	0.010204082	0.103448276
GO:BP	GO:00342: regulation	0.036395	TRUE	"Any proce	87	294	5	20972	0.017006803	0.057471264
GO:BP	GO:00712: cellular res	0.036395	TRUE	"Any proce	87	294	5	20972	0.017006803	0.057471264
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GO:BP	GO:20006: negative re	0.036395	TRUE	"Any proce	29	294	3	20972	0.010204082	0.103448276
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GO:BP	GO:00080: regulation	0.036476	TRUE	"Any proce	147	1172	16	20972	0.013651877	0.108843537
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GO:BP	GO:00069: autophagy	0.036646	TRUE	"The cellul	599	172	12	20972	0.069767442	0.020033389
GO:BP	GO:19046: cellular res	0.036646	TRUE	"Any proce	39	172	3	20972	0.01744186	0.076923077
GO:BP	GO:00619: process uti	0.036646	TRUE	"A cellular	599	172	12	20972	0.069767442	0.020033389
GO:BP	GO:00708: regulation	0.036646	TRUE	"Any proce	39	172	3	20972	0.01744186	0.076923077
GO:BP	GO:00900: regulation	0.03668	TRUE	"Any proce	209	227	7	20972	0.030837004	0.033492823
GO:BP	GO:00513: regulation	0.036709	TRUE	"Any proce	186	1172	19	20972	0.016211604	0.102150538
GO:BP	GO:00604: epithelium	0.036745	TRUE	"The proce	1234	227	23	20972	0.101321586	0.018638574
GO:BP	GO:00519: negative re	0.036788	TRUE	"Any proce	23	682	4	20972	0.005865103	0.173913043
GO:BP	GO:00509: detection	0.036788	TRUE	"The serie	23	682	4	20972	0.005865103	0.173913043
GO:BP	GO:00024: antigen pri	0.036788	TRUE	"The proce	36	682	5	20972	0.007331378	0.138888889
GO:BP	GO:00604: branching	0.036788	TRUE	"The proce	23	682	4	20972	0.005865103	0.173913043
GO:BP	GO:00030: regulation	0.036788	TRUE	"The regul	51	682	6	20972	0.008797654	0.117647059
GO:BP	GO:00030: regionaliza	0.036788	TRUE	"The patte	422	682	24	20972	0.035190616	0.056872038
GO:BP	GO:19033: regulation	0.036788	TRUE	"Any proce	12	682	3	20972	0.004398827	0.25
GO:BP	GO:00723: intrinsic a	0.036788	TRUE	"The serie	84	682	8	20972	0.011730205	0.095238095
GO:BP	GO:00506: positive re	0.036788	TRUE	"Any proce	200	682	14	20972	0.020527859	0.07
GO:BP	GO:00513: regulation	0.036788	TRUE	"Any proce	12	682	3	20972	0.004398827	0.25
GO:BP	GO:00304: sleep	0.036788	TRUE	"Any proce	23	682	4	20972	0.005865103	0.173913043

