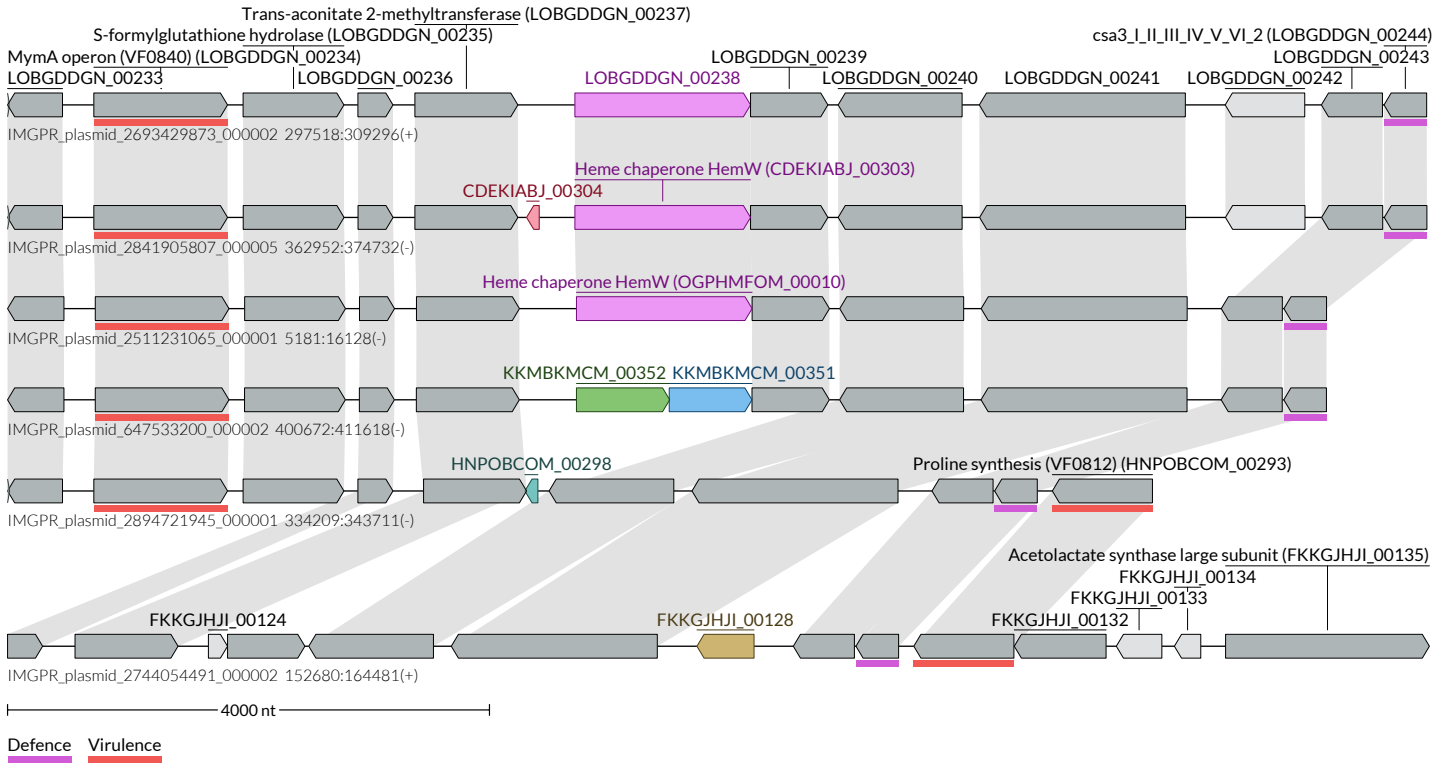
Trans-aconitate 2-methyltransferase (LOBGDDGN_00237)
S-formylglutathione hydrolase (LOBGDDGN_00235)
csa3_I_II_III_IV_V_VI_2 (LOBGDDGN_00244)
MymA operon (VF0840) (LOBGDDGN_00234)
LOBGDDGN_00239
LOBGDDGN_00243
LOBGDDGN_00233
LOBGDDGN_00236
LOBGDDGN_00238
LOBGDDGN_00240
LOBGDDGN_00241
LOBGDDGN_00242
IMGPR_plasmid_2693429873_000002 297518:309296(+)
Heme chaperone HemW (CDEKIABJ_00303)
CDEKIABJ_00304
IMGPR_plasmid_2841905807_000005 362952:374732(-)
Heme chaperone HemW (OGPHMFOM_00010)
IMGPR_plasmid_2511231065_000001 5181:16128(-)
KKMBKMCM_00352
KKMBKMCM_00351
IMGPR_plasmid_647533200_000002 400672:411618(-)
HNPOBCOM_00298
Proline synthesis (VF0812) (HNPOBCOM_00293)
IMGPR_plasmid_2894721945_000001 334209:343711(-)
Acetolactate synthase large subunit (FKKGJHJI_00135)
FKKGJHJI_00134
FKKGJHJI_00133
FKKGJHJI_00124
FKKGJHJI_00128
FKKGJHJI_00132
IMGPR_plasmid_2744054491_000002 152680:164481(+)
4000 nt
Defence
Virulence