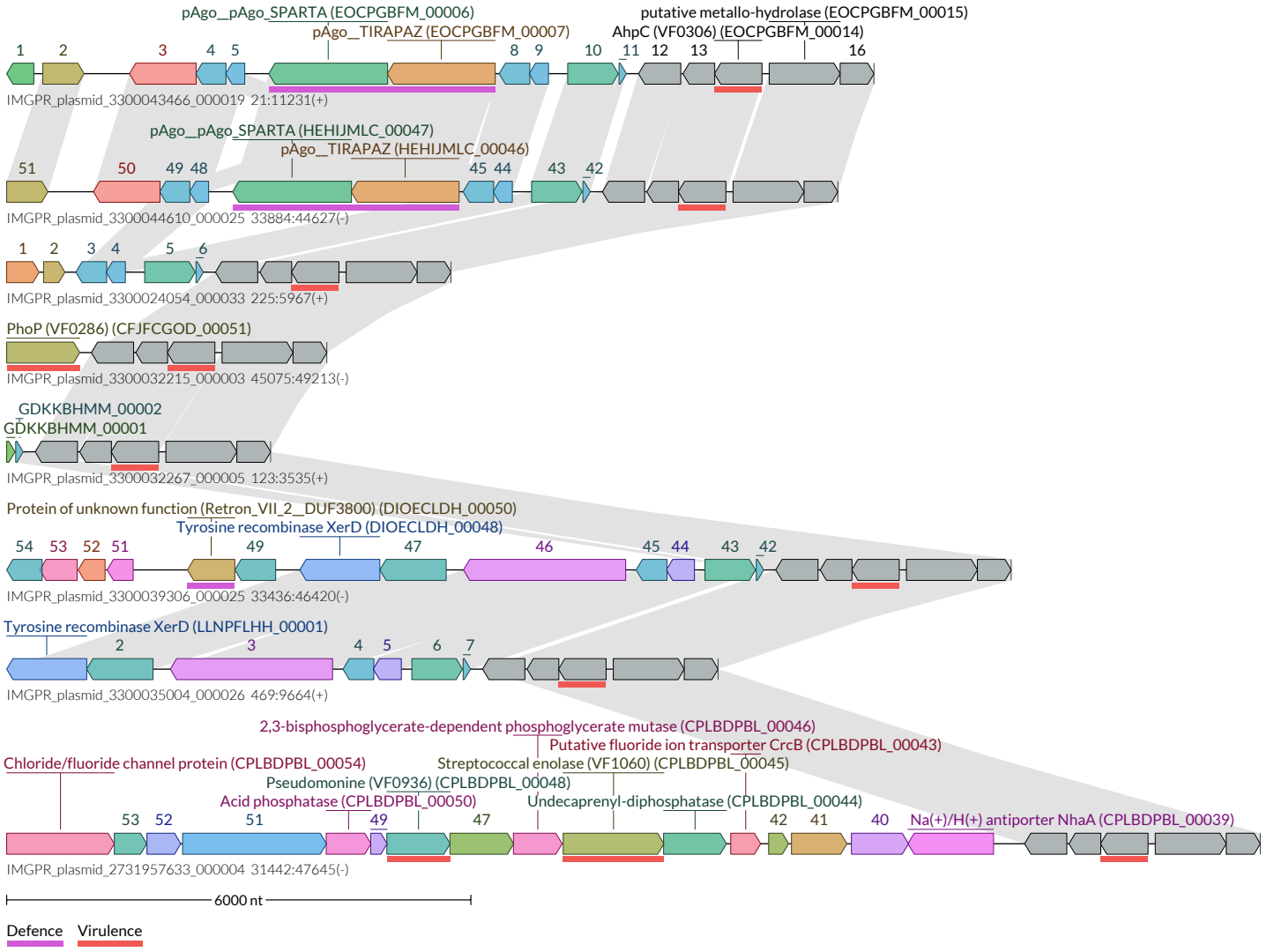
pAgo__pAgo_SPARTA (EOCPGBFM_00006)
pAgo__TIRAPAZ (EOCPGBFM_00007)
putative metallo-hydrolase (EOCPGBFM_00015)
AhpC (VF0306) (EOCPGBFM_00014)
1 2 3 4 5 8 9 10 11 12 13 16
IMGPR_plasmid_3300043466_000019 21:11231(+)
pAgo__pAgo_SPARTA (HEHIJMLC_00047)
pAgo__TIRAPAZ (HEHIJMLC_00046)
51 50 49 48 45 44 43 42
IMGPR_plasmid_3300044610_000025 33884:44627(-)
1 2 3 4 5 6
IMGPR_plasmid_3300024054_000033 225:5967(+)
PhoP (VF0286) (CFJFCGOD_00051)
IMGPR_plasmid_3300032215_000003 45075:49213(-)
GDKKBHMM_00002
GDKKBHMM_00001
IMGPR_plasmid_3300032267_000005 123:3535(+)
Protein of unknown function (Retron_VII_2__DUF3800) (DIOECLDH_00050)
Tyrosine recombinase XerD (DIOECLDH_00048)
54 53 52 51 49 47 46 45 44 43 42
IMGPR_plasmid_3300039306_000025 33436:46420(-)
Tyrosine recombinase XerD (LLNPFLHH_00001)
2 3 4 5 6 7
IMGPR_plasmid_3300035004_000026 469:9664(+)
2,3-bisphosphoglycerate-dependent phosphoglycerate mutase (CPLBDPBL_00046)
Putative fluoride ion transporter CrcB (CPLBDPBL_00043)
Chloride/fluoride channel protein (CPLBDPBL_00054)
Streptococcal enolase (VF1060) (CPLBDPBL_00045)
Pseudomonine (VF0936) (CPLBDPBL_00048)
Acid phosphatase (CPLBDPBL_00050)
Undecaprenyl-diphosphatase (CPLBDPBL_00044)
53 52 51 49 47 42 41 40
Na(+)/H(+) antiporter NhaA (CPLBDPBL_00039)
IMGPR_plasmid_2731957633_000004 31442:47645(-)
6000 nt
Defence
Virulence