

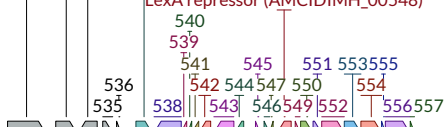
ono(2-hydroxyethyl) terephthalate hydrolase (AMCIDIMH_00532)

Diacetyl reductase [(S)-acetoin forming] (AMCIDIMH_00534)

Right origin-binding protein (AMCIDIMH_00533)

OmpD (VF0969) (AMCIDIMH_00537)

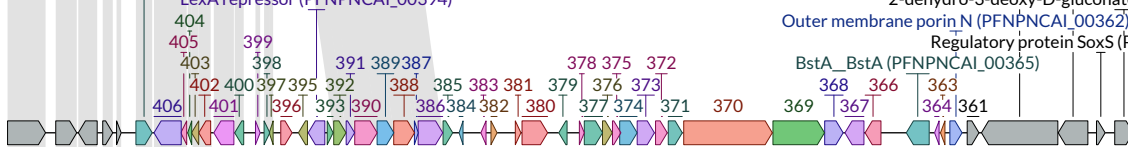
LexA repressor (AMCIDIMH_00548)



IMGPR_plasmid_2919150387_000002 566470:583273(+)

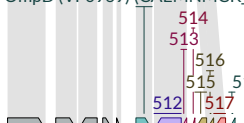
OmpD (VF0969) (PFNPNCAL_00407)

LexA repressor (PFNPNCAL_00394)



IMGPR_plasmid_2927767760_000004 393614:440083(-)

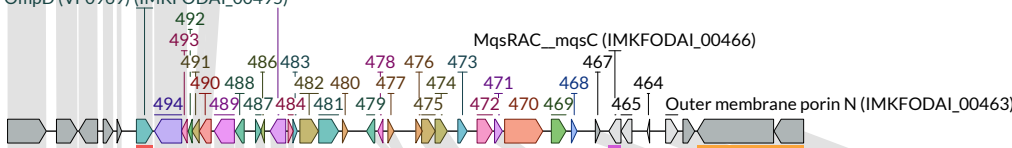
OmpD (VF0969) (CAEMNMCK_00511)



IMGPR_plasmid_2839371029_000001 532609:541999(+)

putative HTH-type transcriptional regulator (IMKFODAI_00485)

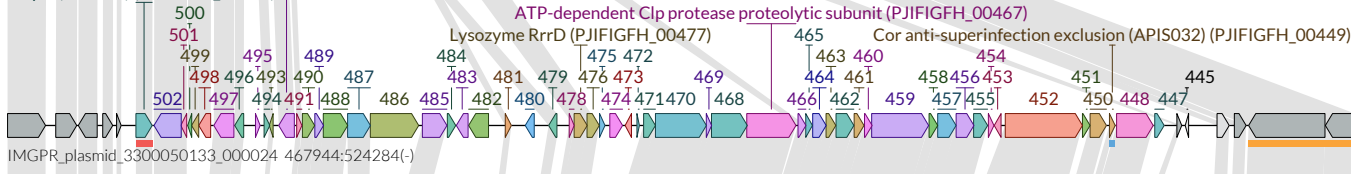
OmpD (VF0969) (IMKFODAI_00495)



IMGPR_plasmid_2895090955_000003 478931:511795(-)

putative HTH-type transcriptional regulator (PJIFIGFH_00492)

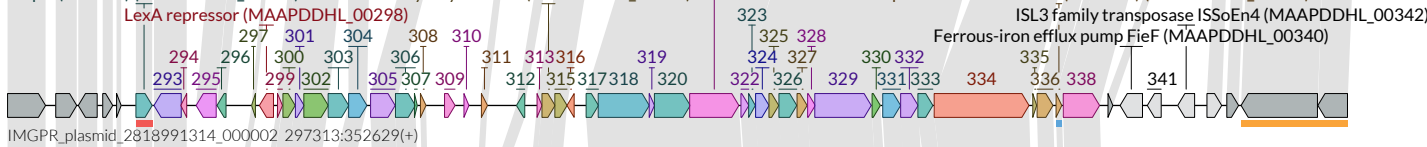
OmpD (VF0969) (PJIFIGFH_00503)



IMGPR_plasmid_3300050133_000024 467944:524284(-)

ATP-dependent Clp protease proteolytic subunit (MAAPDDHL_00321)

OmpD (VF0969) (MAAPDDHL_00292)

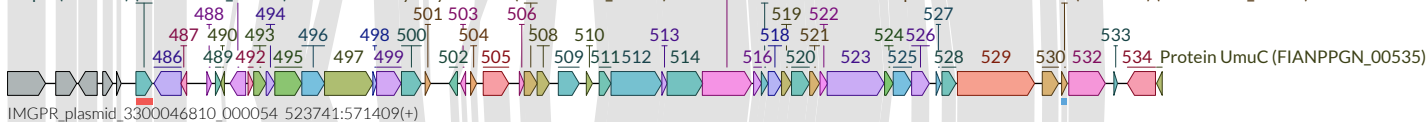


IMGPR_plasmid_2818991314_000002 297313:352629(+)

