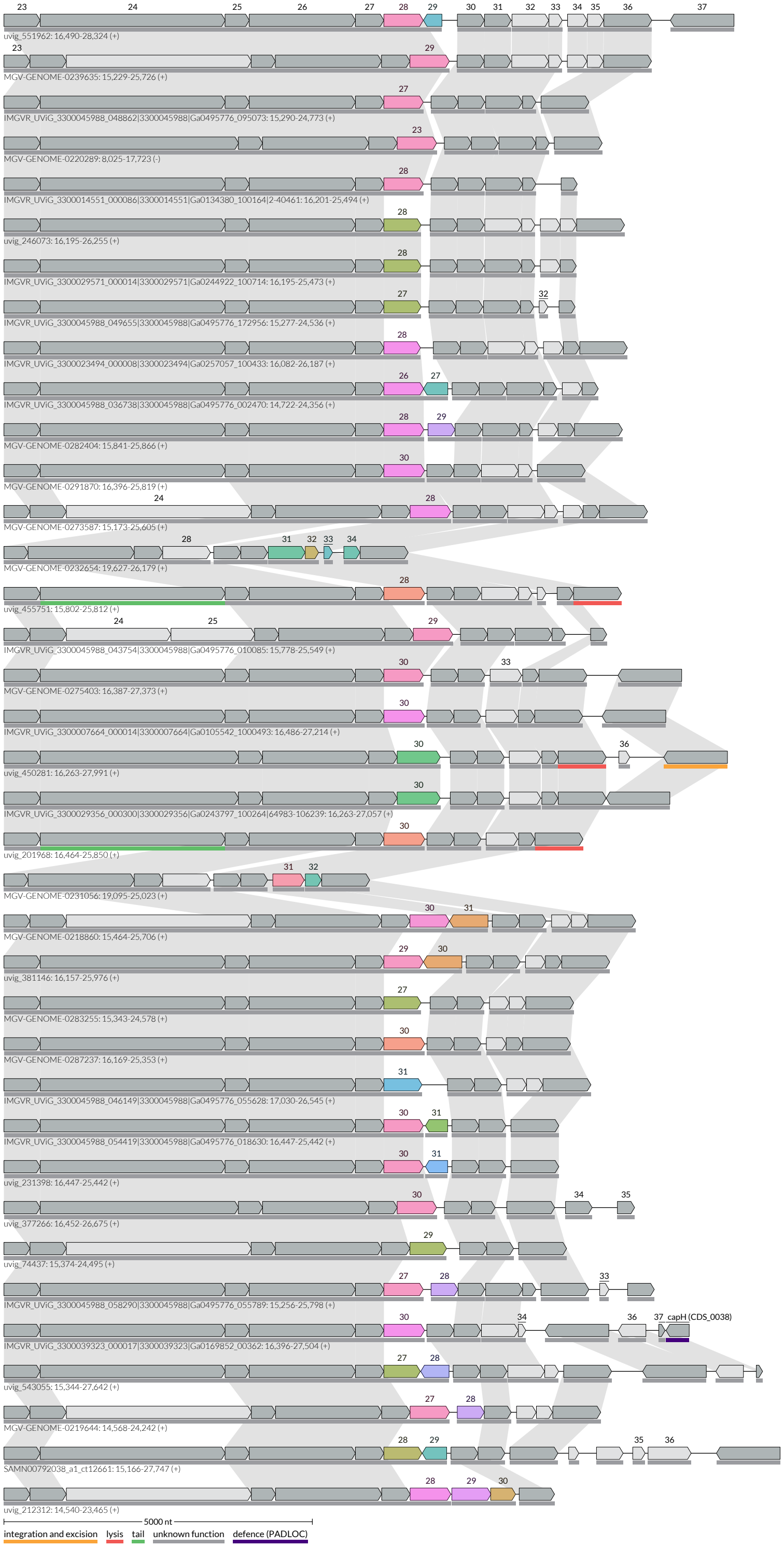
uvig_551962: 16,490-28,324 (+)
MGV-GENOME-0239635: 15,229-25,726 (+)
IMGVR_UViG_3300045988_048862|3300045988|Ga0495776_095073: 15,290-24,773 (+)
MGV-GENOME-0220289: 8,025-17,723 (-)
IMGVR_UViG_3300014551_000086|3300014551|Ga0134380_100164|2-40461: 16,201-25,494 (+)
uvig_246073: 16,195-26,255 (+)
IMGVR_UViG_3300029571_000014|3300029571|Ga0244922_100714: 16,195-25,473 (+)
IMGVR_UViG_3300045988_049655|3300045988|Ga0495776_172956: 15,277-24,536 (+)
IMGVR_UViG_3300023494_000008|3300023494|Ga0257057_100433: 16,082-26,187 (+)
IMGVR_UViG_3300045988_036738|3300045988|Ga0495776_002470: 14,722-24,356 (+)
MGV-GENOME-0282404: 15,841-25,866 (+)
MGV-GENOME-0291870: 16,396-25,819 (+)
MGV-GENOME-0273587: 15,173-25,605 (+)
MGV-GENOME-0232654: 19,627-26,179 (+)
uvig_455751: 15,802-25,812 (+)
IMGVR_UViG_3300045988_043754|3300045988|Ga0495776_010085: 15,778-25,549 (+)
MGV-GENOME-0275403: 16,387-27,373 (+)
IMGVR_UViG_3300007664_000014|3300007664|Ga0105542_1000493: 16,486-27,214 (+)
uvig_450281: 16,263-27,991 (+)
IMGVR_UViG_3300029356_000300|3300029356|Ga0243797_100264|64983-106239: 16,263-27,057 (+)
uvig_201968: 16,464-25,850 (+)
MGV-GENOME-0231056: 19,095-25,023 (+)
MGV-GENOME-0218860: 15,464-25,706 (+)
uvig_381146: 16,157-25,976 (+)
MGV-GENOME-0283255: 15,343-24,578 (+)
MGV-GENOME-0287237: 16,169-25,353 (+)
IMGVR_UViG_3300045988_046149|3300045988|Ga0495776_055628: 17,030-26,545 (+)
IMGVR_UViG_3300045988_054419|3300045988|Ga0495776_018630: 16,447-25,442 (+)
uvig_231398: 16,447-25,442 (+)
uvig_377266: 16,452-26,675 (+)
uvig_74437: 15,374-24,495 (+)
IMGVR_UViG_3300045988_058290|3300045988|Ga0495776_055789: 15,256-25,798 (+)
IMGVR_UViG_3300039323_000017|3300039323|Ga0169852_00362: 16,396-27,504 (+)
37_capH (CDS_0038)
uvig_543055: 15,344-27,642 (+)
MGV-GENOME-0219644: 14,568-24,242 (+)
SAMN00792038_a1_ct12661: 15,166-27,747 (+)
uvig_212312: 14,540-23,465 (+)
5000 nt
integration and excision
lysis
tail
unknown function
defence (PADLOC)