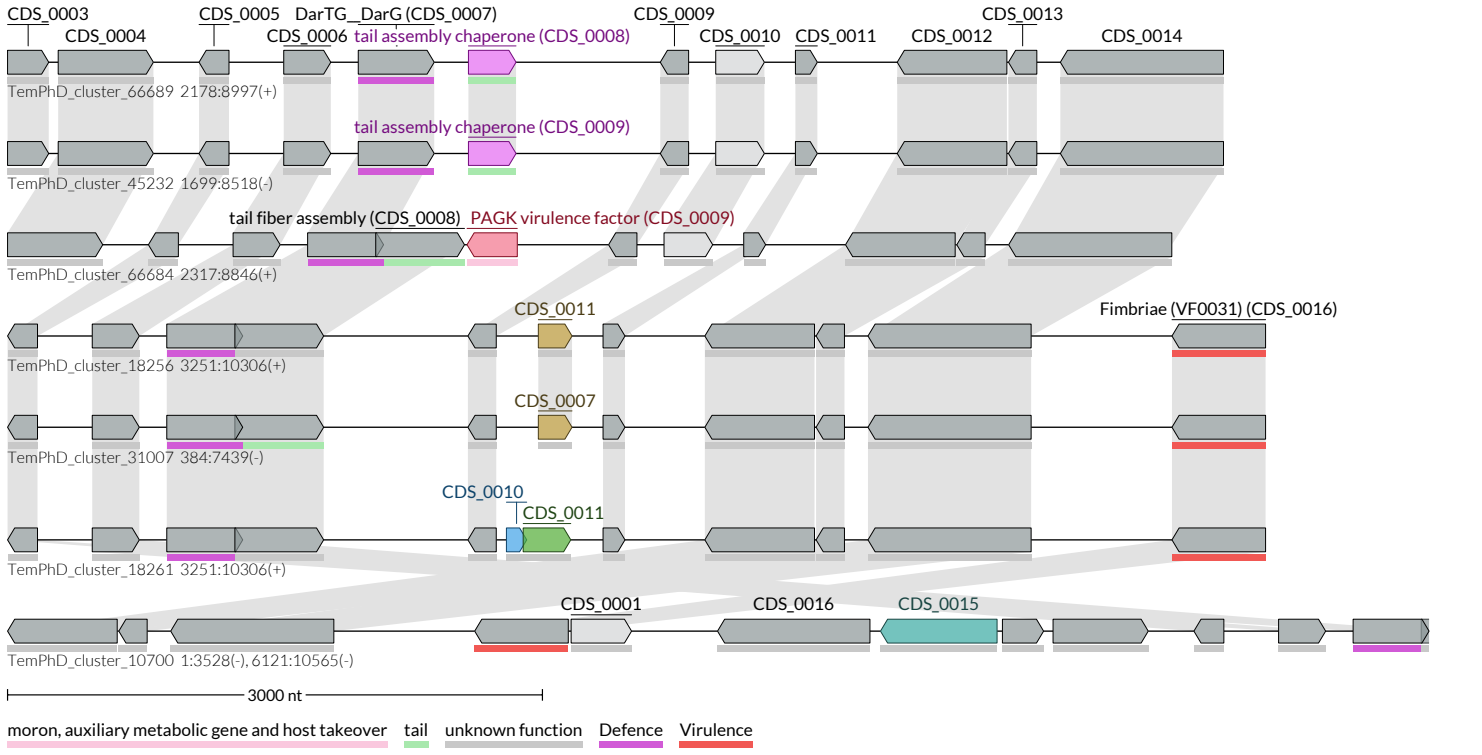
CDS_0003
CDS_0004
CDS_0005
DarTG__DarG (CDS_0007)
CDS_0006
tail assembly chaperone (CDS_0008)
CDS_0009
CDS_0010
CDS_0011
CDS_0012
CDS_0013
CDS_0014
TemPhD_cluster_66689 2178:8997(+)
tail assembly chaperone (CDS_0009)
TemPhD_cluster_45232 1699:8518(-)
tail fiber assembly (CDS_0008)
PAGK virulence factor (CDS_0009)
TemPhD_cluster_66684 2317:8846(+)
CDS_0011
Fimbriae (VF0031) (CDS_0016)
TemPhD_cluster_18256 3251:10306(+)
CDS_0007
TemPhD_cluster_31007 384:7439(-)
CDS_0010
CDS_0011
TemPhD_cluster_18261 3251:10306(+)
CDS_0001
CDS_0016
CDS_0015
TemPhD_cluster_10700 1:3528(-), 6121:10565(-)
3000 nt
moron, auxiliary metabolic gene and host takeover
tail
unknown function
Defence
Virulence