component_9_vs_pathway_expression

Component 9	IC(Δ)	P-val	FDR
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_BIOSYNTHESIS	0.434(0.021)	1.11e-02	3.70e-02
KEGG_LYSINE_DEGRADATION	0.383(0.045)	1.11e-02	3.70e-02
KEGG_OXIDATIVE_PHOSPHORYLATION	0.341(0.035)	1.11e-02	3.70e-02
KEGG_PYRIMIDINE_METABOLISM	0.324(0.013)	1.11e-02	3.70e-02
KEGG_PURINE_METABOLISM	0.308(0.038)	1.11e-02	3.70e-02
KEGG_SELENOAMINO_ACID_METABOLISM	0.303(0.021)	1.11e-02	3.70e-02
KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM	0.293(0.013)	1.11e-02	3.70e-02
KEGG_BETA_ALANINE_METABOLISM	0.283(0.034)	1.11e-02	3.70e-02
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION	0.260(0.052)	1.11e-02	3.70e-02
KEGG_CITRATE_CYCLE_TCA_CYCLE	0.230(0.026)	2.22e-02	5.56e-02
KEGG_GLUTATHIONE_METABOLISM	-0.262(0.020)	1.00e+00	0.00e+00
KEGG_GLYCOLYSIS_GLUCONEOGENESIS	-0.276(0.070)	1.00e+00	0.00e+00
KEGG_GALACTOSE_METABOLISM	-0.296(0.033)	1.00e+00	0.00e+00
KEGG_TYROSINE_METABOLISM	-0.341(0.019)	1.00e+00	0.00e+00
KEGG_PHENYLALANINE_METABOLISM	-0.342(0.015)	1.00e+00	0.00e+00
KEGG_PRIMARY_BILE_ACID_BIOSYNTHESIS	-0.391(0.046)	1.00e+00	0.00e+00
KEGG_ASCORBATE_AND_ALDARATE_METABOLISM	-0.402(0.039)	1.00e+00	0.00e+00
KEGG_PENTOSE_AND_GLUCURONATE_INTERCONVERSIONS	-0.423(0.030)	1.00e+00	0.00e+00
KEGG_STARCH_AND_SUCROSE_METABOLISM	-0.452(0.033)	1.00e+00	0.00e+00
KEGG_STEROID_HORMONE_BIOSYNTHESIS	-0.494(0.042)	1.00e+00	0.00e+00