		- K01623 ALO; fructose-bisphosphate aldolase, class I
		- K01623 ALO; fructose-dispriospriate aldolase, class i - K00850 pfkA, PFK; 6-phosphofructokinase 1
		K00012 UGH, ugd; UPglucose 6-dehydrogenase
		K01052 LIPA; lysosomal acid lipase/cholesteryl ester hydrolase  - K03542 psbS; photosystem II 22ka protein
		- K08912 LHCB1; light-harvesting complex II chlorophyll a/b binding protein 1
		- K00760 hprT, hpt, HPRT1; hypoxanthine phosphoribosyltransferase
		- K12319 GUCY1B; guanylate cyclase soluble subunit beta - K00872 thrB; homoserine kinase
		- K12251 aguB; N-carbamoylputrescine amidase
		- K24138 prx3; glutaredoxin/glutathione-dependent peroxiredoxin
		- K00703 glgA; starch synthase
		K01793 GLCE; heparosan-N-sulfate-glucuronate 5-epimerase  - K00995 pgsA, PGS1; CP-diacylglycerolglycerol-3-phosphate 3-phosphatidyltransferase
		- K13788 pta; phosphate acetyltransferase
		K01602 rbcS, cbbS; ribulose-bisphosphate carboxylase small chain
		- K00814 GPT, ALT; alanine transaminase
		- K01599 hemE, URO; uroporphyrinogen decarboxylase - K15747 LUT5, CYP97A3; beta-ring hydroxylase
		- K05601 hcp; hydroxylamine reductase
		K01870 IARS, ileS; isoleucyl-tRNA synthetase
		- K00898 PK2_3_4; pyruvate dehydrogenase kinase 2/3/4
		K09602 OTUB1; ubiquitin thioesterase protein OTUB1  - K17609 NXN; nucleoredoxin
		- K13216 PPP1R8, NIPP1; nuclear inhibitor of protein phosphatase 1
		- K16075 MRS2, MFM1; magnesium transporter
		- K15987 hppA; K(+)-stimulated pyrophosphate-energized sodium pump - K24175 MFS5; MFS transporter, MFS domain-containing protein family, molybdate-anion tra
		- K24175 MFS5; MFS transporter, MFS domain-containing protein family, molybdate-anion tra - K03442 mscS; small conductance mechanosensitive channel
		K14347 SLC10A7, P7; solute carrier family 10 (sodium/bile acid cotransporter), member 7
		- K05643 ABCA3; ATP-binding cassette, subfamily A (ABC1), member 3
		K09284 AP2; AP2-like factor, euAP2 lineage - K03097 CSNK2A; casein kinase II subunit alpha
		- K14810 X56, BP9; ATP-dependent RNA helicase X56/BP9
		- K22857 EEF1AKMT4; EEF1A lysine methyltransferase 4
		- K04382 PPP2C; serine/threonine-protein phosphatase 2A catalytic subunit
		- K04077 groEL, HSP1; chaperonin GroEL - K17783 ERV1, GFER, ALR; mitochondrial FA-linked sulfhydryl oxidase
		K17783 CIX12) OI EI() MEI() Interest and IV Interest Summy at your States of
		- K12890 SFRS1, ASF, SF2; splicing factor, arginine/serine-rich 1
		- K12896 SFRS7; splicing factor, arginine/serine-rich 7
		- K12736 PPW1; peptidylprolyl isomerase domain and W repeat-containing protein 1 - K03066 PSMC5, RPT6; 26S proteasome regulatory subunit T6
		K03676 grxC, GLRX2; glutaredoxin 3
		- K09529 NAJC9; naJ homolog subfamily C member 9
		- K22803 SMC5; structural maintenance of chromosomes protein 5 - K04459 USP, MKP; dual specificity MAP kinase phosphatase
		- K04371 ERK, MAPK1_3; mitogen-activated protein kinase 1/3
		- K14498 SNRK2; serine/threonine-protein kinase SRK2
		- K07950 ARL5B; AP-ribosylation factor-like protein 5B
		- K03544 clpX, CLPX; ATP-dependent Clp protease ATP-binding subunit ClpX - K08770 UBC; ubiquitin C
		- K16280 RGLG; E3 ubiquitin-protein ligase RGLG
		- K10474 KBTB6_7; kelch repeat and BTB domain-containing protein 6/7
		- K10481 BTB9; BTB/POZ domain-containing protein 9
		- K08511 ATVAMP72; vesicle-associated membrane protein 72 - K07874 RAB1A; Ras-related protein Rab-1A
		- K09503 NAJA2; naJ homolog subfamily A member 2
		- K12486 SMAP; stromal membrane-associated protein
		- K07375 TUBB; tubulin beta
		- K03386 PRX2_4, ahpC; peroxiredoxin 2/4 - K10523 SPOP; speckle-type POZ protein
		- K01322 PREP; prolyl oligopeptidase
		- K10355 ACTF; actin, other eukaryote
		- K10408 NAH; dynein heavy chain, axonemal - K11252 H2B; histone H2B
		- K11232 HZB, HIStorie HZB - K03453 TC.BASS; bile acid:Na+ symporter, BASS family
		- K23518 MACRO, ymdB; O-acetyl-AP-ribose deacetylase
		- K17893 AOX1, AOX2; ubiquinol oxidase
		- K20416 FA5; palmitoyl-[glycerolipid] 7-desaturase - K07034 K07034; uncharacterized protein
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