

[Purine metabolism; n = 16]

K04765 mazG; nucleoside triphosphate diphosphatase [EC:3.6.1.9]

| triphosphate,3'-diphosphate pyrophosphatase [EC:3.6.1.11 3.6.1.40] |

K00942 gmk; guanylate kinase [EC:2.7.4.8]

midophosphoribosyltransferase [EC:2.4.2.14]

K01756 purB; adenylosuccinate lyase [EC:4.3.2.2]

imidazole-succinocarboxamide synthase [EC:6.3.2.6]

K00945 cmk; CMP/dCMP kinase [EC:2.7.4.25]

xanthine phosphoribosyltransferase [EC:2.4.2.8]

K00769 gpt; xanthine phosphoribosyltransferase [EC:2.4.2.22]

K00769 gpt; xanthine phosphoribosyltransferase [EC:2.4.2.22]

0759 APRT; adenine phosphoribosyltransferase [EC:2.4.2.7]

K01939 purA; adenylosuccinate synthase [EC:6.3.4.4]

K00088 IMPDH; IMP dehydrogenase [EC:1.1.1.205]

SAR116_0223 mazg protein

SAR116_1993 Ppx/GppA pho

SAR116_0636 Guanylate kinase

SAR116_0651 purF; Glutamine p

SAR116_0667 adenylosuccinate lyase

SAR116_0668 phosphoribosylaminoimi

SAR116_1251 Cytidylate kinase

SAR116_1426 HGPR

SAR116_1505 Xanthine

SAR116_1505 Xgpt

SAR116_1601 Aden

SAR116_1668 Adenine phosphori

SAR116_2308 adenylosuccinate synthase

SAR116_2483 inosine-5'-monophosphate dehydrog