GO:BP	GO:00355: protein K2	0.036788	TRUE	"A protein	12	682	3	20972	0.004398827	0.25
GO:BP	GO:00457: positive re	0.036788	TRUE	"Any proce	23	682	4	20972	0.005865103	0.173913043
GO:BP	GO:00344: microtubu	0.036788	TRUE	"Any proce	12	682	3	20972	0.004398827	0.25
GO:BP	GO:20011: regulation	0.036788	TRUE	"Any proce	12	682	3	20972	0.004398827	0.25
GO:BP	GO:00350: regulation	0.036788	TRUE	"Any proce	84	682	8	20972	0.011730205	0.095238095
GO:BP	GO:00329: protein-DN	0.036788	TRUE	"The disag	23	682	4	20972	0.005865103	0.173913043
GO:BP	GO:00219: olfactory li	0.036788	TRUE	"The progr	36	682	5	20972	0.007331378	0.138888889
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GO:BP	GO:00459: positive re	0.036788	TRUE	"Any proce	84	682	8	20972	0.011730205	0.095238095
GO:BP	GO:01400: dendrite a	0.036788	TRUE	"The proce	12	682	3	20972	0.004398827	0.25
GO:BP	GO:19907: response t	0.036788	TRUE	"Any proce	36	682	5	20972	0.007331378	0.138888889
GO:BP	GO:00329: circadian r	0.036788	TRUE	"Any proce	67	682	7	20972	0.01026393	0.104477612
GO:BP	GO:00160: synaptic ve	0.036788	TRUE	"The initial	12	682	3	20972	0.004398827	0.25
GO:BP	GO:00611: establishm	0.036788	TRUE	"The speci	23	682	4	20972	0.005865103	0.173913043
GO:BP	GO:19030: regulation	0.036788	TRUE	"Any proce	375	682	22	20972	0.032258065	0.058666667
GO:BP	GO:00065: ubiquitin-c	0.036813	TRUE	"The chem	680	294	18	20972	0.06122449	0.026470588
GO:BP	GO:00468: regulation	0.03682	TRUE	"Any proce	262	227	8	20972	0.035242291	0.030534351
GO:BP	GO:00466: lymphocyt	0.036976	TRUE	"The expai	295	1172	27	20972	0.023037543	0.091525424
GO:BP	GO:00100: specificati	0.03698	TRUE	"The regio	38	227	3	20972	0.013215859	0.078947368
GO:BP	GO:00991: presynapti	0.03698	TRUE	"Any proce	38	227	3	20972	0.013215859	0.078947368
GO:BP	GO:00488: brain morp	0.03698	TRUE	"The proce	38	227	3	20972	0.013215859	0.078947368
GO:BP	GO:00511: regulation	0.03698	TRUE	"Any proce	38	227	3	20972	0.013215859	0.078947368
GO:BP	GO:19048: regulation	0.037005	TRUE	"Any proce	86	1172	11	20972	0.009385666	0.127906977
GO:BP	GO:00458: negative re	0.037033	TRUE	"Any proce	1275	172	20	20972	0.11627907	0.015686275
GO:BP	GO:00161: vesicle-me	0.037048	TRUE	"A cellular	1572	294	34	20972	0.115646259	0.021628499
GO:BP	GO:00198: stem cell p	0.037089	TRUE	"The proce	180	682	13	20972	0.019061584	0.072222222
GO:BP	GO:00347: positive re	0.03712	TRUE	"Any proce	183	172	6	20972	0.034883721	0.032786885
GO:BP	GO:00070: spindle org	0.037316	TRUE	"A process	210	227	7	20972	0.030837004	0.033333333
GO:BP	GO:00228: regulation	0.037334	TRUE	"Any proce	121	682	10	20972	0.014662757	0.082644628
GO:BP	GO:00352: organ grov	0.03737	TRUE	"The incre	160	682	12	20972	0.017595308	0.075
GO:BP	GO:00320: negative re	0.037493	TRUE	"Any proce	73	227	4	20972	0.017621145	0.054794521
GO:BP	GO:00074: neuroblast	0.037493	TRUE	"The expai	73	227	4	20972	0.017621145	0.054794521

GO:BP	GO:00903!	cellular ser	0.037585	TRUE	"A cell agir	98	1172	12	20972	0.010238908	0.12244898
GO:BP	GO:19001!	negative re	0.037766	TRUE	"Any proce	12	172	2	20972	0.011627907	0.166666667
GO:BP	GO:00463!	negative re	0.037766	TRUE	"Any proce	12	172	2	20972	0.011627907	0.166666667
GO:BP	GO:00068!	sodium ior	0.037766	TRUE	"The direc	245	172	7	20972	0.040697674	0.028571429
GO:BP	GO:00105!	positive re	0.037766	TRUE	"Any proce	12	172	2	20972	0.011627907	0.166666667
GO:BP	GO:00160!	synaptic ve	0.037766	TRUE	"The initial	12	172	2	20972	0.011627907	0.166666667
GO:BP	GO:00217!	hippocam	0.037863	TRUE	"The progr	88	294	5	20972	0.017006803	0.056818182
GO:BP	GO:00615!	actin filam	0.038009	TRUE	"A process	162	294	7	20972	0.023809524	0.043209877
GO:BP	GO:00513!	positive re	0.038078	TRUE	"Any proce	161	227	6	20972	0.026431718	0.037267081
GO:BP	GO:00715!	response t	0.038078	TRUE	"Any proce	264	227	8	20972	0.035242291	0.03030303
GO:BP	GO:00611!	regulation	0.038146	TRUE	"Any proce	200	1172	20	20972	0.017064846	0.1
GO:BP	GO:00508!	positive re	0.038175	TRUE	"Any proce	380	172	9	20972	0.052325581	0.023684211
GO:BP	GO:00900!	regulation	0.038226	TRUE	"Any proce	498	227	12	20972	0.052863436	0.024096386
GO:BP	GO:01060!	regulation	0.038342	TRUE	"Any proce	40	172	3	20972	0.01744186	0.075
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GO:BP	GO:00987!	maintenan	0.038342	TRUE	"Any proce	185	172	6	20972	0.034883721	0.032432432
GO:BP	GO:00219!	hindbrain	0.038388	TRUE	"The radia	10	294	2	20972	0.006802721	0.2
GO:BP	GO:00465!	retinal rod	0.038388	TRUE	"Developm	10	294	2	20972	0.006802721	0.2
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GO:BP	GO:00192!	neuronal a	0.038388	TRUE	"The prop	10	294	2	20972	0.006802721	0.2
GO:BP	GO:00330!	positive re	0.038388	TRUE	"Any proce	10	294	2	20972	0.006802721	0.2
GO:BP	GO:00508!	progester	0.038474	TRUE	"A nuclear	13	227	2	20972	0.008810573	0.153846154
GO:BP	GO:00511!	positive re	0.038474	TRUE	"Any proce	13	227	2	20972	0.008810573	0.153846154
GO:BP	GO:00340!	maintenan	0.038474	TRUE	"The proce	13	227	2	20972	0.008810573	0.153846154
GO:BP	GO:00363!	atrioventri	0.038474	TRUE	"The progr	13	227	2	20972	0.008810573	0.153846154

