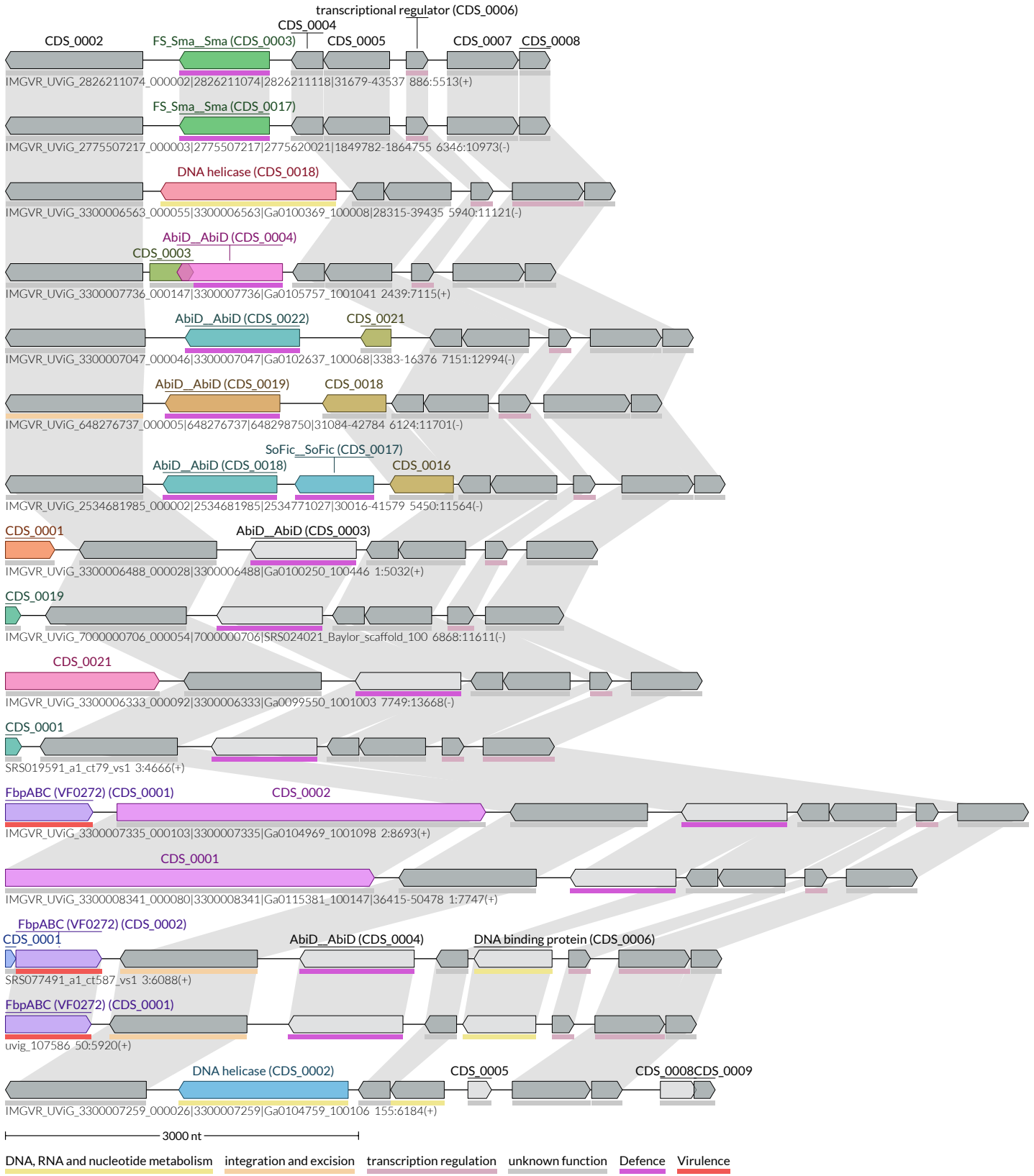
transcriptional regulator (CDS_0006)
CDS_0004
CDS_0002
FS_Sma__Sma (CDS_0003)
CDS_0005
CDS_0007
CDS_0008
IMGVR_UViG_2826211074_000002|2826211074|2826211118|31679-43537 886:5513(+)
FS_Sma__Sma (CDS_0017)
IMGVR_UViG_2775507217_000003|2775507217|2775620021|1849782-1864755 6346:10973(-)
DNA helicase (CDS_0018)
IMGVR_UViG_3300006563_000055|3300006563|Ga0100369_100008|28315-39435 5940:11121(-)
AbiD__AbiD (CDS_0004)
CDS_0003
IMGVR_UViG_3300007736_000147|3300007736|Ga0105757_1001041 2439:7115(+)
AbiD__AbiD (CDS_0022)
CDS_0021
IMGVR_UViG_3300007047_000046|3300007047|Ga0102637_100068|3383-16376 7151:12994(-)
AbiD__AbiD (CDS_0019)
CDS_0018
IMGVR_UViG_648276737_000005|648276737|648298750|31084-42784 6124:11701(-)
SoFic__SoFic (CDS_0017)
AbiD__AbiD (CDS_0018)
CDS_0016
IMGVR_UViG_2534681985_000002|2534681985|2534771027|30016-41579 5450:11564(-)
CDS_0001
AbiD__AbiD (CDS_0003)
IMGVR_UViG_3300006488_000028|3300006488|Ga0100250_100446 1:5032(+)
CDS_0019
IMGVR_UViG_7000000706_000054|7000000706|SRS024021_Baylor_scaffold_100 6868:11611(-)
CDS_0021
IMGVR_UViG_3300006333_000092|3300006333|Ga0099550_1001003 7749:13668(-)
CDS_0001
SRS019591_a1_ct79_vs1 3:4666(+)
FbpABC (VF0272) (CDS_0001)
CDS_0002
IMGVR_UViG_3300007335_000103|3300007335|Ga0104969_1001098 2:8693(+)
CDS_0001
IMGVR_UViG_3300008341_000080|3300008341|Ga0115381_100147|36415-50478 1:7747(+)
FbpABC (VF0272) (CDS_0002)
CDS_0001
AbiD__AbiD (CDS_0004)
DNA binding protein (CDS_0006)
SRS077491_a1_ct587_vs1 3:6088(+)
FbpABC (VF0272) (CDS_0001)
uvig_107586 50:5920(+)
DNA helicase (CDS_0002)
CDS_0005
CDS_0008CDS_0009
IMGVR_UViG_3300007259_000026|3300007259|Ga0104759_100106 155:6184(+)
3000 nt
DNA, RNA and nucleotide metabolism
integration and excision
transcription regulation
unknown function
Defence
Virulence