

feature_Taxon

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|--|------|------|------|------|------|------|------|------|------|
| K03783__"punA, PNP; purine–nucleoside phosphorylase [EC:2.4.2.1]" | 1231 | 1022 | 1226 | 541 | 808 | 304 | 139 | 8 | 2258 |
| K02879__"RP–L17, MRPL17, rplQ; large subunit ribosomal protein L17" | 151 | 2863 | 1183 | 75 | 411 | 836 | 1030 | 18 | 388 |
| K03309__"TC.DAACS; dicarboxylate/amino acid:cation (Na+ or H+) symporter, DAACS family" | 242 | 611 | 1213 | 2890 | 543 | 10 | 475 | 572 | 80 |
| K01923__purC; phosphoribosylaminoimidazole–succinocarboxamide synthase [EC:6.3.2.6]" | 2555 | 141 | 842 | 23 | 4 | 642 | 546 | 310 | 995 |
| K07097__K07097; uncharacterized protein | 913 | 1013 | 117 | 28 | 129 | 2868 | 415 | 30 | 350 |
| K13925__pfbA; plasmin and fibronectin–binding protein A | 852 | 947 | 19 | 717 | 58 | 2819 | 97 | 291 | 5 |
| K09818__ABC.MN.S; manganese/iron transport system substrate–binding protein | 5 | 310 | 1200 | 243 | 215 | 923 | 13 | 9 | 2876 |
| K00134__"GAPDH, gapA; glyceraldehyde 3–phosphate dehydrogenase [EC:1.2.1.12]" | 331 | 75 | 24 | 607 | 119 | 556 | 384 | 961 | 2544 |
| K04794__"PTH2; peptidyl–tRNA hydrolase, PTH2 family [EC:3.1.1.29]" | 3 | 942 | 66 | 2879 | 980 | 335 | 37 | 174 | 55 |
| K07052__K07052; uncharacterized protein | 1166 | 853 | 1182 | 452 | 705 | 813 | 56 | 138 | 20 |
| K02226__"cobC, phpB; alpha–ribazole phosphatase [EC:3.1.3.73]" | 120 | 25 | 211 | 13 | 1237 | 50 | 4 | 2955 | 548 |
| K07448__mrr; restriction system protein | 2896 | 112 | 14 | 375 | 169 | 518 | 87 | 691 | 292 |
| K09748__rimP; ribosome maturation factor RimP | 484 | 401 | 171 | 12 | 1261 | 20 | 79 | 754 | 1937 |
| K18427__hpdB; 4–hydroxyphenylacetate decarboxylase large subunit [EC:4.1.1.83]" | 63 | 572 | 885 | 134 | 55 | 2835 | 15 | 143 | 97 |
| K02013__ABC.FEV.A; iron complex transport system ATP–binding protein [EC:3.6.3.34]" | 1357 | 734 | 192 | 351 | 1936 | 68 | 26 | 52 | 46 |
| K05988__dexA; dextranase [EC:3.2.1.11]" | 2 | 137 | 1857 | 642 | 872 | 21 | 439 | 64 | 724 |
| K07705__"lytT, lytR; two–component system, LytTR family, response regulator LytT" | 685 | 35 | 2107 | 500 | 1299 | 5 | 68 | 23 | 8 |
| K07344__trbL; type IV secretion system protein TrbL | 1112 | 45 | 89 | 9 | 422 | 2305 | 76 | 190 | 444 |
| K18122__"cat2, abfT; 4–hydroxybutyrate CoA–transferase [EC:2.8.3.–]" | 159 | 163 | 30 | 170 | 743 | 415 | 2293 | 61 | 294 |
| K13016__wbpB; UDP–N–acetyl–2–amino–2–deoxyglucuronate dehydrogenase [EC:1.1.1.335]" | 11 | 467 | 3 | 73 | 2806 | 228 | 20 | 178 | 123 |
| K10678__nfsA; nitroreductase [EC:1.–.–.–]" | 89 | 21 | 91 | 121 | 2800 | 6 | 31 | 501 | 92 |
| K01847__MUT; methylmalonyl–CoA mutase [EC:5.4.99.2]" | 282 | 30 | 26 | 151 | 56 | 849 | 217 | 317 | 1750 |
| K07032__K07032; uncharacterized protein | 16 | 347 | 6 | 123 | 2688 | 159 | 3 | 302 | 23 |
| K00175__"korB, oorB, oforB; 2–oxoglutarate/2–oxoacid ferredoxin oxidoreductase subunit beta [EC:1.2.7.3 1.2.7.11]" | 219 | 132 | 59 | 17 | 211 | 151 | 133 | 2247 | 442 |
| K07091__lptF; lipopolysaccharide export system permease protein | 603 | 312 | 18 | 534 | 293 | 552 | 602 | 429 | 210 |
| K02119__"ATPVC, ntpC, atpC; V/A–type H+/Na+–transporting ATPase subunit C" | 732 | 508 | 321 | 91 | 916 | 410 | 92 | 29 | 427 |
| K13829__aroKB; shikimate kinase / 3–dehydroquate synthase [EC:2.7.1.71 4.2.3.4]" | 2256 | 156 | 131 | 86 | 27 | 11 | 12 | 60 | 2 |
| K07704__"lytS; two–component system, LytTR family, sensor histidine kinase LytS [EC:2.7.13.3]" | 958 | 204 | 559 | 11 | 142 | 3 | 325 | 13 | 19 |
| K17948__nanM; N–acetylneuraminate epimerase [EC:5.1.3.24]" | 244 | 338 | 129 | 195 | 377 | 221 | 351 | 19 | 310 |
| K00785__"lst; beta–galactosamide–alpha–2,3–sialyltransferase [EC:2.4.99.–]" | 80 | 97 | 28 | 208 | 74 | 56 | 126 | 421 | 105 |
| K12240__pchF; pyochelin synthetase | 6 | 4 | 9 | 90 | 12 | 137 | 2 | 3 | 13 |
| K12998__rgpE; glucosyltransferase [EC:2.4.1.–]" | 1 | 1 | 2 | 1 | 40 | 1 | 1 | 2 | 1 |

