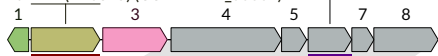


Michele_2A_hits_grouped (CCDHKIHN_00006)

FadE29 (VF0845) (CCDHKIHN_00002)



IMGPR_plasmid_3300022693_000326: 41-7,710 (+)



IMGPR_plasmid_3300026098_000059: 49-5,734 (+)



IMGPR_plasmid_3300025216_000061: 49,587-53,204 (-)

Succinyl-CoA--L-malate CoA-transferase beta subunit (MNGJGLMD_00072)

Glutaconate CoA-transferase subunit B (MNGJGLMD_00063)

Trans-feruloyl-CoA synthase FCS1 (MNGJGLMD_00070)

HTH-type transcriptional regulator KipR (MNGJGLMD_00056)

Trans-feruloyl-CoA synthase FCS1 (MNGJGLMD_00071)

FadE29 (VF0845) (MNGJGLMD_00057)

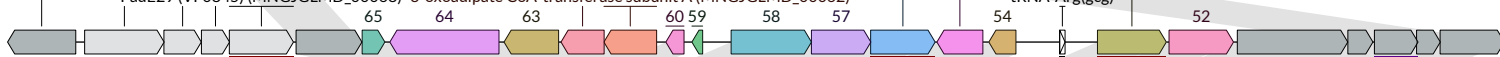
Bifunctional protein PaaZ (MNGJGLMD_00069)

FadE29 (VF0845) (MNGJGLMD_00053)

FadE29 (VF0845) (MNGJGLMD_00068)

3-oxoadipate CoA-transferase subunit A (MNGJGLMD_00062)

tRNA-Arg(gcg)



IMGPR_plasmid_3300009578_000113: 44,194-70,855 (-)

HTH-type transcriptional regulator KipR (HGACHAIA_00010)

Crotonobetainyl-CoA reductase (HGACHAIA_00008)

Glutaconate CoA-transferase subunit B (HGACHAIA_00003)

FadE28 (VF0844) (HGACHAIA_00012)

FadE29 (VF0845) (HGACHAIA_00009)

3-oxoadipate CoA-transferase subunit A (HGACHAIA_00004)

FadE29 (VF0845) (HGACHAIA_00013)

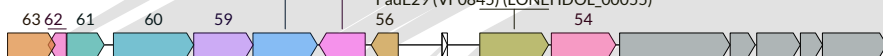


IMGPR_plasmid_2788500509_000016: 24-19,871 (+)

HTH-type transcriptional regulator KipR (LONEHDOL_00058)

FadE29 (VF0845) (LONEHDOL_00059)

FadE29 (VF0845) (LONEHDOL_00055)

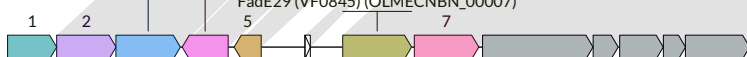


IMGPR_plasmid_3300017702_000072: 44,867-60,526 (-)

HTH-type transcriptional regulator KipR (OLMECNBN_00004)

FadE29 (VF0845) (OLMECNBN_00003)

FadE29 (VF0845) (OLMECNBN_00007)



IMGPR_plasmid_3300025180_000037: 41-13,259 (+)

10000 nt

defence (DefenseFinder) **virulence (VFDB)** **tRNA**