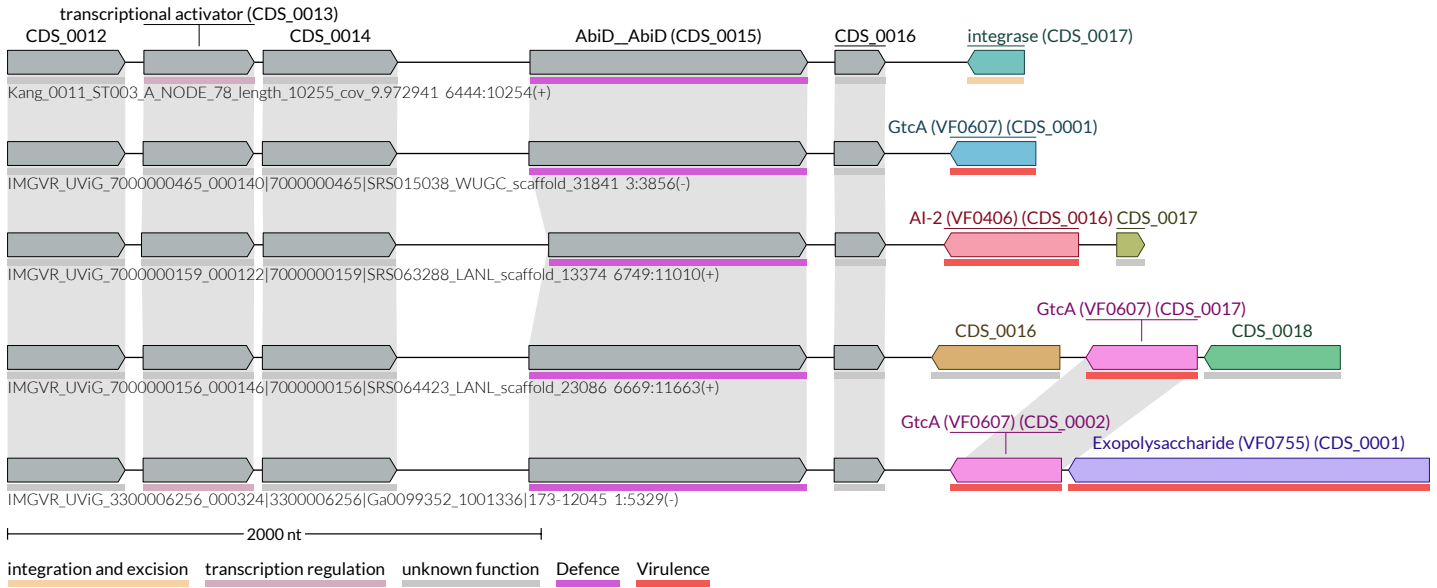
transcriptional activator (CDS_0013)
CDS_0012
CDS_0014
AbiD__AbiD (CDS_0015)
CDS_0016
integrase (CDS_0017)
Kang_0011_ST003_A_NODE_78_length_10255_cov_9.972941 6444:10254(+)
GtcA (VF0607) (CDS_0001)
IMGVR_UViG_7000000465_000140|7000000465|SRS015038_WUGC_scaffold_31841 3:3856(-)
AI-2 (VF0406) (CDS_0016)
CDS_0017
IMGVR_UViG_7000000159_000122|7000000159|SRS063288_LANL_scaffold_13374 6749:11010(+)
GtcA (VF0607) (CDS_0017)
CDS_0016
CDS_0018
IMGVR_UViG_7000000156_000146|7000000156|SRS064423_LANL_scaffold_23086 6669:11663(+)
GtcA (VF0607) (CDS_0002)
Exopolysaccharide (VF0755) (CDS_0001)
IMGVR_UViG_3300006256_000324|3300006256|Ga0099352_1001336|173-12045 1:5329(-)
2000 nt
integration and excision
transcription regulation
unknown function
Defence
Virulence