GO:BP	GO:00340: maintenance of	0.038474	TRUE	"The process of maintaining the structure and function of the organism"	13	227	2	20972	0.008810573	0.153846154
GO:BP	GO:00224: reproductive process	0.038551	TRUE	"A biological process that results in the production of new individuals"	1592	1172	111	20972	0.094709898	0.069723618
GO:BP	GO:19018: negative regulation of	0.038604	TRUE	"Any process that results in the inhibition of a biological process"	75	1172	10	20972	0.008532423	0.133333333
GO:BP	GO:19035: regulation of	0.038604	TRUE	"Any process that results in the regulation of a biological process"	75	1172	10	20972	0.008532423	0.133333333
GO:BP	GO:00601: inner ear development	0.038604	TRUE	"The process of developing the inner ear"	75	1172	10	20972	0.008532423	0.133333333
GO:BP	GO:00331: regulation of	0.038604	TRUE	"Any process that results in the regulation of a biological process"	148	1172	16	20972	0.013651877	0.108108108
GO:BP	GO:00329: positive regulation of	0.038604	TRUE	"Any process that results in the activation of a biological process"	148	1172	16	20972	0.013651877	0.108108108
GO:BP	GO:19028: regulation of	0.038604	TRUE	"Any process that results in the regulation of a biological process"	75	1172	10	20972	0.008532423	0.133333333
GO:BP	GO:00901: tissue migration	0.038678	TRUE	"The process of moving from one location to another"	103	682	9	20972	0.013196481	0.087378641
GO:BP	GO:00610: somite development	0.038678	TRUE	"The process of developing the somites"	85	682	8	20972	0.011730205	0.094117647
GO:BP	GO:00902: negative regulation of	0.038678	TRUE	"Any process that results in the inhibition of a biological process"	103	682	9	20972	0.013196481	0.087378641
GO:BP	GO:00075: heart development	0.038696	TRUE	"The process of developing the heart"	591	682	31	20972	0.045454545	0.052453469
GO:BP	GO:00509: positive regulation of	0.038696	TRUE	"Any process that results in the activation of a biological process"	141	682	11	20972	0.016129032	0.078014184
GO:BP	GO:00424: inner ear development	0.038871	TRUE	"The process of developing the inner ear"	53	1172	8	20972	0.006825939	0.150943396
GO:BP	GO:00516: mitochondrial	0.038871	TRUE	"Any process that results in the regulation of a biological process"	53	1172	8	20972	0.006825939	0.150943396
GO:BP	GO:00020: regulation of	0.03891	TRUE	"Any process that results in the regulation of a biological process"	68	682	7	20972	0.01026393	0.102941176
GO:BP	GO:00075: sex difference	0.03891	TRUE	"The establishment of sex differences"	288	682	18	20972	0.026392962	0.0625
GO:BP	GO:00717: response to	0.03891	TRUE	"Any process that results in the response to a stimulus"	161	682	12	20972	0.017595308	0.074534161
GO:BP	GO:00480: antigen presentation	0.03891	TRUE	"The process of presenting antigens to T cells"	68	682	7	20972	0.01026393	0.102941176
GO:BP	GO:00313: positive regulation of	0.03891	TRUE	"Any process that results in the activation of a biological process"	68	682	7	20972	0.01026393	0.102941176
GO:BP	GO:00420: T cell proliferation	0.038947	TRUE	"The expansion of T cells"	214	1172	21	20972	0.017918089	0.098130841
GO:BP	GO:01200: regulation of	0.038947	TRUE	"Any process that results in the regulation of a biological process"	201	1172	20	20972	0.017064846	0.099502488
GO:BP	GO:00434: regulation of	0.038947	TRUE	"Any process that results in the regulation of a biological process"	64	1172	9	20972	0.007679181	0.140625
GO:BP	GO:00074: peripheral	0.038947	TRUE	"The process of developing the peripheral nervous system"	87	1172	11	20972	0.009385666	0.126436782
GO:BP	GO:00322: regulation of	0.038947	TRUE	"Any process that results in the regulation of a biological process"	111	1172	13	20972	0.01109215	0.117117117
GO:BP	GO:00456: regulation of	0.038947	TRUE	"Any process that results in the regulation of a biological process"	214	1172	21	20972	0.017918089	0.098130841
GO:BP	GO:19026: negative regulation of	0.038977	TRUE	"Any process that results in the inhibition of a biological process"	1284	172	20	20972	0.11627907	0.015576324
GO:BP	GO:00860: cell communication	0.039055	TRUE	"Any process that results in the regulation of a biological process"	57	294	4	20972	0.013605442	0.070175439
GO:BP	GO:00989: regulation of	0.039055	TRUE	"Any process that results in the regulation of a biological process"	30	294	3	20972	0.010204082	0.1
GO:BP	GO:00063: regulation of	0.039055	TRUE	"Any process that results in the regulation of a biological process"	30	294	3	20972	0.010204082	0.1
GO:BP	GO:00514: positive regulation of	0.039055	TRUE	"Any process that results in the activation of a biological process"	30	294	3	20972	0.010204082	0.1
GO:BP	GO:19043: cellular response	0.039055	TRUE	"Any process that results in the regulation of a biological process"	30	294	3	20972	0.010204082	0.1

