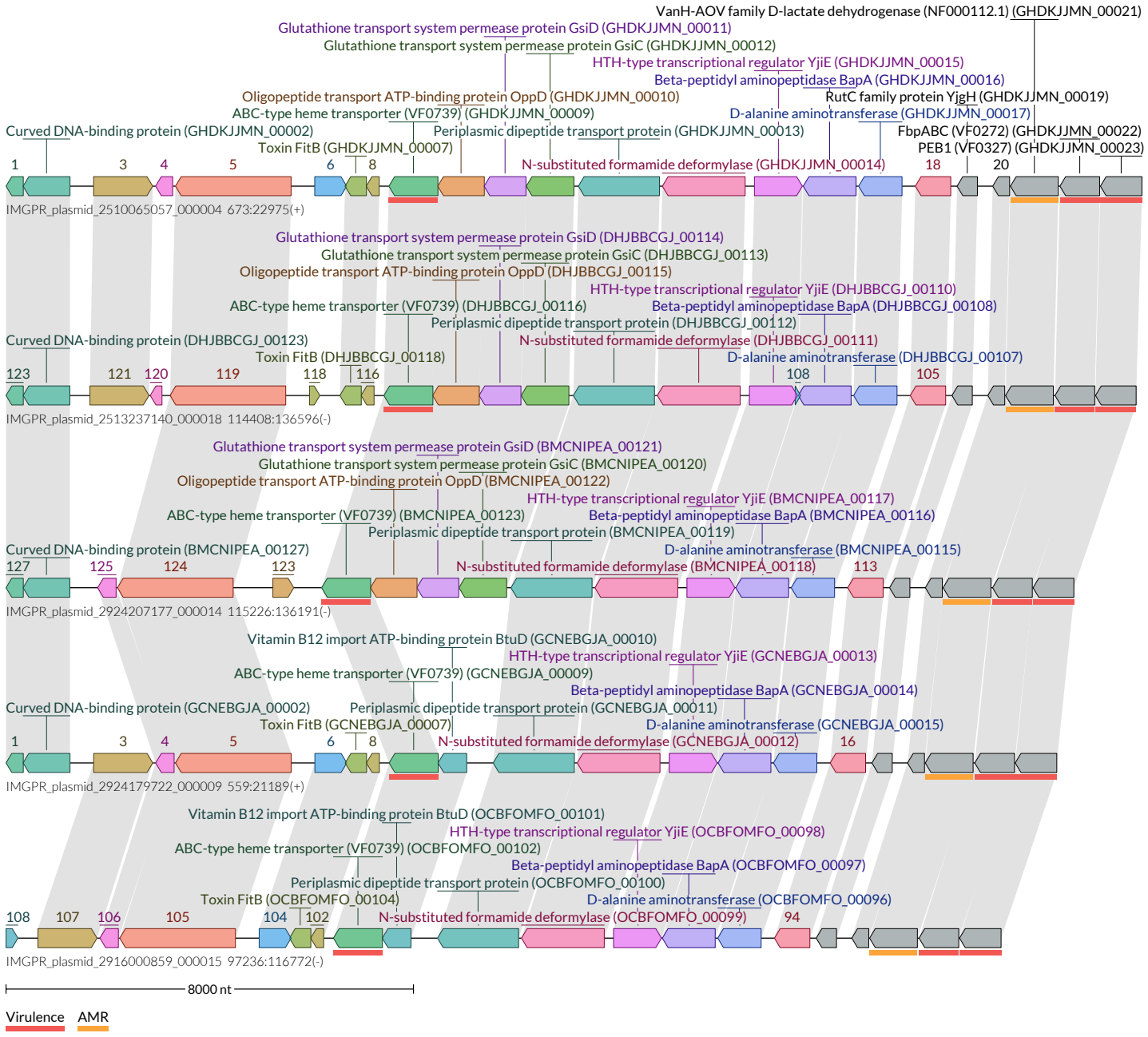
VanH-AOV family D-lactate dehydrogenase (NF000112.1) (GHDKJJMN_00021)
Glutathione transport system permease protein GsiD (GHDKJJMN_00011)
Glutathione transport system permease protein GsiC (GHDKJJMN_00012)
HTH-type transcriptional regulator YjiE (GHDKJJMN_00015)
Beta-peptidyl aminopeptidase BapA (GHDKJJMN_00016)
Oligopeptide transport ATP-binding protein OppD (GHDKJJMN_00010)
RutC family protein YjgH (GHDKJJMN_00019)
ABC-type heme transporter (VF0739) (GHDKJJMN_00009)
D-alanine aminotransferase (GHDKJJMN_00017)
Curved DNA-binding protein (GHDKJJMN_00002)
Periplasmic dipeptide transport protein (GHDKJJMN_00013)
FbpABC (VF0272) (GHDKJJMN_00022)
Toxin FitB (GHDKJJMN_00007)
PEB1 (VF0327) (GHDKJJMN_00023)
N-substituted formamide deformylase (GHDKJJMN_00014)
1 3 4 5 6 8 18 20
IMGPR_plasmid_2510065057_000004 673:22975(+)
Glutathione transport system permease protein GsiD (DHJBBCGJ_00114)
Glutathione transport system permease protein GsiC (DHJBBCGJ_00113)
Oligopeptide transport ATP-binding protein OppD (DHJBBCGJ_00115)
HTH-type transcriptional regulator YjiE (DHJBBCGJ_00110)
ABC-type heme transporter (VF0739) (DHJBBCGJ_00116)
Beta-peptidyl aminopeptidase BapA (DHJBBCGJ_00108)
Periplasmic dipeptide transport protein (DHJBBCGJ_00112)
Curved DNA-binding protein (DHJBBCGJ_00123)
N-substituted formamide deformylase (DHJBBCGJ_00111)
Toxin FitB (DHJBBCGJ_00118)
D-alanine aminotransferase (DHJBBCGJ_00107)
123 121 120 119 118 116 108 105
IMGPR_plasmid_2513237140_000018 114408:136596(-)
Glutathione transport system permease protein GsiD (BMCNIPEA_00121)
