

Pathway				Gene ranks	NES	pval	padj
MODULE_5					2.23	3.5e-04	1.7e-01
LEI_MYB_TARGETS					1.98	1.6e-03	1.7e-01
CELL_PROJECTION_ORGANIZATION					1.86	2.2e-03	1.7e-01
OF_PROTEIN_METABOLIC_PROCESS					1.94	2.8e-03	1.7e-01
GENESIS_PPARG_RXRA_BOUND_8D					1.80	5.7e-03	1.7e-01
MODULE_60					1.75	8.5e-03	2.1e-01
CELLULAR_ORGANISMAL_PROCESS					1.78	9.0e-03	2.1e-01
PROTEIN_MODIFICATION_PROCESS					1.78	9.0e-03	2.1e-01
ATION_OF_RESPONSE_TO_STRESS					1.78	9.0e-03	2.1e-01
DELYS_THYROID_CANCER_UP					1.77	1.2e-02	2.3e-01
KEGG_OXIDATIVE_PHOSPHORYLATION					-1.80	5.9e-03	1.7e-01
KEGG_HUNTINGTONS_DISEASE					-1.80	5.9e-03	1.7e-01
KEGG_ALZHEIMERS_DISEASE					-1.80	5.9e-03	1.7e-01
RY_ELECTRON_TRANSPORT_CHAIN					-1.80	5.9e-03	1.7e-01
GO_OXIDATIVE_PHOSPHORYLATION					-1.80	5.9e-03	1.7e-01
ATION_OF_ORGANIC_COMPOUNDS					-1.80	5.9e-03	1.7e-01
GO_ELECTRON_TRANSPORT_CHAIN					-1.80	5.9e-03	1.7e-01
GO_CELLULAR_RESPIRATION					-1.80	5.9e-03	1.7e-01
COUPLED_ELECTRON_TRANSPORT					-1.80	5.9e-03	1.7e-01
GO_MEMBRANE_PROTEIN_COMPLEX					-1.88	2.5e-03	1.7e-01
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