GO:BP	GO:00432' positive re	0.039114	TRUE	"Any proce	186	172	6	20972	0.034883721	0.032258065
GO:BP	GO:00609' heart form	0.039123	TRUE	"The devel	33	1172	6	20972	0.005119454	0.181818182
GO:BP	GO:00309' regulation	0.039123	TRUE	"Any proce	33	1172	6	20972	0.005119454	0.181818182
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GO:BP	GO:00190' viral gene	0.039434	TRUE	"A process	81	172	4	20972	0.023255814	0.049382716
GO:BP	GO:00033' cardiac mu	0.039434	TRUE	"The enlar	81	172	4	20972	0.023255814	0.049382716
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all	['GO:0033043', 'GO:0097106', 'GO:0099175']
all	['GO:0051094', 'GO:0060996', 'GO:0060998']
score2	['GO:0030154', 'GO:0050793', 'GO:0050794']
score1	['GO:0009987']
all	['GO:0002009', 'GO:0035239']
all	['GO:0032501', 'GO:0048519', 'GO:0051239']
all	['GO:0001654', 'GO:0090596']
score1	['GO:0051649', 'GO:0099003']

score2	['GO:0008150']
all	['GO:0009653', 'GO:0009888']
syndromic	['GO:0021537', 'GO:0048856']
score2	['GO:0016477', 'GO:2000145']
all	['GO:0006887', 'GO:0009987']
score1	['GO:0007049', 'GO:0050794']
score3	['GO:0098916']
score3	['GO:0099537']
score1	['GO:0007611']
score2	['GO:0071705']
all	['GO:0019725', 'GO:0048878']
score2	['GO:0021953', 'GO:0048666']
all	['GO:0015844', 'GO:0015850']
all	['GO:0046903', 'GO:0051046', 'GO:0051050']
all	['GO:0048729', 'GO:0060429']
syndromic	['GO:0016043', 'GO:0048522', 'GO:0051128']
syndromic	['GO:0032502', 'GO:0048519', 'GO:0050793']
all	['GO:0032940', 'GO:0048522', 'GO:0051047', 'GO:1903530']
syndromic	['GO:0007417', 'GO:0030182']
syndromic	['GO:0016358', 'GO:0048856']
score3	['GO:0099536']
all	['GO:0034765', 'GO:0098655']
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all	['GO:0061024', 'GO:0099173']
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score1	['GO:0007165', 'GO:0009966', 'GO:0010647', 'GO:0023056', 'GO:0048584']
score1	['GO:0033036']

