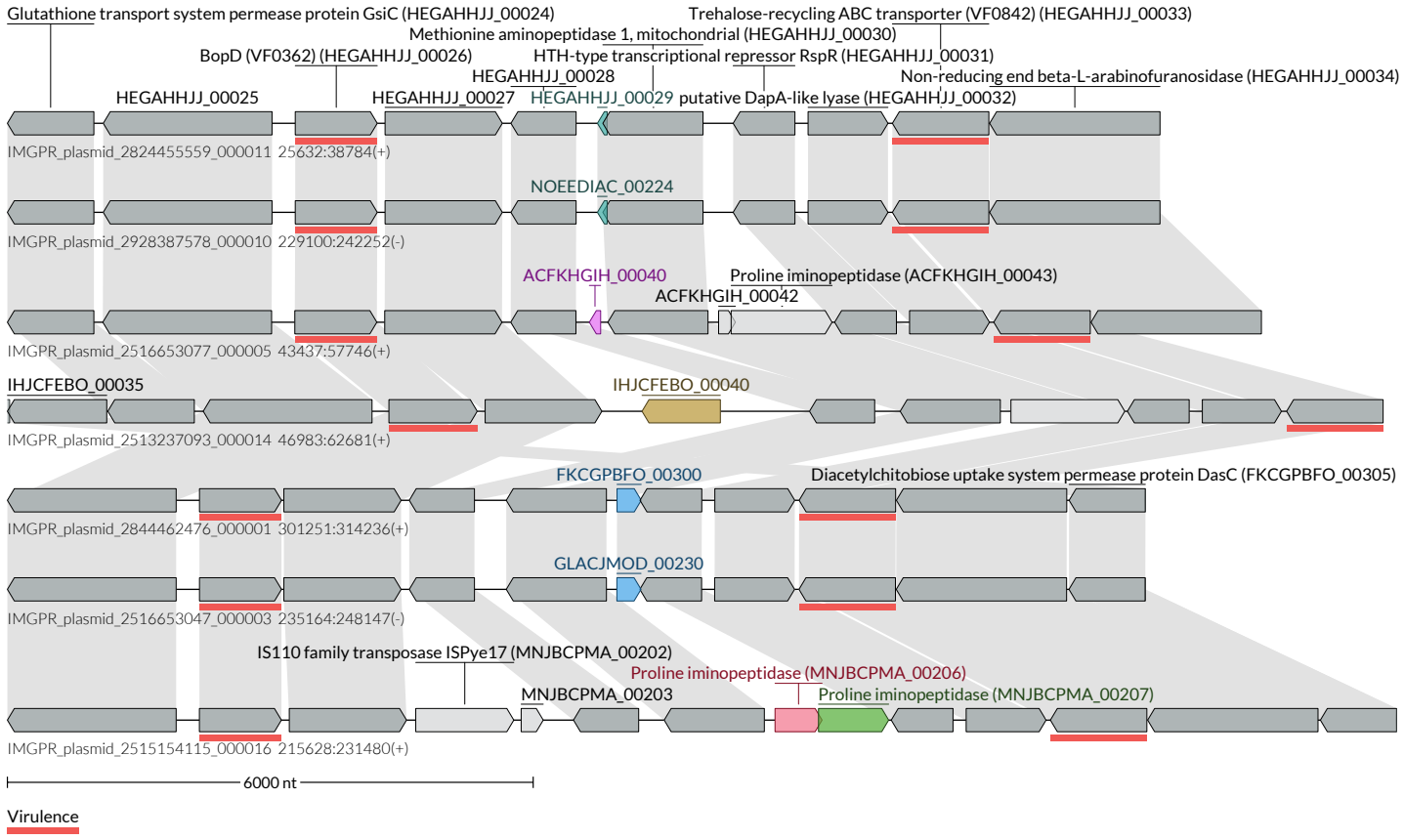
Glutathione transport system permease protein GsiC (HEGAHHJJ_00024)
Trehalose-recycling ABC transporter (VF0842) (HEGAHHJJ_00033)
Methionine aminopeptidase 1, mitochondrial (HEGAHHJJ_00030)
BopD (VF0362) (HEGAHHJJ_00026)
HTH-type transcriptional repressor RspR (HEGAHHJJ_00031)
HEGAHHJJ_00028
Non-reducing end beta-L-arabinofuranosidase (HEGAHHJJ_00034)
HEGAHHJJ_00025
HEGAHHJJ_00027
HEGAHHJJ_00029
putative DapA-like lyase (HEGAHHJJ_00032)
IMGPR_plasmid_2824455559_000011 25632:38784(+)
NOEEDIAC_00224
IMGPR_plasmid_2928387578_000010 229100:242252(-)
ACFKHGIH_00040
Proline iminopeptidase (ACFKHGIH_00043)
ACFKHGIH_00042
IMGPR_plasmid_2516653077_000005 43437:57746(+)
IHJCFEBO_00035
IHJCFEBO_00040
IMGPR_plasmid_2513237093_000014 46983:62681(+)
FKCGPBFO_00300
Diacetylchitobiose uptake system permease protein DasC (FKCGPBFO_00305)
IMGPR_plasmid_2844462476_000001 301251:314236(+)
GLACJMOD_00230
IMGPR_plasmid_2516653047_000003 235164:248147(-)
IS110 family transposase ISPye17 (MNJBCPMA_00202)
Proline iminopeptidase (MNJBCPMA_00206)
MNJBCPMA_00203
Proline iminopeptidase (MNJBCPMA_00207)
IMGPR_plasmid_2515154115_000016 215628:231480(+)
6000 nt
Virulence