Pathway	Gene ranks	NES	pval	padj
intracellular anatomical structure	Immuram na maran semene e com monero a comunicación de la comunicación de la comunicación de la comunicación de	1.73	8.7e-04	3.5e-02
cellular response to heat	If the contract of the contrac	1.66	5.2e-03	1.4e-01
cellular response to hydrogen peroxide	Here there is a second of the	1.64	1.2e-02	2.9e-01
cell adhesion	l e de la companya d	1.57	2.0e-02	4.0e-01
fungal-type cell wall	10 mm ( ) m m ( )	1.39	3.3e-02	5.5e-01
nucleotide-excision repair		1.47	6.4e-02	8.2e-01
negative regulation of G2/M transition of mitotic cell cycle	The control of the co	1.46	6.6e-02	8.2e-01
SCF ubiquitin ligase complex	The state of the s	1.44	7.7e-02	8.9e-01
peroxisome	1.1	1.37	7.8e-02	8.9e-01
oxidoreductase activity	The first of the second of the	1.44	8.2e-02	9.0e-01
cellular response to pH		1.41	8.3e-02	9.0e-01
zinc ion binding	THE CONTRACTOR OF THE CONTRACT	1.43	8.6e-02	9.2e-01
phagophore assembly site		1.40	9.5e-02	9.8e-01
cellular response to farnesol		1.39	1.0e-01	1.0e+00
protein histidine kinase activity		1.38	1.1e-01	1.0e+00
piecemeal microautophagy of the nucleus	0 101 11 10 10 1	1.37	1.2e-01	1.0e+00
glutathione metabolic process		1.36	1.3e-01	1.0e+00
protein homodimerization activity	The first of the second of the	1.35	1.3e-01	1.0e+00
membrane raft		1.34	1.4e-01	1.0e+00
asexual sporulation resulting in formation of a cellular spore	The first manner of the first of the second	1.23	1.6e-01	1.0e+00
endonucleolytic cleavage in 5'-ETS of tricistronic rRNA transcript (SSU-rRNA. 5.8S rRNA. LSU-rRNA)		-1.76	2.7e-03	8.5e-02
ribosomal small subunit biogenesis		-1.81	2.7e-03	8.5e-02
endonucleolytic cleavage to generate mature 5'-end of SSU-rRNA from (SSU-rRNA. 5.8S rRNA. LSU-rRNA)		-1.76	2.6e-03	8.5e-02
shamixanthone biosynthetic process	term of the second of the sec	-1.82	9.6e-04	3.5e-02
SSU-rRNA from 5.8S rRNA and LSU-rRNA from tricistronic rRNA transcript (SSU-rRNA. 5.8S rRNA. LSU-rRNA)	e e e e e e e e e e e e e e e e e e e	-1.92	9.4e-04	3.5e-02
preribosome. large subunit precursor	e e e e e e e e e e e e e e e e e e e	-1.98	7.1e-04	3.1e-02
helvolic acid biosynthetic process		-1.76	6.5e-04	3.0e-02
cytosolic large ribosomal subunit		-1.91	4.6e-04	2.3e-02
mitochondrial ribosome	Control of the second of the s	-1.92	2.2e-04	1.2e-02
ribosomal large subunit biogenesis	· · · · · · · · · · · · · · · · · · ·	-2.06	1.2e-04	6.9e-03
maturation of SSU-rRNA from tricistronic rRNA transcript (SSU-rRNA. 5.8S rRNA. LSU-rRNA)		-2.10	2.7e-05	1.8e-03
pseurotin A biosynthetic process	1 - 1 - <b>H</b>	-1.88	2.6e-05	1.8e-03
fumitremorgin B biosynthetic process	to the	-1.88	1.6e-05	1.3e-03
rRNA processing	And the second of the second o	-2.17	5.3e-06	4.9e-04
secondary metabolic process	l limitation and a second community of the second community of the second community of the second community of	-2.07	4.4e-06	4.8e-04
prenyltransferase activity	· · · · · · · · · · · · · · · · · · ·	-1.92	3.6e-06	4.7e-04
small-subunit processome		-2.28	1.3e-06	2.1e-04
structural constituent of ribosome	The second of th	-2.25	6.3e-08	1.4e-05
fumagillin biosynthetic process	en e	-2.24	3.2e-09	1.0e-06
nucleolus	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1	-2.34	2.3e-10	1.5e-07
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