score1	['GO:0007417', 'GO:0030182']
all	['GO:0008037', 'GO:0048666']
syndromic	['GO:0045445', 'GO:0045595']
all	['GO:0032880', 'GO:0045184']
syndromic	['GO:0048167']
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score2	['GO:0048878']
all	['GO:0048871']
all	['GO:0010605', 'GO:0019538', 'GO:0051246']
all	['GO:0008016', 'GO:0065008']
score1	['GO:0009653', 'GO:0048513']
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all	['GO:0007409', 'GO:1990138']
syndromic	['GO:0023051', 'GO:0023052', 'GO:0048518']
syndromic	['GO:0000278', 'GO:0051726']
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score2	['GO:0032940', 'GO:0048522', 'GO:0051047', 'GO:1903530']
all	['GO:0141124']
score2	['GO:0040011', 'GO:0050789']
score2	['GO:0055085', 'GO:0098657']
score1	['GO:0021543', 'GO:0048856']
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syndromic	['GO:0042692']
score2	['GO:0001964', 'GO:0032102']
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all	['GO:0050804']
score2	['GO:0044085', 'GO:0050794']
score3	['GO:0008150']
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all	['GO:0043434']
score1	['GO:0007610', 'GO:0009416']

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all	['GO:0022607', 'GO:0044091', 'GO:0061024']
all	['GO:0016043']
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score3	['GO:0007267']
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all	['GO:0009306', 'GO:0051223', 'GO:1903530']
all	['GO:0002027', 'GO:0061337']
score1	['GO:0099173', 'GO:0106027']
all	['GO:0006874']
all	['GO:0032501', 'GO:0042592']
all	['GO:0006814', 'GO:0098655']
all	['GO:0044772', 'GO:0044843']
score1	['GO:0009653', 'GO:0010171', 'GO:0060322']
score1	['GO:0007416', 'GO:0050807', 'GO:1901888']
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score1	['GO:0008150']

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all	['GO:0010975', 'GO:0016358', 'GO:0050793']
score2	['GO:0006915', 'GO:0043067']
score1	['GO:0007416']
syndromic	['GO:0048255']
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all	['GO:0055080', 'GO:0098771']
syndromic	['GO:0060537']
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score2	['GO:0006874']
score2	['GO:0048731']
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all	['GO:0009888']
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all	['GO:0007169', 'GO:0032869']
score3	['GO:0050789', 'GO:0051179']
all	['GO:0090304']
syndromic	['GO:0008150']
score2	['GO:0006351', 'GO:0006355', 'GO:1902680']
syndromic	['GO:0040007', 'GO:0050789']
all	['GO:0048278', 'GO:0140029']
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syndromic	['GO:0048513', 'GO:0072359']
syndromic	['GO:0001558', 'GO:0016049', 'GO:0045927', 'GO:0048522']
all	['GO:0006576', 'GO:0009712']
all	['GO:0018958']
all	['GO:0001508', 'GO:0042391', 'GO:0050789']
score2	['GO:0040012', 'GO:0048870', 'GO:0050794']

