

lysine/arginine/ornithine-binding periplasmic protein (DDPKDNGI_00298)

4-methylaminobutanoate oxidase (formaldehyde-forming) (DDPKDNGI_00302)

Glycine cleavage system transcriptional activator (DDPKDNGI_00301)

1'-deoxyxylulose-5-phosphate synthase YajO (DDPKDNGI_00299)

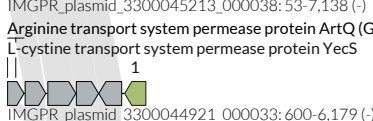
Formyltetrahydrofolate deformylase (DDPKDNGI_00300)



IMGPR_plasmid_3300045213_000038: 53-7,138 (-)

Arginine transport system permease protein ArtQ (GJKLKDGL_00006)

L-cystine transport system permease protein YecS



tet_MFS_A_B_C_D-NCBIFAM Tet(A)/Tet(B)/Tet(C) family tetracycline efflux MFS transporter (NF012174.0) (JMCOIAOA_00311)

TRAP-T-associated universal stress protein TeaD (JMCOIAOA_00308)

putative 3-hydroxybutyryl-CoA dehydrogenase (JMCOIAOA_00305)

Transcriptional regulatory protein ros (JMCOIAOA_00312)

Electron transfer flavoprotein subunit alpha (JMCOIAOA_00304)

Electron transfer flavoprotein subunit beta (JMCOIAOA_00303)

O-acetylserine sulfhydrylase (JMCOIAOA_00307)

L-methionine gamma-lyase (JMCOIAOA_00309)

Protein translocase subunit SecA (JMCOIAOA_00318)

Glutaredoxin 3 (JMCOIAOA_00326)

Aspartate kinase (JMCOIAOA_00331)

T6SS-II' (VF0783) (JMCOIAOA_00336)

IMGPR_plasmid_3300045096_000061: 335,135-381,499 (+)

MFS_efflux_tcmA-NCBIFAM tetracenomycin C efflux MFS transporter (NF000025.1) (OALOJIOK_00309)

Flagellar motor switch protein FlIN (OALOJIOK_00316)

Flagellar motor switch protein FlIG (OALOJIOK_00315)

putative ATP synthase YscN (OALOJIOK_00322)

Flagellar L-ring protein (OALOJIOK_00331)

Flagellar P-ring protein (OALOJIOK_00329)

Flagellar M-ring protein (OALOJIOK_00335)

Flagella (VF0685) (OALOJIOK_00325)

Flagella (VF0373) (OALOJIOK_00321)

Flagella (VF0373) (OALOJIOK_00327)

RegX3 (VF0858) (OALOJIOK_00341)

Flagella (VF0373) (OALOJIOK_00342)

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