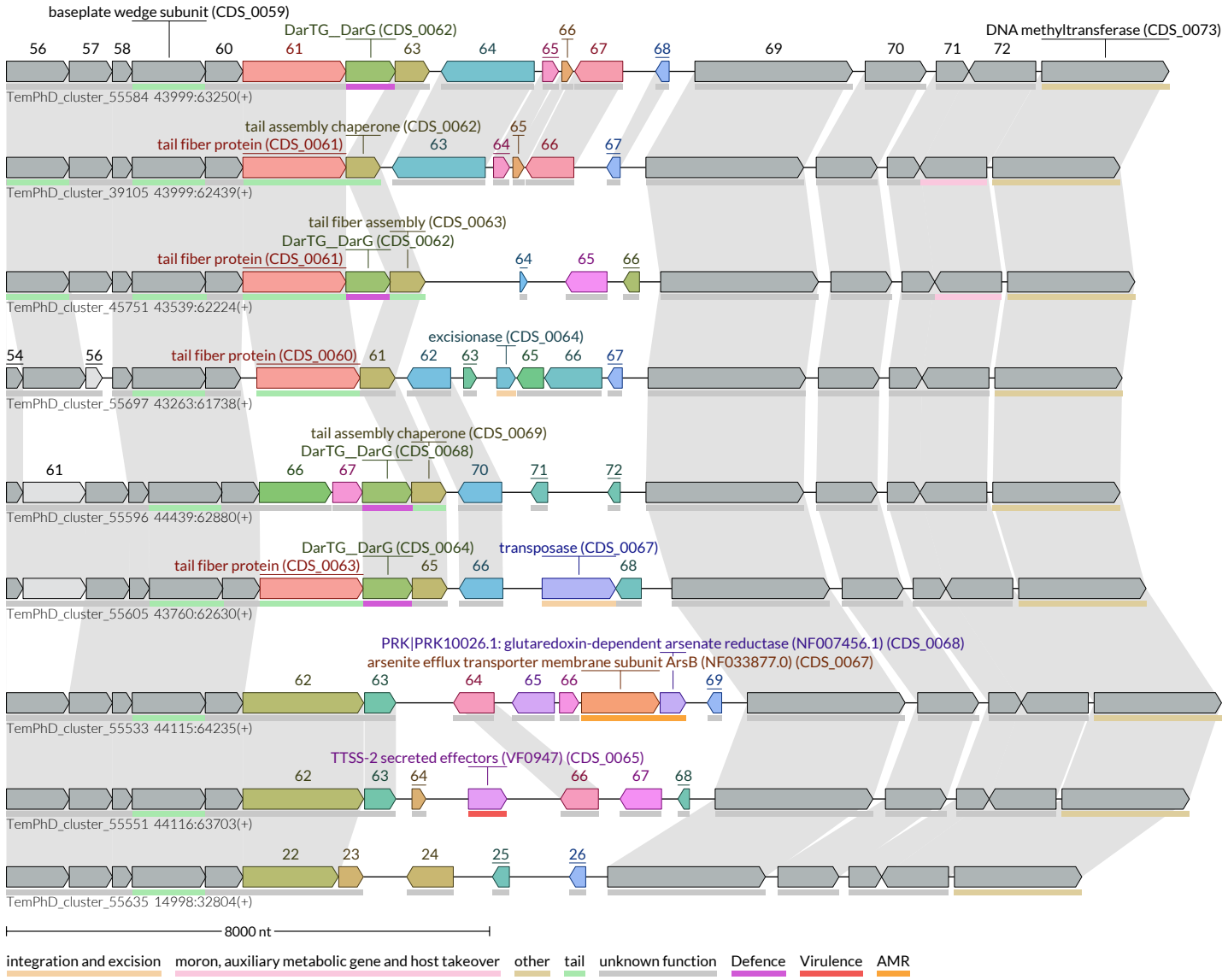
baseplate wedge subunit (CDS_0059)
DarTG__DarG (CDS_0062)
DNA methyltransferase (CDS_0073)
56 57 58 60 61 63 64 65 66 67 68 69 70 71 72
TemPhD_cluster_55584 43999:63250(+)
tail assembly chaperone (CDS_0062)
tail fiber protein (CDS_0061)
63 64 65 66 67
TemPhD_cluster_39105 43999:62439(+)
tail fiber assembly (CDS_0063)
DarTG__DarG (CDS_0062)
tail fiber protein (CDS_0061)
64 65 66
TemPhD_cluster_45751 43539:62224(+)
excisionase (CDS_0064)
54 56 tail fiber protein (CDS_0060) 61 62 63 65 66 67
TemPhD_cluster_55697 43263:61738(+)
tail assembly chaperone (CDS_0069)
DarTG__DarG (CDS_0068)
61 66 67 70 71 72
TemPhD_cluster_55596 44439:62880(+)
DarTG__DarG (CDS_0064)
transposase (CDS_0067)
tail fiber protein (CDS_0063)
65 66 68
TemPhD_cluster_55605 43760:62630(+)
PRK|PRK10026.1: glutaredoxin-dependent arsenate reductase (NF007456.1) (CDS_0068)
arsenite efflux transporter membrane subunit ArsB (NF033877.0) (CDS_0067)
62 63 64 65 66 69
TemPhD_cluster_55533 44115:64235(+)
TTSS-2 secreted effectors (VF0947) (CDS_0065)
62 63 64 66 67 68
TemPhD_cluster_55551 44116:63703(+)
22 23 24 25 26
TemPhD_cluster_55635 14998:32804(+)
8000 nt
integration and excision
moron, auxiliary metabolic gene and host takeover
other
tail
unknown function
Defence
Virulence
AMR