

feature_Taxon

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

