component_9_vs_pathway_expression

Pathway	Component 9	IC(Δ)	P-val	FDR
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_BIOSYNTHESIS		0.434(0.046)	1.11e-02	2.78e-02
KEGG_LYSINE_DEGRADATION		0.383(0.041)	1.11e-02	2.78e-02
KEGG_OXIDATIVE_PHOSPHORYLATION		0.341(0.053)	1.11e-02	2.78e-02
KEGG_PYRIMIDINE_METABOLISM		0.324(0.018)	1.11e-02	2.78e-02
KEGG_PURINE_METABOLISM		0.308(0.015)	1.11e-02	2.78e-02
KEGG_SELENOAMINO_ACID_METABOLISM		0.303(0.019)	1.11e-02	2.78e-02
KEGG_GLYCINE_SERINE_AND_THREONINE_METABOLISM		0.293(0.035)	1.11e-02	2.78e-02
KEGG_BETA_ALANINE_METABOLISM		0.283(0.018)	1.11e-02	2.78e-02
KEGG_VALINE_LEUCINE_AND_ISOLEUCINE_DEGRADATION		0.260(0.012)	1.11e-02	2.78e-02
KEGG_CITRATE_CYCLE_TCA_CYCLE		0.230(0.019)	1.11e-02	2.78e-02
KEGG_GLUTATHIONE_METABOLISM		-0.262(0.033)	1.00e+00	0.00e+00
KEGG_GLYCOLYSIS_GLUCONEOGENESIS		-0.276(0.042)	1.00e+00	0.00e+00
KEGG_GALACTOSE_METABOLISM		-0.296(0.033)	1.00e+00	0.00e+00
KEGG_TYROSINE_METABOLISM		-0.341(0.010)	1.00e+00	0.00e+00
KEGG_PHENYLALANINE_METABOLISM		-0.342(0.028)	1.00e+00	0.00e+00
KEGG_PRIMARY_BILE_ACID_BIOSYNTHESIS		-0.391(0.009)	1.00e+00	0.00e+00
KEGG_ASCORBATE_AND_ALDARATE_METABOLISM		-0.402(0.036)	1.00e+00	0.00e+00
KEGG_PENTOSE_AND_GLUCURONATE_INTERCONVERSIONS		-0.423(0.040)	1.00e+00	0.00e+00
KEGG_STARCH_AND_SUCROSE_METABOLISM		-0.452(0.039)	1.00e+00	0.00e+00
KEGG_STEROID_HORMONE_BIOSYNTHESIS		-0.494(0.055)	1.00e+00	0.00e+00