

GO term	Description	2-value	FDR q-value	Enrichment (N. B. n. b)	Genes
(20-0000000)	ogulation of primary metabolic process	1.478.4	1200	1.22 (9174,2563,981,333)	+1 Show panes
GD-0090504	nucleic axid metabolic process	1.90-4	9.952-1	1.25 (9174,1997,980,267)	+1 Show pency
00:0051226	direnosens organization	2.88E-4	1200	1.62 (9174,384,1005,68)	+1 Show genes
00:0051547	ogulation of kentinocyte migration	4.32.4	1200	4,587.00 (9174,2,1,1)	+1 Show paner
GD-005[549	positrie regulation of keratinocyte migration	4.32.4	9.02E-1	4,587.00 (9174,2,1,1)	+1 Show person
(30:00)9222	ogulation of metabolic process	4.77E-4	8.34E-1	1.18 (9174,2891,981,366)	+1 Show panes
GD-0031325	regulation of cellular metabolic process	4.778.4	7.15E-1	1.19 (9174,2634,1032,353)	+1 Show pency
00:00(156)	popistic process involved in Interlysis	5.84E-4	7.66E-1	51.54 (9174,2,178,2)	+1 Show gener
CO:1901560	organic cyclic compound metabolic process	7.34E-4	8.56E-1	1.20 (9174,2473,980,317)	+1 Show paner
GD-00W0245	regulation of macroenolecule metabolic process	7.568.4	7.93E-1	1.20 (9174,2436,981,313)	+1 Show ponce
CD-0000002	segative regulation of gone ellencing by RNA	7.63E-4	7.28E-1	1,310.57 (9174,1,7,1)	a) Show gener
CD (050965	to guive regulation of gone silencing by mRNA	7.63E-4	6.67E-1	1,310.57 (9174,1,7,1)	+1 Show panes
(3D-009008)	regulation of heart induction by regulation of canonical Was signaling pathway	7.638.4	6.168-1	1,310.57 (9174,1,7,1)	+1 Show ponce
(20-00001-9)	to guive regulation of posttram-criptional gene silencing	7.63E-4	5.72E-1	1,310.57 (9174,1,7,1)	+1 Show gener
CO-0010468	regulation of gene expression	7.968.4	5.57E-1	1.22 (9174,2016,1036,278)	+1 Show paner
GD-000K139	suckerbase-containing compound metabolic process	8.150-4	5.34E-1	1.21 (9174,2283,980,295)	al Show person
GD-0006259	DNA metabolic process	8.22E-4	5.08E-1	1.58 (9174,394,974,66)	+1 Show panes
GD:1901576	organic substance biosynthetic process	8.52.4	4.96E-1	1.22 (9174,2127,980,277)	+1 Show ponce
00:0071500	cellular empense to exogenous delNA	8.63E-4	4.77E-1	9.74 (9174,6,628,4)	+1 Show gener
CONTRACT CON	seguive regulation of fooding behavior	8.72E-4	4.58E-1	1,146.75 (9174,1,8,1)	+1 Show pomes
CO-CO (CONTO)	segative regulation of macroenolecule metabolic process	8.77E-4	4.39E-1	1.68 (9174,853,352,55)	(+) Show poncy
QD 0034641	collatur airrogem compound metabolic process	92:-4	4.29E-1	1.20 (9174,2497,980,319)	+1 Show genes
00.0045415	se gariou negatation of interleakin-8 biosynthetic process	9.63E-4	4.39E-1	14.36 (9174,3,639,3)	+1 Show panes
00:0019219	regulation of nucleobase-containing compound metabolic process	9.78.4	4.34E-1	1.22 (9174,2054,1032,281)	(+) Show perior
	negalation of cellular macromolecule biosymhetic process	9.82E-4	4.12E-1	1.23 (9174,1817,1032,252)	+1 Show pancy

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