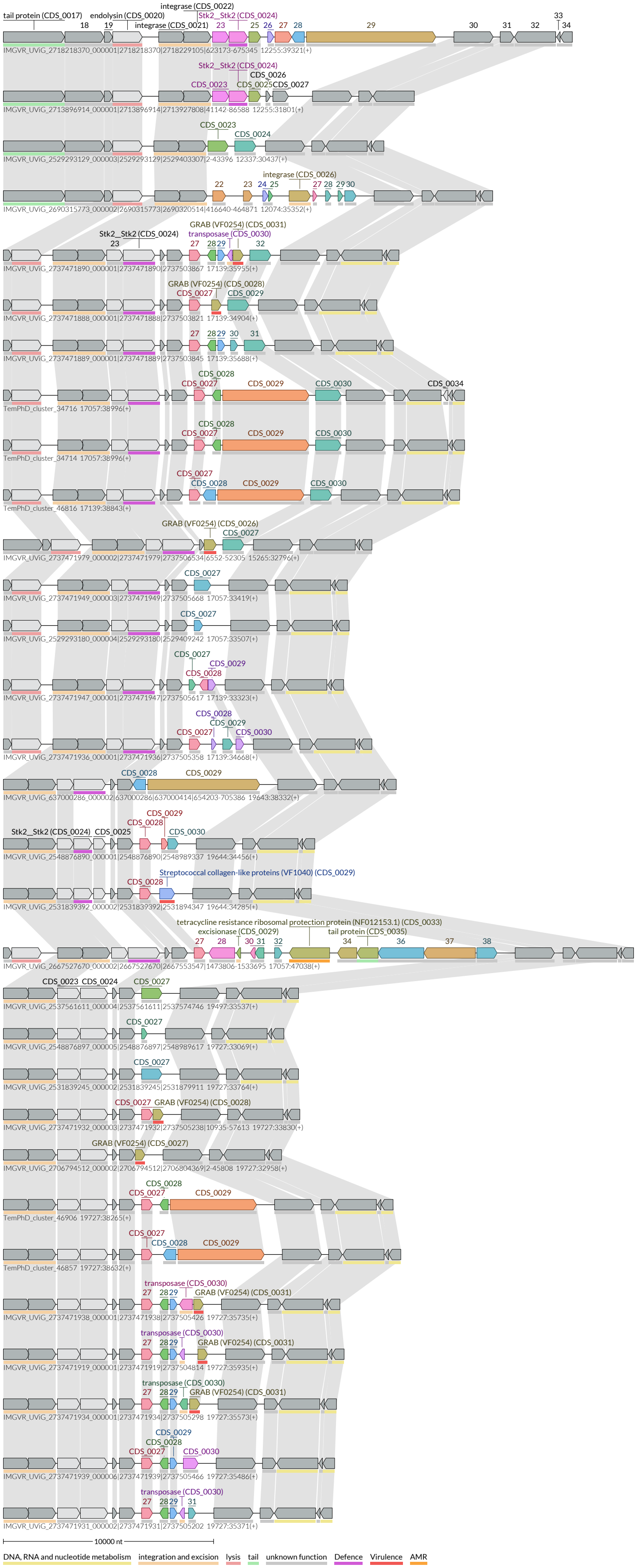
tail protein (CDS_0017)
endolysin (CDS_0020)
integrase (CDS_0022)
Stk2__Stk2 (CDS_0024)
integrase (CDS_0021)
IMGVR_UViG_2718218370_000001|2718218370|2718229105|623173-675345 12255:39321(+)
Stk2__Stk2 (CDS_0024)
CDS_0026
CDS_0023
CDS_0025
CDS_0027
IMGVR_UViG_2713896914_000001|2713896914|2713927808|41142-86588 12255:31801(+)
CDS_0023
CDS_0024
IMGVR_UViG_2529293129_000003|2529293129|2529403307|2-43396 12337:30437(+)
integrase (CDS_0026)
IMGVR_UViG_2690315773_000002|2690315773|2690320514|416640-464871 12074:35352(+)
GRAB (VF0254) (CDS_0031)
Stk2__Stk2 (CDS_0024)
transposase (CDS_0030)
IMGVR_UViG_2737471890_000001|2737471890|2737503867 17139:35955(+)
GRAB (VF0254) (CDS_0028)
CDS_0027
CDS_0029
IMGVR_UViG_2737471888_000001|2737471888|2737503821 17139:34904(+)
IMGVR_UViG_2737471889_000001|2737471889|2737503845 17139:35688(+)
CDS_0028
CDS_0027
CDS_0029
CDS_0030
CDS_0034
TemPhD_cluster_34716 17057:38996(+)
TemPhD_cluster_34714 17057:38996(+)
CDS_0027
CDS_0028
TemPhD_cluster_46816 17139:38843(+)
GRAB (VF0254) (CDS_0026)
IMGVR_UViG_2737471979_000002|2737471979|2737506534|6552-52305 15265:32796(+)
IMGVR_UViG_2737471949_000003|2737471949|2737505668 17057:33419(+)
IMGVR_UViG_2529293180_000004|2529293180|2529409242 17057:33507(+)
IMGVR_UViG_2737471947_000001|2737471947|2737505617 17139:33323(+)
IMGVR_UViG_2737471936_000001|2737471936|2737505358 17139:34668(+)
IMGVR_UViG_637000286_000002|637000286|637000414|654203-705386 19643:38332(+)
Stk2__Stk2 (CDS_0024)
IMGVR_UViG_2548876890_000001|2548876890|2548989337 19644:34456(+)
Streptococcal collagen-like proteins (VF1040) (CDS_0029)
IMGVR_UViG_2531839392_000002|2531839392|2531894347 19644:34285(+)
tetracycline resistance ribosomal protection protein (NF012153.1) (CDS_0033)
excisionase (CDS_0029)
tail protein (CDS_0035)
IMGVR_UViG_2667527670_000002|2667527670|2667553547|1473806-1533695 17057:47038(+)
CDS_0023
CDS_0024
IMGVR_UViG_2537561611_000004|2537561611|2537574746 19497:33537(+)
IMGVR_UViG_2548876897_000005|2548876897|2548989617 19727:33069(+)
IMGVR_UViG_2531839245_000002|2531839245|2531879911 19727:33764(+)
IMGVR_UViG_2737471932_000003|2737471932|2737505238|10935-57613 19727:33830(+)
GRAB (VF0254) (CDS_0027)
IMGVR_UViG_2706794512_000002|2706794512|2706804369|2-45808 19727:32958(+)
TemPhD_cluster_46906 19727:38265(+)
TemPhD_cluster_46857 19727:38632(+)
transposase (CDS_0030)
GRAB (VF0254) (CDS_0031)
IMGVR_UViG_2737471938_000001|2737471938|2737505426 19727:35735(+)
IMGVR_UViG_2737471919_000001|2737471919|2737504814 19727:35935(+)
IMGVR_UViG_2737471934_000001|2737471934|2737505298 19727:35573(+)
IMGVR_UViG_2737471939_000006|2737471939|2737505466 19727:35486(+)
IMGVR_UViG_2737471931_000002|2737471931|2737505202 19727:35371(+)
10000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
tail
unknown function
Defence
Virulence
AMR