

protein-methionine-sulfoxide reductase catalytic subunit MsrP (LOPJJDKA_00029)

Nitrilotriacetate monooxygenase component A (LOPJJDKA_00028)

HTH-type transcriptional regulator DmIR (LOPJJDKA_00031)

SoFic_SoFic (LOPJJDKA_00030)

Bifunctional hemolysin/adenylate cyclase (LOPJJDKA_00034)

S-(hydroxymethyl)glutathione dehydrogenase (LOPJJDKA_00035)

IMGPR_plasmid_2921282425_000036: 26,269-36,140 (+)

Bifunctional hemolysin/adenylate cyclase (GKKCOEDK_00002)

S-(hydroxymethyl)glutathione dehydrogenase (GKKCOEDK_00001)

IMGPR_plasmid_2806310410_000017: 17-9,579 (-)

Bifunctional hemolysin/adenylate cyclase (FJOHEBEO_00003)

IMGPR_plasmid_2551306090_000046: 92-9,981 (-)

Bifunctional hemolysin/adenylate cyclase (FLFCNILM_00030)

Putative ring-cleaving dioxygenase MhqQ (FLFCNILM_00037)

HTH-type transcriptional regulator DmIR (FLFCNILM_00038)

N-substituted formamide deformylase (FLFCNILM_00034)

Enterobactin exporter EntS (FLFCNILM_00032)

putative hydrolase YcaC (FLFCNILM_00036)

IMGPR_plasmid_2551306084_000040: 24,502-42,083 (+)

Bifunctional hemolysin/adenylate cyclase (OIIEKEDC_00010)

Putative ring-cleaving dioxygenase MhqQ (OIIEKEDC_00003)

HTH-type transcriptional regulator DmIR (OIIEKEDC_00002)

N-substituted formamide deformylase (OIIEKEDC_00006)

Enterobactin exporter EntS (OIIEKEDC_00008)

putative hydrolase YcaC (OIIEKEDC_00004)

IMGPR_plasmid_2513237086_000044: 330-17,911 (-)

Bifunctional hemolysin/adenylate cyclase (HKIGGFLE_00010)

Putative ring-cleaving dioxygenase MhqQ (HKIGGFLE_00003)

HTH-type transcriptional regulator DmIR (HKIGGFLE_00002)

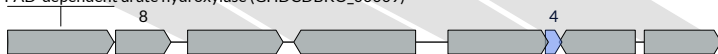
N-substituted formamide deformylase (HKIGGFLE_00006)

Enterobactin exporter EntS (HKIGGFLE_00008)

putative hydrolase YcaC (HKIGGFLE_00004)

IMGPR_plasmid_2551306085_000045: 105-17,686 (-)

FAD-dependent urate hydroxylase (GHDCDBKO_00009)



IMGPR_plasmid_2630968934_000030: 548-8,929 (-)

7000 nt

defence (DefenseFinder)