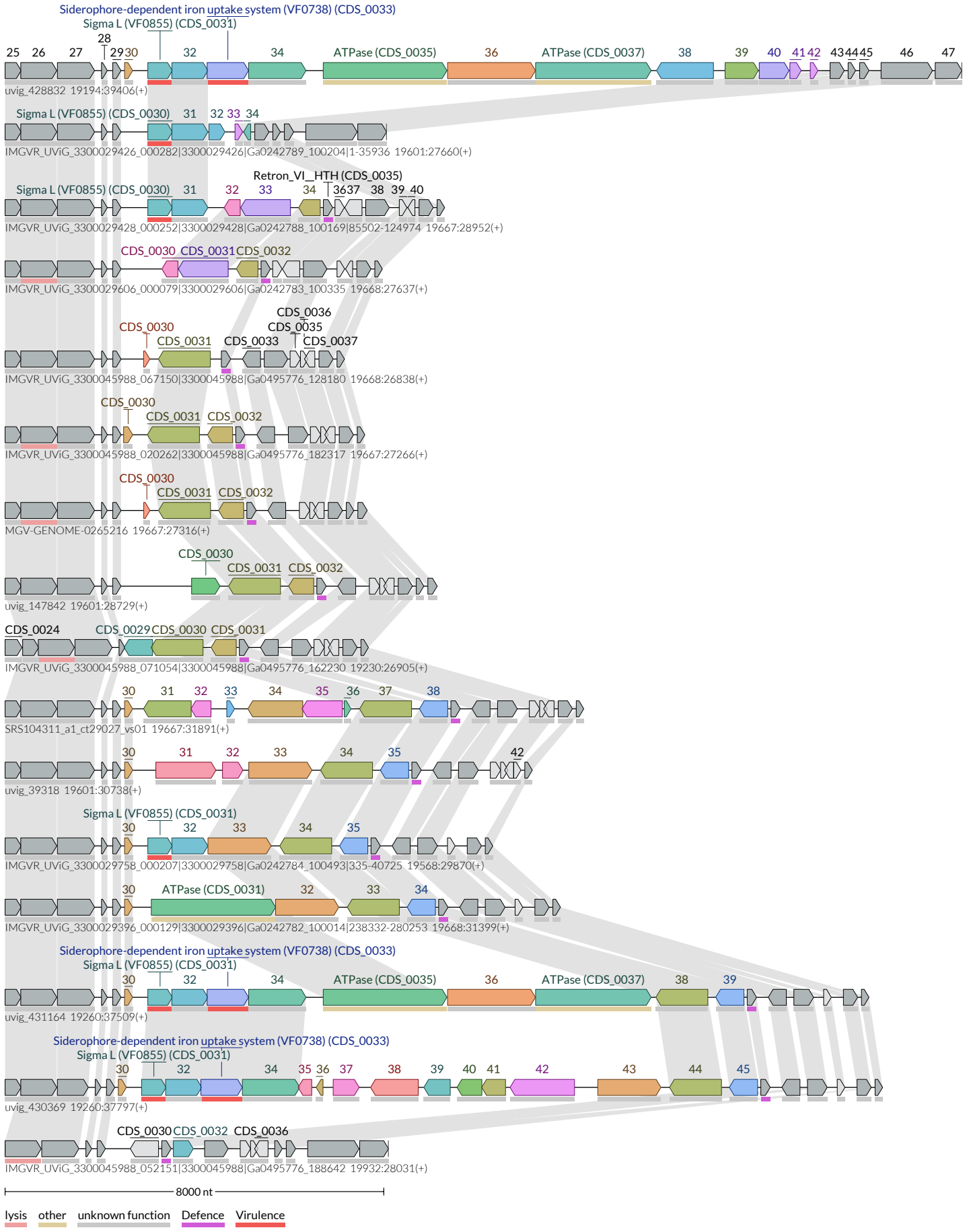
Siderophore-dependent iron uptake system (VF0738) (CDS_0033)
Sigma L (VF0855) (CDS_0031)
25 26 27 28 29 30 32 34 ATPase (CDS_0035) 36 ATPase (CDS_0037) 38 39 40 41 42 43 44 45 46 47
uvig_428832 19194:39406(+)
Sigma L (VF0855) (CDS_0030) 31 32 33 34
IMGVR_UViG_3300029426_000282|3300029426|Ga0242789_100204|1-35936 19601:27660(+)
Retron_VI_HTH (CDS_0035)
Sigma L (VF0855) (CDS_0030) 31 32 33 34 36 37 38 39 40
IMGVR_UViG_3300029428_000252|3300029428|Ga0242788_100169|85502-124974 19667:28952(+)
CDS_0030 CDS_0031 CDS_0032
IMGVR_UViG_3300029606_000079|3300029606|Ga0242783_100335 19668:27637(+)
CDS_0036
CDS_0030 CDS_0035
CDS_0031 CDS_0033 CDS_0037
IMGVR_UViG_3300045988_067150|3300045988|Ga0495776_128180 19668:26838(+)
CDS_0030
CDS_0031 CDS_0032
IMGVR_UViG_3300045988_020262|3300045988|Ga0495776_182317 19667:27266(+)
CDS_0030
CDS_0031 CDS_0032
MGV-GENOME-0265216 19667:27316(+)
CDS_0030
CDS_0031 CDS_0032
uvig_147842 19601:28729(+)
CDS_0024 CDS_0029 CDS_0030 CDS_0031
IMGVR_UViG_3300045988_071054|3300045988|Ga0495776_162230 19230:26905(+)
30 31 32 33 34 35 36 37 38
SRS104311_a1_ct29027_vs01 19667:31891(+)
30 31 32 33 34 35 42
uvig_39318 19601:30738(+)
Sigma L (VF0855) (CDS_0031)
30 32 33 34 35
IMGVR_UViG_3300029758_000207|3300029758|Ga0242784_100493|335-40725 19568:29870(+)
30 ATPase (CDS_0031) 32 33 34
IMGVR_UViG_3300029396_000129|3300029396|Ga0242782_100014|238332-280253 19668:31399(+)
Siderophore-dependent iron uptake system (VF0738) (CDS_0033)
Sigma L (VF0855) (CDS_0031)
30 32 34 ATPase (CDS_0035) 36 ATPase (CDS_0037) 38 39
uvig_431164 19260:37509(+)
Siderophore-dependent iron uptake system (VF0738) (CDS_0033)
Sigma L (VF0855) (CDS_0031)
30 32 34 35 36 37 38 39 40 41 42 43 44 45
uvig_430369 19260:37797(+)
CDS_0030 CDS_0032 CDS_0036
IMGVR_UViG_3300045988_052151|3300045988|Ga0495776_188642 19932:28031(+)
8000 nt
lysis
other
unknown function
Defence
Virulence