Pathway	Gene ranks	NES	pval	padj
TRNA SPLICING	THE COURT OF THE CONTROL OF THE CONT	1.79	1.3e-04	3.1e-02
ALL- <itrans< biosynthesis<="" diphosphate="" i-decaprenyl="" td=""><td>THE COURT OF THE C</td><td>1.71</td><td>1.8e-03</td><td>2.0e-01</td></itrans<>	THE COURT OF THE C	1.71	1.8e-03	2.0e-01
NADH REPAIR		1.58	7.7e-03	2.9e-01
4-HYDROXYPROLINE DEGRADATION I		1.54	9.1e-03	2.9e-01
ANDROGEN BIOSYNTHESIS	THE CONTRACT OF THE CONTRACT O	1.53	2.6e-02	5.5e-01
SPERMINE AND SPERMIDINE DEGRADATION I	THE TELEVISION OF SOME OF STREET	1.48	2.6e-02	5.5e-01
THIAMIN SALVAGE III	The manner manner of the control of	1.44	3.0e-02	5.5e-01
ESTROGEN BIOSYNTHESIS	THE REPORT OF THE PROPERTY OF	1.47	4.2e-02	6.3e-01
THIOSULFATE DISPROPORTIONATION III (RHODANESE)		1.43	4.9e-02	6.3e-01
METHYLMALONYL PATHWAY	THE RELEASE TO THE CONTROL OF THE CO	1.39	5.0e-02	6.3e-01
GLUCONEOGENESIS I	The many transfer of the second of the secon	-1.35	6.3e-02	7.4e-01
RETINOL BIOSYNTHESIS	The state of the s	-1.45	4.7e-02	6.3e-01
GDP-MANNOSE BIOSYNTHESIS		-1.45	4.4e-02	6.3e-01
SERINE BIOSYNTHESIS		-1.38	3.9e-02	6.3e-01
SUPEROXIDE RADICALS DEGRADATION		-1.42	3.0e-02	5.5e-01
PENTOSE PHOSPHATE PATHWAY (OXIDATIVE BRANCH)		-1.52	2.0e-02	5.4e-01
3-PHOSPHOINOSITIDE DEGRADATION		-1.53	9.7e-03	2.9e-01
HISTAMINE BIOSYNTHESIS		-1.49	9.0e-03	2.9e-01
CDP-DIACYLGLYCEROL BIOSYNTHESIS I		-1.68	4.8e-03	2.9e-01
PENTOSE PHOSPHATE PATHWAY (NON-OXIDATIVE BRANCH)		-1.72	2.5e-03	2.0e-01
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