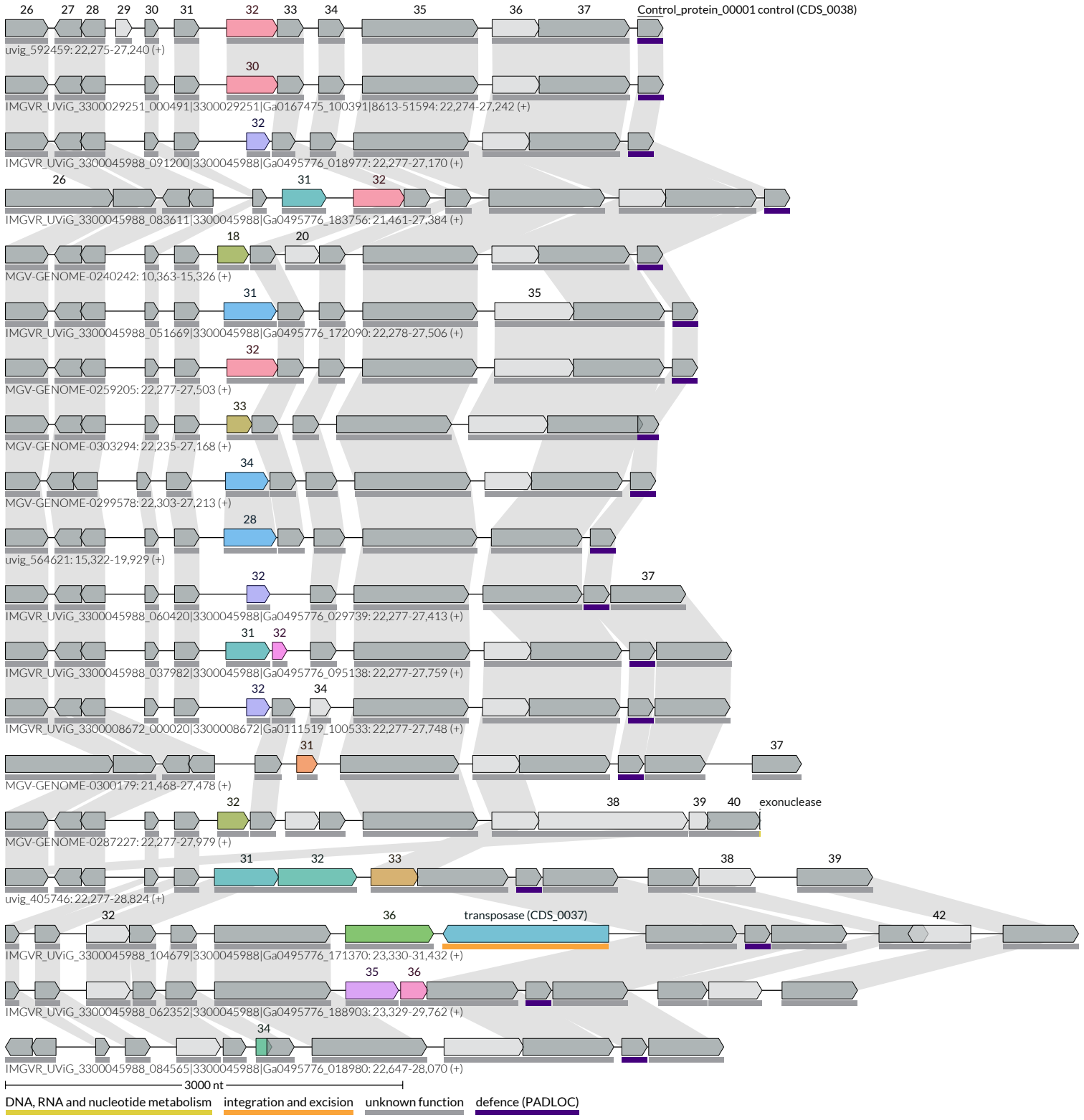
26 27 28 29 30 31 32 33 34 35 36 37 Control_protein_00001 control (CDS_0038)
uvig_592459: 22,275-27,240 (+)
30
IMGVR_UViG_3300029251_000491|3300029251|Ga0167475_100391|8613-51594: 22,274-27,242 (+)
32
IMGVR_UViG_3300045988_091200|3300045988|Ga0495776_018977: 22,277-27,170 (+)
26 31 32
IMGVR_UViG_3300045988_083611|3300045988|Ga0495776_183756: 21,461-27,384 (+)
18 20
MGV-GENOME-0240242: 10,363-15,326 (+)
31 35
IMGVR_UViG_3300045988_051669|3300045988|Ga0495776_172090: 22,278-27,506 (+)
32
MGV-GENOME-0259205: 22,277-27,503 (+)
33
MGV-GENOME-0303294: 22,235-27,168 (+)
34
MGV-GENOME-0299578: 22,303-27,213 (+)
28
uvig_564621: 15,322-19,929 (+)
32 37
IMGVR_UViG_3300045988_060420|3300045988|Ga0495776_029739: 22,277-27,413 (+)
31 32
IMGVR_UViG_3300045988_037982|3300045988|Ga0495776_095138: 22,277-27,759 (+)
32 34
IMGVR_UViG_3300008672_000020|3300008672|Ga0111519_100533: 22,277-27,748 (+)
31 37
MGV-GENOME-0300179: 21,468-27,478 (+)
32 38 39 40 exonuclease
MGV-GENOME-0287227: 22,277-27,979 (+)
31 32 33 38 39
uvig_405746: 22,277-28,824 (+)
32 36 transposase (CDS_0037) 42
IMGVR_UViG_3300045988_104679|3300045988|Ga0495776_171370: 23,330-31,432 (+)
35 36
IMGVR_UViG_3300045988_062352|3300045988|Ga0495776_188903: 23,329-29,762 (+)
34
IMGVR_UViG_3300045988_084565|3300045988|Ga0495776_018980: 22,647-28,070 (+)
3000 nt
DNA, RNA and nucleotide metabolism
integration and excision
unknown function
defence (PADLOC)