



Node 1	Node 2	Node 1 annotation	Node 2 annotation	Confidence score
Mdh2	Cs	Malate dehydrogenase 2	Cytrate synthase	0.996
PDH-A1	DLD	Pyruvate dehydrogenase E1	Dihydrolipoyl dehydrogenase, m	0.991
Acadvl	Hadha	VLC-specific acyl-CoA dehydrogenase	Trifunctional enzyme subunit alpha	0.967
Grpel1	HSPA9	GrpE protein homolog 1	Stress-70 protein, mitochondrial	0.964
Acadvl	Cpt2	VLC-specific acyl-CoA dehydrogenase	Carnitine O-palmitoyltransferase 2	0.961
Ivd	Hadha	Isovaleryl-CoA dehydrogenase	Trifunctional enzyme subunit alpha	0.954
Hsd17b10	Hadha	3-hydroxyacyl-CoA dehydrogenase	Trifunctional enzyme subunit alpha	0.931
Mdh2	DLD	Malate dehydrogenase 2	Dihydrolipoyl dehydrogenase	0.89
Hmgcl	Cs	Hydroxymethylglutaryl-CoA lyase	Cytrate synthase	0.877
I79_015301	Hspg2	Nidogen 1.2	Heparan sulfate core protein	0.867
Tufm	HSPA9	Mitochondrial elongation factor Tu	Stress-70 protein, mitochondrial	0.859
Col6a1	Sparc	Collagen alpha-1(VI) chain	Kazal-like domain-containing protein	0.84
Tufm	Mdh2	Mitochondrial elongation factor Tu	Malate dehydrogenase 2	0.828
Shmt2	DLD	Serine hydroxymethyltransferase	Dihydrolipoyl dehydrogenase, m	0.823
Tufm	DLD	Mitochondrial elongation factor Tu	Dihydrolipoyl dehydrogenase, m	0.776
Tufm	Grpel1	Mitochondrial elongation factor Tu	GrpE protein homolog 1	0.768
I79_015301	Sparc	Nidogen 1.2	Kazal-like domain-containing protein	0.76
App	Clu	Amyloid-beta A4 protein	Clusterin	0.736
Hsp90b1	HSPA9	HSP90, beta	Stress-70 protein, mitochondrial	0.734
Mdh2	Hmgcl	Malate dehydrogenase 2	Hydroxymethylglutaryl-CoA lyase	0.711
Cpt2	PDH-A1	Carnitine O-palmitoyltransferase 2	Pyruvate dehydrogenase E1, subunit α	0.703
Shmt2	Tufm	Serine hydroxymethyltransferase	Mitochondrial elongation factor Tu	0.692