syndromic	['GO:0007399', 'GO:0051094', 'GO:0051240', 'GO:0051960']
syndromic	['GO:0051146', 'GO:0051147']
all	['GO:0043010', 'GO:0048856']
all	['GO:0050807', 'GO:0099172']
all	['GO:0043010', 'GO:0048592']
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score1	['GO:0050807', 'GO:0099173']
all	['GO:0007166']
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all	['GO:0030003', 'GO:0055074']
all	['GO:0009894', 'GO:0030163', 'GO:0051246']
score2	['GO:0009653', 'GO:0032502']
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syndromic	['GO:0000288', 'GO:1900151', 'GO:1902373']
score1	['GO:0007165']
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score1	['GO:0006810']
score2	['GO:0051179']
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score1	['GO:0048518', 'GO:0048583', 'GO:0050896']
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score2	['GO:0006836', 'GO:0007268', 'GO:0051649', 'GO:0099643']
score2	['GO:0023061', 'GO:0099536']
all	['GO:0051234']
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score1	['GO:0000278', 'GO:0022402']
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score2	['GO:1901698']
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syndromic	['GO:0006338', 'GO:0071824']
syndromic	['GO:0010564', 'GO:0045023']
syndromic	['GO:0002181', 'GO:0006417']
syndromic	['GO:0009888']
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score1	['GO:0006338', 'GO:0010468']
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syndromic	['GO:0003008', 'GO:0051239']
syndromic	['GO:0048856', 'GO:0060322']
syndromic	['GO:0050793', 'GO:0060996']
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all	['GO:0009628']
score1	['GO:0031175', 'GO:0120035']
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syndromic	['GO:0022402']
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syndromic	['GO:0006937', 'GO:0006941']
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all	['GO:0009987', 'GO:0032501']
score2	['GO:0009987']
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all	['GO:0006873', 'GO:0055080']
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score1	['GO:0009653', 'GO:0009790']
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syndromic	['GO:0022008']
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score2	['GO:0051640']
all	['GO:0001775', 'GO:0002376']
all	['GO:0033043', 'GO:0051276']
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syndromic	['GO:0023061', 'GO:0099536']
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all	['GO:1901698']
score2	['GO:0007268', 'GO:0010647', 'GO:0023056', 'GO:0050804']
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syndromic	['GO:0050896']
syndromic	['GO:0000956', 'GO:0061157']
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all	['GO:0007154', 'GO:0061337']
all	['GO:0007405', 'GO:0050769', 'GO:1902692', 'GO:2000179']
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score1	['GO:0048856', 'GO:0060322']
score1	['GO:0009653', 'GO:0048589']
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all	['GO:0007010', 'GO:0010639', 'GO:0051493']
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syndromic	['GO:0048518', 'GO:0048583', 'GO:0050896']
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all	['GO:0000082', 'GO:1901990', 'GO:1902806']
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all	['GO:0044087', 'GO:0051017', 'GO:0110053']
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score1	['GO:0040007', 'GO:0040008', 'GO:0048518']
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syndromic	['GO:0046649', 'GO:1903131']
all	['GO:0032501']
all	['GO:0022402', 'GO:0051276']
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score2	['GO:0006813', 'GO:0098655']
score1	['GO:0007409', 'GO:1990138']

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all	['GO:0009308']
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score3	['GO:0030154', 'GO:0050793', 'GO:0050794']
score1	['GO:0032879', 'GO:0050794', 'GO:0051641']
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score2	['GO:0007416', 'GO:0022607', 'GO:0099173']
score2	['GO:0048511']
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syndromic	['GO:0042692', 'GO:0045595']
score1	['GO:0008104']
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score1	['GO:0035556']
score1	['GO:0007269', 'GO:0050804', 'GO:0051588', 'GO:1903530']
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syndromic	['GO:0003012', 'GO:0044057']
score3	['GO:0016043']
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score1	['GO:0032501', 'GO:0098727']
all	['GO:0010604', 'GO:0019538', 'GO:0051246']
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score1	['GO:0007585', 'GO:0051239']
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syndromic	['GO:0050896']

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score2	['GO:0007186']
score3	['GO:0042221']
score1	['GO:0009653', 'GO:0032502']
score3	['GO:0006351', 'GO:0006355', 'GO:1902680']
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all	['GO:0050905']
all	['GO:0071869']
all	['GO:0006367', 'GO:0141137']
score3	['GO:0010557', 'GO:0032774', 'GO:0051254', 'GO:2001141']
score2	['GO:0006357', 'GO:0006366', 'GO:0045893']

all	['GO:0022411', 'GO:0043933']
score1	['GO:0001101']
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score2	['GO:0033555']
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syndromic	['GO:0009966', 'GO:0035556']
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all	['GO:0042221']
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all	['GO:0051276', 'GO:0098813']
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score2	['GO:0050804']
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score1	['GO:0042221', 'GO:0051716']
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all	['GO:0043583', 'GO:0048562', 'GO:0090596']
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syndromic	['GO:0009628']
syndromic	['GO:0033993', 'GO:0070887']
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all	['GO:0021696', 'GO:0048646']
all	['GO:0007169', 'GO:0038179']
all	['GO:0031112', 'GO:0031113', 'GO:0032273', 'GO:0046785', 'GO:1902905']
score3	['GO:0030029']
score3	['GO:0065003']
syndromic	['GO:0032535']
all	['GO:0008104', 'GO:0015031', 'GO:0046907']

