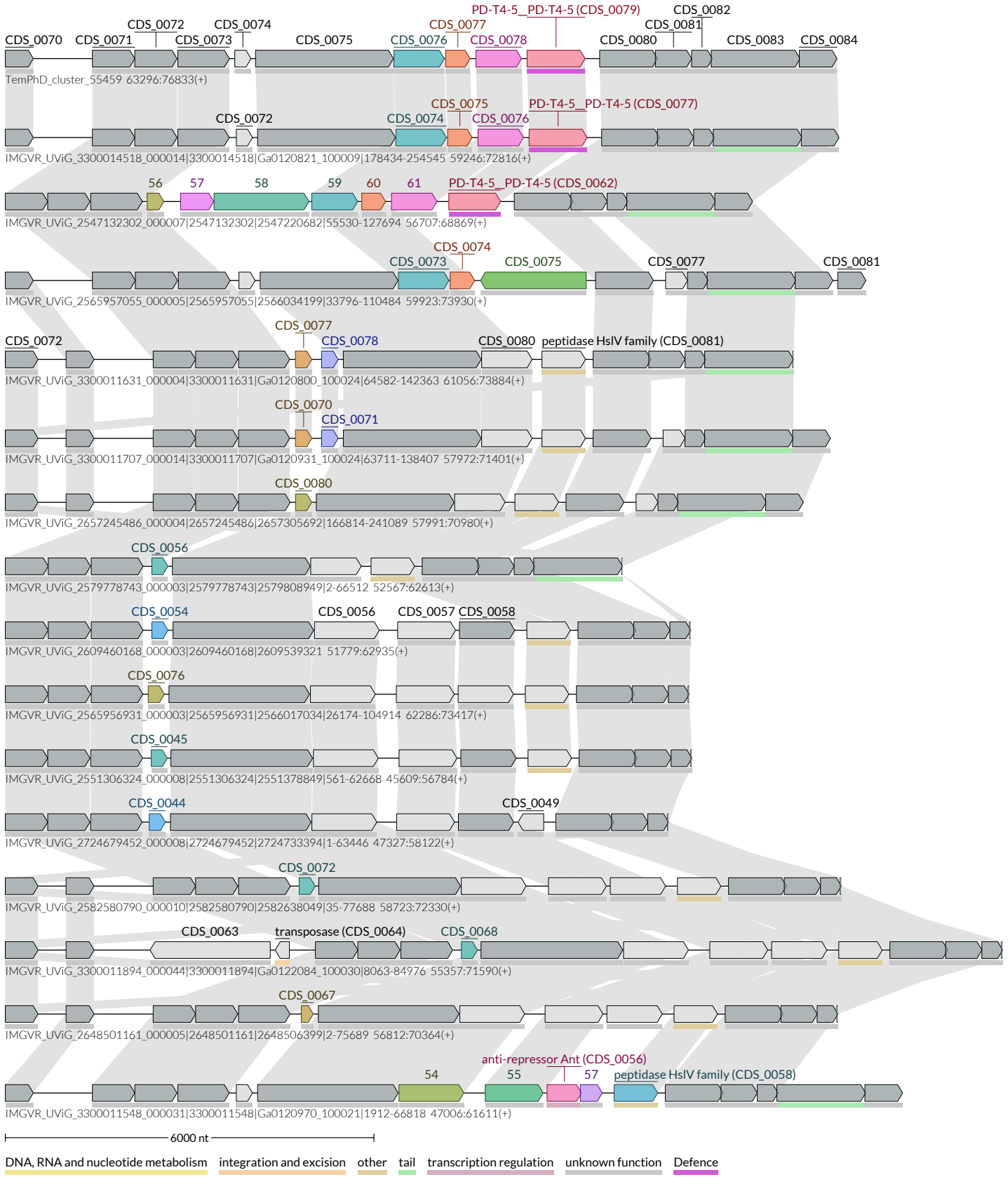
CDS_0070
CDS_0071
CDS_0072
CDS_0073
CDS_0074
CDS_0075
CDS_0076
CDS_0077
CDS_0078
PD-T4-5__PD-T4-5 (CDS_0079)
CDS_0080
CDS_0081
CDS_0082
CDS_0083
CDS_0084
TemPhD_cluster_55459 63296:76833(+)
CDS_0072
CDS_0074
CDS_0075
CDS_0076
PD-T4-5__PD-T4-5 (CDS_0077)
IMGVR_UViG_3300014518_000014|3300014518|Ga0120821_100009|178434-254545 59246:72816(+)
56
57
58
59
60
61
PD-T4-5__PD-T4-5 (CDS_0062)
IMGVR_UViG_2547132302_000007|2547132302|2547220682|55530-127694 56707:68869(+)
CDS_0073
CDS_0074
CDS_0075
CDS_0077
CDS_0081
IMGVR_UViG_2565957055_000005|2565957055|2566034199|33796-110484 59923:73930(+)
CDS_0072
CDS_0077
CDS_0078
CDS_0080
peptidase HslV family (CDS_0081)
IMGVR_UViG_3300011631_000004|3300011631|Ga0120800_100024|64582-142363 61056:73884(+)
CDS_0070
CDS_0071
IMGVR_UViG_3300011707_000014|3300011707|Ga0120931_100024|63711-138407 57972:71401(+)
CDS_0080
IMGVR_UViG_2657245486_000004|2657245486|2657305692|166814-241089 57991:70980(+)
CDS_0056
IMGVR_UViG_2579778743_000003|2579778743|2579808949|2-66512 52567:62613(+)
CDS_0054
CDS_0056
CDS_0057
CDS_0058
IMGVR_UViG_2609460168_000003|2609460168|2609539321 51779:62935(+)
CDS_0076
IMGVR_UViG_2565956931_000003|2565956931|2566017034|26174-104914 62286:73417(+)
CDS_0045
IMGVR_UViG_2551306324_000008|2551306324|2551378849|561-62668 45609:56784(+)
CDS_0044
CDS_0049
IMGVR_UViG_2724679452_000008|2724679452|2724733394|1-63446 47327:58122(+)
CDS_0072
IMGVR_UViG_2582580790_000010|2582580790|2582638049|35-77688 58723:72330(+)
CDS_0063
transposase (CDS_0064)
CDS_0068
IMGVR_UViG_3300011894_000044|3300011894|Ga0122084_100030|8063-84976 55357:71590(+)
CDS_0067
IMGVR_UViG_2648501161_000005|2648501161|2648506399|2-75689 56812:70364(+)
anti-repressor Ant (CDS_0056)
54
55
57
peptidase HslV family (CDS_0058)
IMGVR_UViG_3300011548_000031|3300011548|Ga0120970_100021|1912-66818 47006:61611(+)
6000 nt
DNA, RNA and nucleotide metabolism
integration and excision
other
tail
transcription regulation
unknown function
Defence