

VanR-ABDEGL family response regulator transcription factor (NF033117.2) (NHODGFGJ_00013)

ANT(9) family aminoglycoside nucleotidyltransferase (NF012212.1) (NHODGFGJ_00008)

Inositol 2-dehydrogenase/D-chiro-inositol 3-dehydrogenase (NHODGFGJ_00007)

Potassium-transporting ATPase ATP-binding subunit (NHODGFGJ_00016)

Potassium-transporting ATPase KdpC subunit (NHODGFGJ_00015)

HitABC (VF0268) (NHODGFGJ_00006)

Sensor protein KdpD (NHODGFGJ_00014)

Gabija_GajB_2 (NHODGFGJ_00010)

NHODGFGJ_00011

NHODGFGJ_00012

NHODGFGJ_00009

Potassium-transporting ATPase potassium-binding subunit (NHODGFGJ_00017)



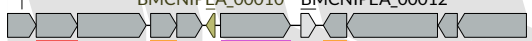
IMGPR_plasmid_2921263974_000022 6234:22465(+)



IMGPR_plasmid_2916055098_000018 6234:22465(+)

L-arabinose transport system permease protein AraQ (BMCNIEPA_00005)

BMCNIEPA_00010 BMCNIEPA_00012



IMGPR_plasmid_2924207177_000014 5386:20745(+)

High-affinity branched-chain amino acid transport system permease protein LivH (GLGILKLF_00031)

High-affinity branched-chain amino acid transport system permease protein LivH (GLGILKLF_00030)

putative fructoselysine utilization operon transcriptional repressor (GLGILKLF_00010)

Leucine-, isoleucine-, valine-, threonine-, and alanine-binding protein (GLGILKLF_00012)

N-formyl-4-amino-5-aminomethyl-2-methylpyrimidine deformylase (GLGILKLF_00019)

F420-dependent glucose-6-phosphate dehydrogenase (GLGILKLF_00020)

NAD(P) transhydrogenase subunit alpha part 1 (GLGILKLF_00018)

Bifunctional F420 biosynthesis protein FbiB (GLGILKLF_00026)

Phosphoenolpyruvate guanylyltransferase (GLGILKLF_00024)

NAD(P) transhydrogenase subunit alpha (GLGILKLF_00017)

Glutaconate CoA-transferase subunit B (GLGILKLF_00021)

Glutaconate CoA-transferase subunit A (GLGILKLF_00022)

NAD(P) transhydrogenase subunit beta (GLGILKLF_00016)

ABC-type heme transporter (VF0739) (GLGILKLF_00029)

Diaminopropionate ammonia-lyase (GLGILKLF_00011)

Phosphoenolpyruvate transferase (GLGILKLF_00025)

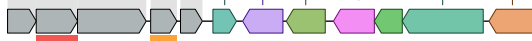
Caffeine dehydrogenase subunit alpha (GLGILKLF_00015)

FbpABC (VF0272) (GLGILKLF_00028)

13 14

FO synthase (GLGILKLF_00023) 27

Dihydroorotase (GLGILKLF_00032)



IMGPR_plasmid_2921208531_000014 5747:47838(+)

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Defence Virulence AMR