ATP-dependent Clp protease proteolytic subunit (FIANPPGN_00515)

putative HTH-type transcriptional regulator (FIANPPGN_00491)

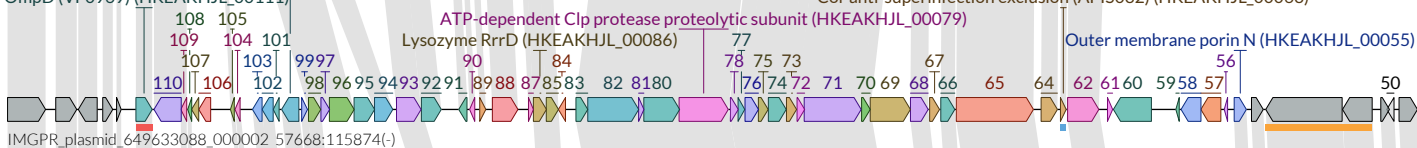
OmpD (VF0969) (FIANPPGN_00485)



IMGPR_plasmid_3300046810_000054 523741:571409(+)

HTH-type transcriptional regulator PrtR (HKEAKHJL_00100)

OmpD (VF0969) (HKEAKHJL_00111)

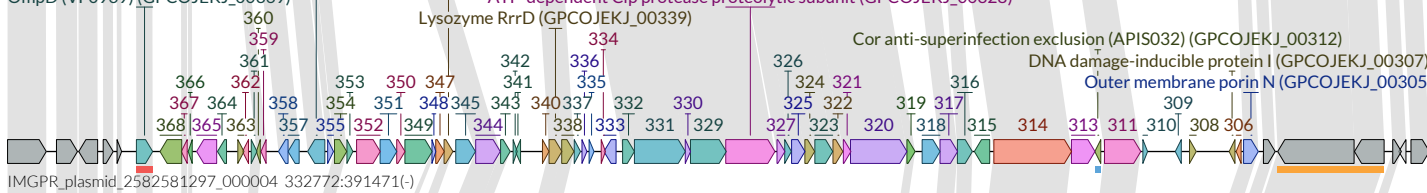


IMGPR_plasmid_649633088_000002 57668:115874(-)

Crossover junction endodeoxyribonuclease RusA (GPCOJEKJ_00346)

HTH-type transcriptional regulator PrtR (GPCOJEKJ_00356)

OmpD (VF0969) (GPCOJEKJ_00369)

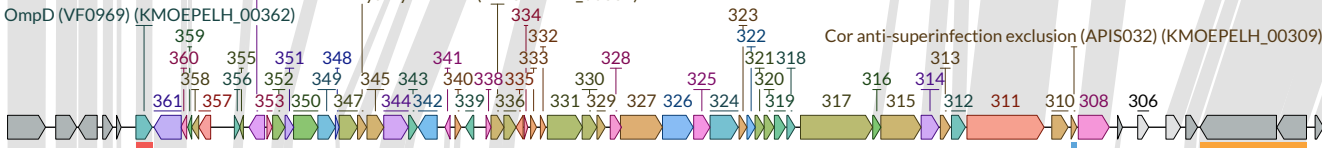


IMGPR_plasmid_2582581297_000004 332772:391471(-)

Crossover junction endodeoxyribonuclease RusA (KMOEPELH_00346)

putative HTH-type transcriptional regulator (KMOEPELH_00354)

OmpD (VF0969) (KMOEPELH_00362)

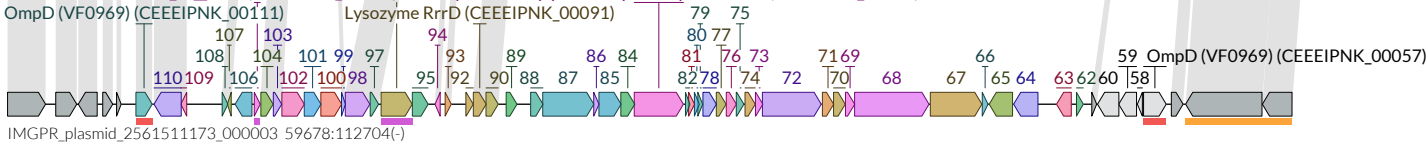


IMGPR_plasmid_2895116614_000001 333424:387760(-)

AAA domain, putative AbiEii toxin, Type IV TA system (CEEEIPNK_00096)

Retron_V1_HTH (CEEEIPNK_00105)

OmpD (VF0969) (CEEEIPNK_00111)



IMGPR_plasmid_2561511173_000003 59678:112704(-)

20000 nt

Defence Virulence Anti-defence AMR