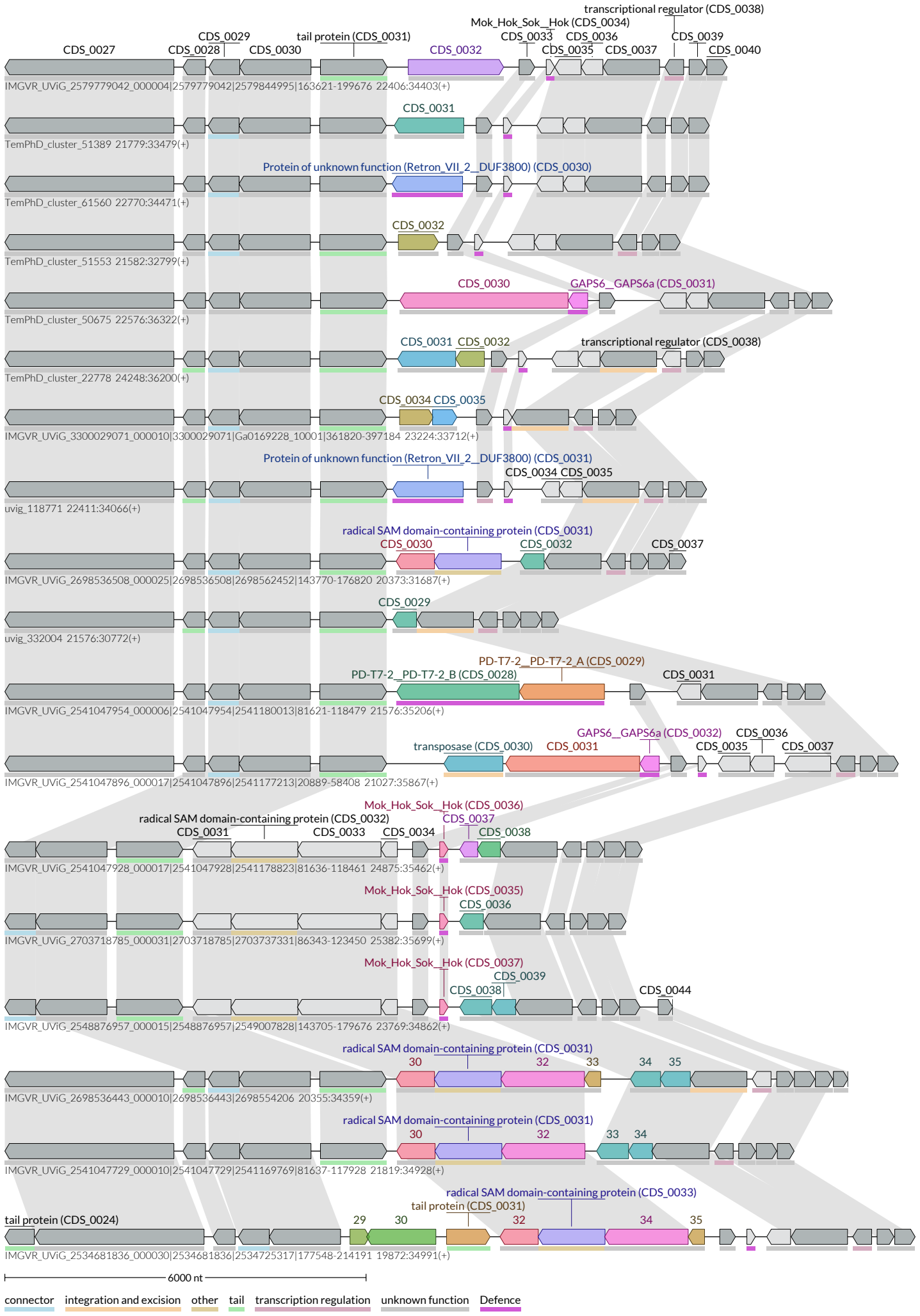
transcriptional regulator (CDS_0038)
Mok_Hok_Sok__Hok (CDS_0034)
CDS_0029
tail protein (CDS_0031)
CDS_0033
CDS_0036
CDS_0039
CDS_0027
CDS_0028
CDS_0030
CDS_0032
CDS_0035
CDS_0037
CDS_0040
IMGVR_UViG_2579779042_000004|2579779042|2579844995|163621-199676 22406:34403(+)
CDS_0031
TemPhD_cluster_51389 21779:33479(+)
Protein of unknown function (Retron_VII_2__DUF3800) (CDS_0030)
TemPhD_cluster_61560 22770:34471(+)
CDS_0032
TemPhD_cluster_51553 21582:32799(+)
CDS_0030
GAPS6__GAPS6a (CDS_0031)
TemPhD_cluster_50675 22576:36322(+)
CDS_0031
CDS_0032
transcriptional regulator (CDS_0038)
TemPhD_cluster_22778 24248:36200(+)
CDS_0034
CDS_0035
IMGVR_UViG_3300029071_000010|3300029071|Ga0169228_10001|361820-397184 23224:33712(+)
Protein of unknown function (Retron_VII_2__DUF3800) (CDS_0031)
CDS_0034
CDS_0035
uvig_118771 22411:34066(+)
radical SAM domain-containing protein (CDS_0031)
CDS_0030
CDS_0032
CDS_0037
IMGVR_UViG_2698536508_000025|2698536508|2698562452|143770-176820 20373:31687(+)
CDS_0029
uvig_332004 21576:30772(+)
PD-T7-2__PD-T7-2_A (CDS_0029)
PD-T7-2__PD-T7-2_B (CDS_0028)
CDS_0031
IMGVR_UViG_2541047954_000006|2541047954|2541180013|81621-118479 21576:35206(+)
GAPS6__GAPS6a (CDS_0032)
CDS_0036
transposase (CDS_0030)
CDS_0031
CDS_0035
CDS_0037
IMGVR_UViG_2541047896_000017|2541047896|2541177213|20889-58408 21027:35867(+)
Mok_Hok_Sok__Hok (CDS_0036)
radical SAM domain-containing protein (CDS_0032)
CDS_0037
CDS_0031
CDS_0033
CDS_0034
CDS_0038
IMGVR_UViG_2541047928_000017|2541047928|2541178823|81636-118461 24875:35462(+)
Mok_Hok_Sok__Hok (CDS_0035)
CDS_0036
IMGVR_UViG_2703718785_000031|2703718785|2703737331|86343-123450 25382:35699(+)
Mok_Hok_Sok__Hok (CDS_0037)
CDS_0039
CDS_0038
CDS_0044
IMGVR_UViG_2548876957_000015|2548876957|2549007828|143705-179676 23769:34862(+)
radical SAM domain-containing protein (CDS_0031)
30
32
33
34
35
IMGVR_UViG_2698536443_000010|2698536443|2698554206 20355:34359(+)
radical SAM domain-containing protein (CDS_0031)
30
32
33
34
IMGVR_UViG_2541047729_000010|2541047729|2541169769|81637-117928 21819:34928(+)
radical SAM domain-containing protein (CDS_0033)
tail protein (CDS_0031)
tail protein (CDS_0024)
29
30
32
34
35
IMGVR_UViG_2534681836_000030|2534681836|2534725317|177548-214191 19872:34991(+)
6000 nt
connector
integration and excision
other
tail
transcription regulation
unknown function
Defence