



Metabolism:Carbohydrate Metabolism:Glycolysis / Gluconeogenesis;K01810 – glucose-6-phosphate isomerase [EC:5.3.1.9];
 Metabolism:Carbohydrate Metabolism:Glycolysis / Gluconeogenesis;K01834 – 2,3-bisphosphoglycerate-dependent phosphoglycerate mutase [EC:5.4.2.1];
 Metabolism:Carbohydrate Metabolism:Galactose metabolism;K00849 – galactokinase [EC:2.7.1.6];
 Metabolism:Carbohydrate Metabolism:Pyruvate metabolism;K11662 – acetyl-CoA carboxylase / biotin carboxylase [EC:6.4.1.2,6.3.4.14];
 Metabolism:Energy Metabolism:Oxidative phosphorylation;K00329 – NADH dehydrogenase [EC:1.6.5.3];
 Metabolism:Energy Metabolism:Oxidative phosphorylation;K00412 – ubiquinol-cytochrome c reductase cytochrome b subunit;
 Metabolism:Energy Metabolism:Oxidative phosphorylation;K02262 – cytochrome c oxidase subunit 3;
 Metabolism:Energy Metabolism:Oxidative phosphorylation;K02128 – F-type H+ transporting ATPase subunit c [EC:3.6.3.14];
 Metabolism:Lipid Metabolism:Glycerolipid metabolism;K00763 – phospholipid:diacylglycerol acyltransferase [EC:2.3.1.158];
 Metabolism:Amino Acid Metabolism:Leucine degradation;K11433 – histone-lysine N-methyltransferase SETMAR [EC:2.1.1.43];
 Metabolism:Amino Acid Metabolism:Leucine degradation;K09186 – histone-lysine N-methyltransferase ML13 [EC:2.1.1.43];
 Metabolism:Metabolism of Other Amino Acids:Glutathione metabolism;K11204 – glutamate—cysteine ligase catalytic subunit [EC:6.3.1.10];
 Genetic Information Processing:Transcription:Basal transcription factors;K03128 – transcription initiation factor TFIID subunit 2;
 Genetic Information Processing:Transcription:Splicesome;K12827 – RNA-binding protein 25;
 Genetic Information Processing:Transcription:Splicesome;K11984 – U4/U11/U5 snRNP-associated protein 1;
 Genetic Information Processing:Transcription:Splicesome;K12850 – pre-mRNA-splicing factor 3B8;
 Genetic Information Processing:Transcription:Splicesome;K12854 – ATP-dependent RNA helicase DDX23/PRP28 [EC:3.6.4.13];
 Genetic Information Processing:Transcription:Splicesome;K12879 – THO complex subunit 2;
 Genetic Information Processing:Transcription:Splicesome;K12741 – heterogeneous nuclear ribonucleoprotein A1/A3;
 Genetic Information Processing:Translation:Ribosome;K02976 – small subunit ribosomal protein S26;
 Genetic Information Processing:Translation:Ribosome;K02930 – large subunit ribosomal protein L4e;
 Genetic Information Processing:Translation:Ribosome;K02936 – large subunit ribosomal protein L7a/e;
 Genetic Information Processing:Translation:Ribosome;K02940 – large subunit ribosomal protein L9e;
 Genetic Information Processing:Translation:Ribosome;K02877 – large subunit ribosomal protein L15e;
 Genetic Information Processing:Translation:Ribosome;K02891 – large subunit ribosomal protein L22e;
 Genetic Information Processing:Translation:RNA transport;K02524 – translation initiation factor 3 subunit A;
 Genetic Information Processing:Folding, Sorting and Degradation:Ubiquitin mediated proteolysis;K0350 – anaphase-promoting complex subunit 3;
 Genetic Information Processing:Folding, Sorting and Degradation:RNA degradation;K12580 – CCR4-NOT transcription complex subunit 3;
 Environmental Information Processing:Membrane Transport:ABC transporters;K06558 – ATP-binding cassette, subfamily B (MDR/TAP), member 1;
 Cellular Processes:Transport and Catabolism:Endocytosis;K12188 – ESCRT-II complex subunit VPS52;
 Cellular Processes:Cell Growth and Death:Cell cycle;K06628 – cell division control protein 45;
 Cellular Processes:Cell Communication:Tight junction;K10352 – myosin heavy chain;

Comparison

KOs