score1	['GO:0022402', 'GO:0051276']
score1	['GO:0033365', 'GO:1903539']
score1	['GO:0031503', 'GO:0097120', 'GO:0099072', 'GO:0099633']
syndromic	['GO:0016079', 'GO:0017156']
syndromic	['GO:0048167', 'GO:0060292']
all	['GO:0016477', 'GO:0030334', 'GO:2000147']
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score1	['GO:0007626', 'GO:0030534']
all	['GO:0007507', 'GO:0014706']
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score3	['GO:0035556']
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all	['GO:0006468']
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score2	['GO:0007154', 'GO:0050877']
all	['GO:0051146', 'GO:0051147']
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score2	['GO:0006887', 'GO:0009987']

score2	['GO:0120036']
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score2	['GO:0007416', 'GO:0051962', 'GO:0051963', 'GO:1901890']
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all	['GO:0072659', 'GO:1904375', 'GO:1905475']
all	['GO:0007269', 'GO:0046928', 'GO:0050806', 'GO:0051590', 'GO:1903532']
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all	['GO:0007169']
all	['GO:0007159', 'GO:0022407']
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score2	['GO:0010594', 'GO:0035767', 'GO:0050920']
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score2	['GO:0008038', 'GO:0061564', 'GO:0106030']
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score2	['GO:0006810']
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syndromic	['GO:0031667', 'GO:0051716']
syndromic	['GO:0072657', 'GO:1990778']
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all	['GO:0048856', 'GO:0060041']
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all	['GO:0007163']
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all	['GO:0015031', 'GO:0032940', 'GO:0035592', 'GO:0071692']
score2	['GO:0042221']
score2	['GO:0042127', 'GO:0050673']
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syndromic	['GO:0045814']
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score1	['GO:0044770']
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all	['GO:0048167', 'GO:0060291']
all	['GO:0051963', 'GO:0099068', 'GO:0099175']
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score2	['GO:0042391']
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syndromic	['GO:0010629', 'GO:0043488', 'GO:0050779', 'GO:0061014']
syndromic	['GO:0044772', 'GO:0044784']
all	['GO:0051259']
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score1	['GO:0022008']
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syndromic	['GO:0007632', 'GO:0008306']
all	['GO:0033554', 'GO:0050794', 'GO:0080134']
all	['GO:0006810']
syndromic	['GO:0042110', 'GO:0050863', 'GO:0051251', 'GO:1903039']
all	['GO:0030098', 'GO:0051249', 'GO:1902105']
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score2	['GO:0007610', 'GO:0048609']
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score1	['GO:0019538', 'GO:0060255', 'GO:0080090']
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score3	['GO:0016192']
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score1	['GO:0007154', 'GO:0010646', 'GO:0048523']
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syndromic	['GO:0099173', 'GO:0106027']
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syndromic	['GO:0016043', 'GO:0048523', 'GO:0051128']
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syndromic	['GO:0072497', 'GO:2000736']
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syndromic	['GO:0000226', 'GO:0032886', 'GO:0051493']
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score2	['GO:0001935', 'GO:0001936', 'GO:0050679']
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syndromic	['GO:0009132']
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score1	['GO:0001764']
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score1	['GO:0006356', 'GO:0042790']

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score3	['GO:0007269', 'GO:0050804', 'GO:0051588', 'GO:1903530']
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all	['GO:0048856']
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score3	['GO:0006338', 'GO:0071824']
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syndromic	['GO:0006793', 'GO:0019222']
syndromic	['GO:0006195', 'GO:0009150', 'GO:0009261']
all	['GO:0007265', 'GO:0046578', 'GO:0051058']
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score1	['GO:0099587']
syndromic	['GO:0055085', 'GO:0098657']
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score2	['GO:0035556', 'GO:0097190']
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score1	['GO:0007169', 'GO:0032869']
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syndromic	['GO:0006606', 'GO:0033157', 'GO:0046822', 'GO:1900180']
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