Glutathione transport system permease protein GsiC (BMCNIPEA_00120)
Oligopeptide transport ATP-binding protein OppD (BMCNIPEA_00122)
HTH-type transcriptional regulator YjiE (BMCNIPEA_00117)
ABC-type heme transporter (VF0739) (BMCNIPEA_00123)
Beta-peptidyl aminopeptidase BapA (BMCNIPEA_00116)
Periplasmic dipeptide transport protein (BMCNIPEA_00119)
Curved DNA-binding protein (BMCNIPEA_00127)
D-alanine aminotransferase (BMCNIPEA_00115)
N-substituted formamide deformylase (BMCNIPEA_00118)
127 125 124 123 113
IMGPR_plasmid_2924207177_000014 115226:136191(-)
Vitamin B12 import ATP-binding protein BtuD (GCNEBGJA_00010)
HTH-type transcriptional regulator YjiE (GCNEBGJA_00013)
ABC-type heme transporter (VF0739) (GCNEBGJA_00009)
Beta-peptidyl aminopeptidase BapA (GCNEBGJA_00014)
Curved DNA-binding protein (GCNEBGJA_00002)
Periplasmic dipeptide transport protein (GCNEBGJA_00011)
Toxin FitB (GCNEBGJA_00007)
D-alanine aminotransferase (GCNEBGJA_00015)
N-substituted formamide deformylase (GCNEBGJA_00012)
1 3 4 5 6 8 16
IMGPR_plasmid_2924179722_000009 559:21189(+)
Vitamin B12 import ATP-binding protein BtuD (OCBFOMFO_00101)
HTH-type transcriptional regulator YjiE (OCBFOMFO_00098)
ABC-type heme transporter (VF0739) (OCBFOMFO_00102)
Beta-peptidyl aminopeptidase BapA (OCBFOMFO_00097)
Periplasmic dipeptide transport protein (OCBFOMFO_00100)
Toxin FitB (OCBFOMFO_00104)
D-alanine aminotransferase (OCBFOMFO_00096)
N-substituted formamide deformylase (OCBFOMFO_00099)
108 107 106 105 104 102 94
IMGPR_plasmid_2916000859_000015 97236:116772(-)
8000 nt
Virulence
AMR