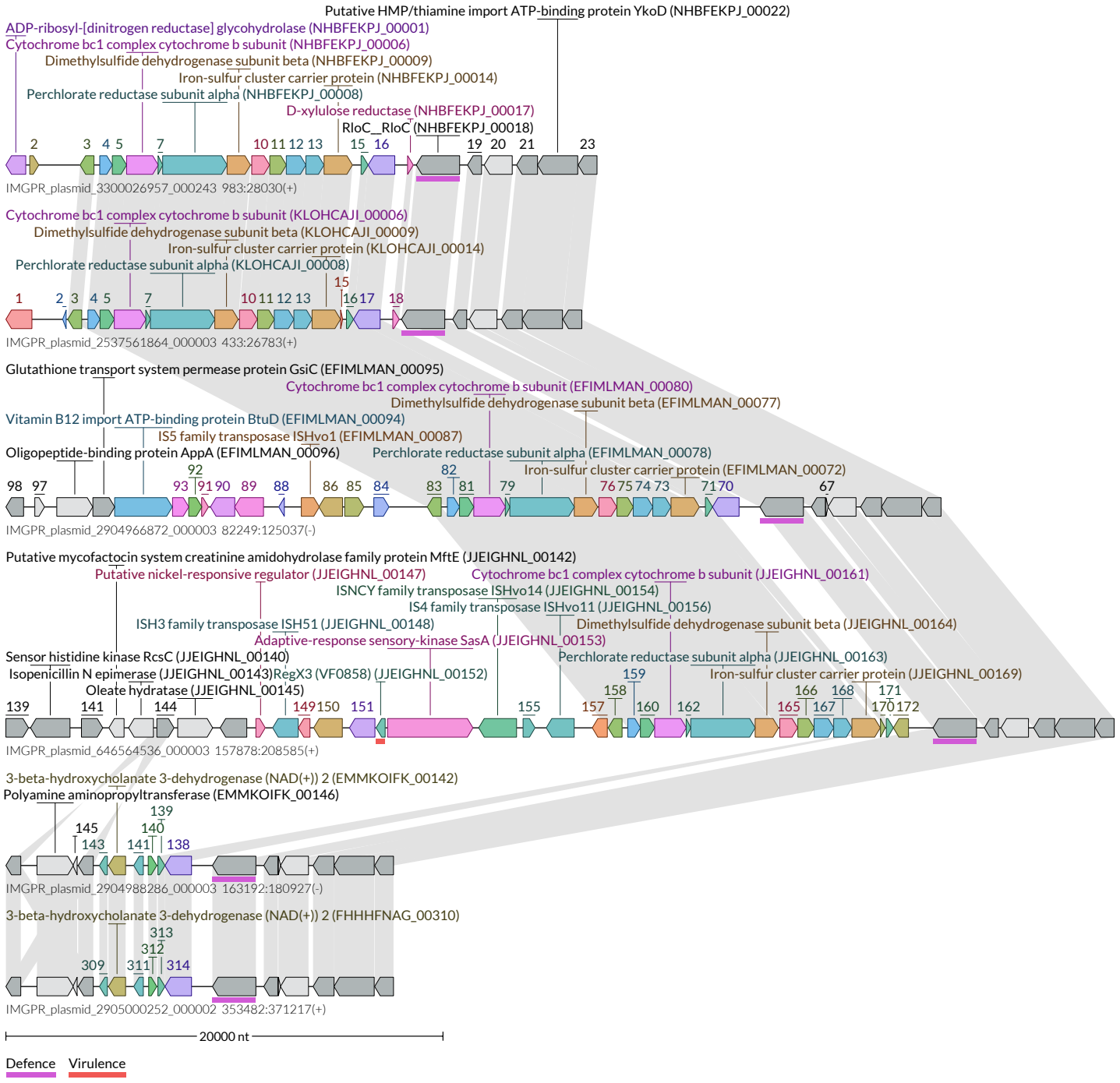
Putative HMP/thiamine import ATP-binding protein YkoD (NHBFEKPJ_00022)
ADP-ribosyl-[dinitrogen reductase] glycohydrolase (NHBFEKPJ_00001)
Cytochrome bc1 complex cytochrome b subunit (NHBFEKPJ_00006)
Dimethylsulfide dehydrogenase subunit beta (NHBFEKPJ_00009)
Iron-sulfur cluster carrier protein (NHBFEKPJ_00014)
Perchlorate reductase subunit alpha (NHBFEKPJ_00008)
D-xylulose reductase (NHBFEKPJ_00017)
RloC__RloC (NHBFEKPJ_00018)
IMGPR_plasmid_3300026957_000243 983:28030(+)
Cytochrome bc1 complex cytochrome b subunit (KLOHCAJI_00006)
Dimethylsulfide dehydrogenase subunit beta (KLOHCAJI_00009)
Iron-sulfur cluster carrier protein (KLOHCAJI_00014)
Perchlorate reductase subunit alpha (KLOHCAJI_00008)
IMGPR_plasmid_2537561864_000003 433:26783(+)
Glutathione transport system permease protein GsiC (EFIMLMAN_00095)
Cytochrome bc1 complex cytochrome b subunit (EFIMLMAN_00080)
Dimethylsulfide dehydrogenase subunit beta (EFIMLMAN_00077)
Vitamin B12 import ATP-binding protein BtuD (EFIMLMAN_00094)
IS5 family transposase ISHvo1 (EFIMLMAN_00087)
Oligopeptide-binding protein AppA (EFIMLMAN_00096)
Perchlorate reductase subunit alpha (EFIMLMAN_00078)
Iron-sulfur cluster carrier protein (EFIMLMAN_00072)
IMGPR_plasmid_2904966872_000003 82249:125037(-)
Putative mycofactocin system creatinine amidohydrolase family protein MftE (JJEIGHNL_00142)
Putative nickel-responsive regulator (JJEIGHNL_00147)
Cytochrome bc1 complex cytochrome b subunit (JJEIGHNL_00161)
ISNCY family transposase ISHvo14 (JJEIGHNL_00154)
IS4 family transposase ISHvo11 (JJEIGHNL_00156)
ISH3 family transposase ISH51 (JJEIGHNL_00148)
Dimethylsulfide dehydrogenase subunit beta (JJEIGHNL_00164)
Adaptive-response sensory-kinase SasA (JJEIGHNL_00153)
Sensor histidine kinase RcsC (JJEIGHNL_00140)
Perchlorate reductase subunit alpha (JJEIGHNL_00163)
Isopenicillin N epimerase (JJEIGHNL_00143)
RegX3 (VF0858) (JJEIGHNL_00152)
Iron-sulfur cluster carrier protein (JJEIGHNL_00169)
Oleate hydratase (JJEIGHNL_00145)
IMGPR_plasmid_646564536_000003 157878:208585(+)
3-beta-hydroxycholanate 3-dehydrogenase (NAD(+)) 2 (EMMKOIFK_00142)
Polyamine aminopropyltransferase (EMMKOIFK_00146)
IMGPR_plasmid_2904988286_000003 163192:180927(-)
3-beta-hydroxycholanate 3-dehydrogenase (NAD(+)) 2 (FHHHFNAG_00310)
IMGPR_plasmid_2905000252_000002 353482:371217(+)
20000 nt
Defence
Virulence