Pathway	Gene ranks	NES	pval	padj
endoplasmic reticulum unfolded protein response	The transfer of the second section of the section of the section of the second section of the	1.63	7.1e-03	1.0e+00
cellular response to unfolded protein	Late the second of the second	1.63	7.1e-03	1.0e+00
regulation of response to external stimulus	1111	1.67	1.4e-02	1.0e+00
cellular response to topologically incorrect protein	The transfer of the second section is	1.57	2.1e-02	1.0e+00
vasculature development		1.58	2.0e-02	1.0e+00
cardiovascular system development		1.58	2.0e-02	1.0e+00
myelin sheath		1.59	1.7e-02	1.0e+00
secretion by cell		1.50	1.3e-02	1.0e+00
secretion		1.49	1.5e-02	1.0e+00
RNA splicing		1.40	1.7e-02	1.0e+00
regulation of cellular protein metabolic process	11.11.11.11.11.11.11.11.11.11.11.11.11.	-1.28	3.3e-02	1.0e+00
chromatin	THE REPORT OF THE PARTY OF THE	-1.43	3.2e-02	1.0e+00
regulation of translation	— III — I	-1.68	7.3e-03	1.0e+00
regulation of cellular amide metabolic process		-1.76	8.1e-03	1.0e+00
nucleolar part		-1.69	9.0e-03	1.0e+00
positive regulation of cellular amide metabolic process	Harmon Contractor of Contractor (Contractor)	-1.53	3.2e-02	1.0e+00
protein-DNA complex	The state of the s	-1.62	3.0e-02	1.0e+00
enzyme activator activity	To the contract of the first	-1.66	2.9e-02	1.0e+00
nuclease activity	The first of the filter	-1.73	2.7e-02	1.0e+00
small GTPase mediated signal transduction		-2.00	7.3e-03	1.0e+00
	0 200 400 600 800			