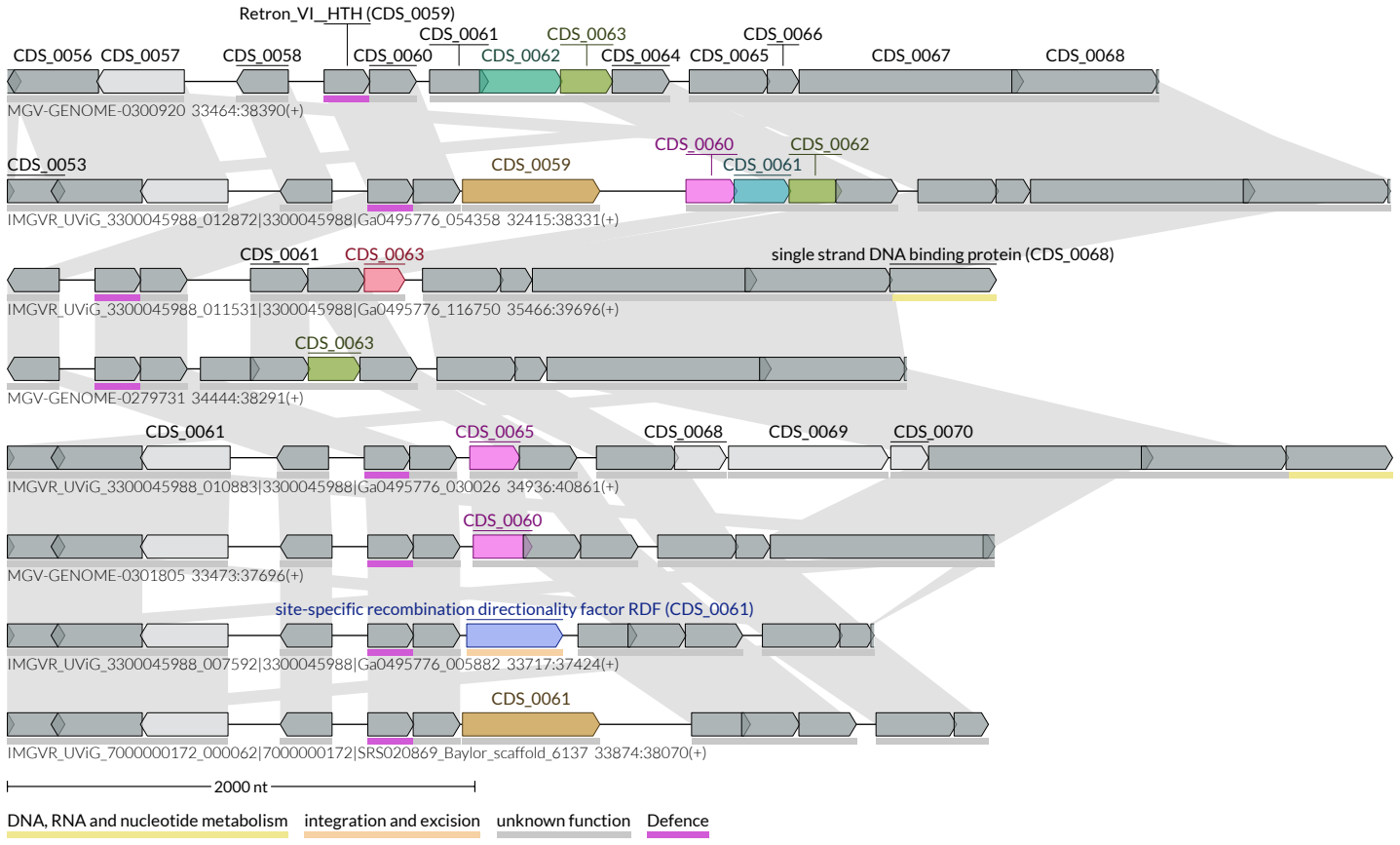
Retron_VI__HTH (CDS_0059)
CDS_0056
CDS_0057
CDS_0058
CDS_0060
CDS_0061
CDS_0062
CDS_0063
CDS_0064
CDS_0065
CDS_0066
CDS_0067
CDS_0068
MGV-GENOME-0300920 33464:38390(+)
CDS_0053
CDS_0059
CDS_0060
CDS_0061
CDS_0062
IMGVR_UViG_3300045988_012872|3300045988|Ga0495776_054358 32415:38331(+)
CDS_0061
CDS_0063
single strand DNA binding protein (CDS_0068)
IMGVR_UViG_3300045988_011531|3300045988|Ga0495776_116750 35466:39696(+)
CDS_0063
MGV-GENOME-0279731 34444:38291(+)
CDS_0061
CDS_0065
CDS_0068
CDS_0069
CDS_0070
IMGVR_UViG_3300045988_010883|3300045988|Ga0495776_030026 34936:40861(+)
CDS_0060
MGV-GENOME-0301805 33473:37696(+)
site-specific recombination directionality factor RDF (CDS_0061)
IMGVR_UViG_3300045988_007592|3300045988|Ga0495776_005882 33717:37424(+)
CDS_0061
IMGVR_UViG_7000000172_000062|7000000172|SRS020869_Baylor_scaffold_6137 33874:38070(+)
2000 nt
DNA, RNA and nucleotide metabolism
integration and excision
unknown function
Defence