

A ▼ 09100 Metabolism

B ▼ 09101 Carbohydrate metabolism

C ▼ 00010 Glycolysis / Gluconeogenesis [PATH:ko00010]

D K00844 HK; hexokinase [EC:2.7.1.1]

K12407 GCK; glucokinase [EC:2.7.1.2]

K00845 glk; glucokinase [EC:2.7.1.2]

K25026 glk; glucokinase [EC:2.7.1.2]

K01810 GPI, pgi; glucose-6-phosphate isomerase [EC:5.3.1.9]

K06859 pgil; glucose-6-phosphate isomerase, archaeal [EC:5.3.1.9]

K13810 tal-pgi; transaldolase / glucose-6-phosphate isomerase [EC:2.2.1.2 5.3.1.9]

K15916 pgi-pmi; glucose/mannose-6-phosphate isomerase [EC:5.3.1.9 5.3.1.8]

K24182 PFK9; 6-phosphofructokinase [EC:2.7.1.11]

K00850 pfkA, PFK; 6-phosphofructokinase 1 [EC:2.7.1.11]

K16370 pfkB; 6-phosphofructokinase 2 [EC:2.7.1.11]

K21071 pfk, pfp; ATP-dependent phosphofructokinase / diphosphate-dependent phosphofructokinase [EC:2.7.1.11 2.7.1.90]

K00918 pfkC; ADP-dependent phosphofructokinase/glucokinase [EC:2.7.1.146 2.7.1.147]

K00895 pfp, PFP; diphosphate-dependent phosphofructokinase [EC:2.7.1.90]

K03841 FBP, fbp; fructose-1,6-bisphosphatase I [EC:3.1.3.11]

K02446 glpX; fructose-1,6-bisphosphatase II [EC:3.1.3.11]

K11532 glpX-SEBP; fructose-1,6-bisphosphatase II / sedoheptulose-1,7-bisphosphatase [EC:3.1.3.11 3.1.3.37]

K01086 fbp-SEBP; fructose-1,6-bisphosphatase I / sedoheptulose-1,7-bisphosphatase [EC:3.1.3.11 3.1.3.37]

K04041 fbp3; fructose-1,6-bisphosphatase III [EC:3.1.3.11]

K01623 ALDO; fructose-bisphosphate aldolase, class I [EC:4.1.2.13]

K11645 fbaB; fructose-bisphosphate aldolase, class I [EC:4.1.2.13]

K01624 FBA, fbaA; fructose-bisphosphate aldolase, class II [EC:4.1.2.13]

K01622 K1622; fructose 1,6-bisphosphate aldolase/phosphatase [EC:4.1.2.13 3.1.3.11]

K16305 K16305; fructose-bisphosphate aldolase / 6-deoxy-5-ketofructose 1-phosphate synthase [EC:4.1.2.13 2.2.1.11]

K16306 K16306; fructose-bisphosphate aldolase / 2-amino-3,7-dideoxy-D-threo-hept-6-ulose synthase [EC:4.1.2.13 2.2.1.10]

K01803 TPI, tpiA; triosephosphate isomerase (TIM) [EC:5.3.1.1]

K00134 GAPDH, gapA; glyceraldehyde 3-phosphate dehydrogenase [EC:1.2.1.12]

K10705 GAPDHS; glyceraldehyde-3-phosphate dehydrogenase, spermatogenic [EC:1.2.1.12]

K00150 gap2; glyceraldehyde-3-phosphate dehydrogenase (NAD(P)) [EC:1.2.1.59]

K00927 PGK, pgk; phosphoglycerate kinase [EC:2.7.2.3]