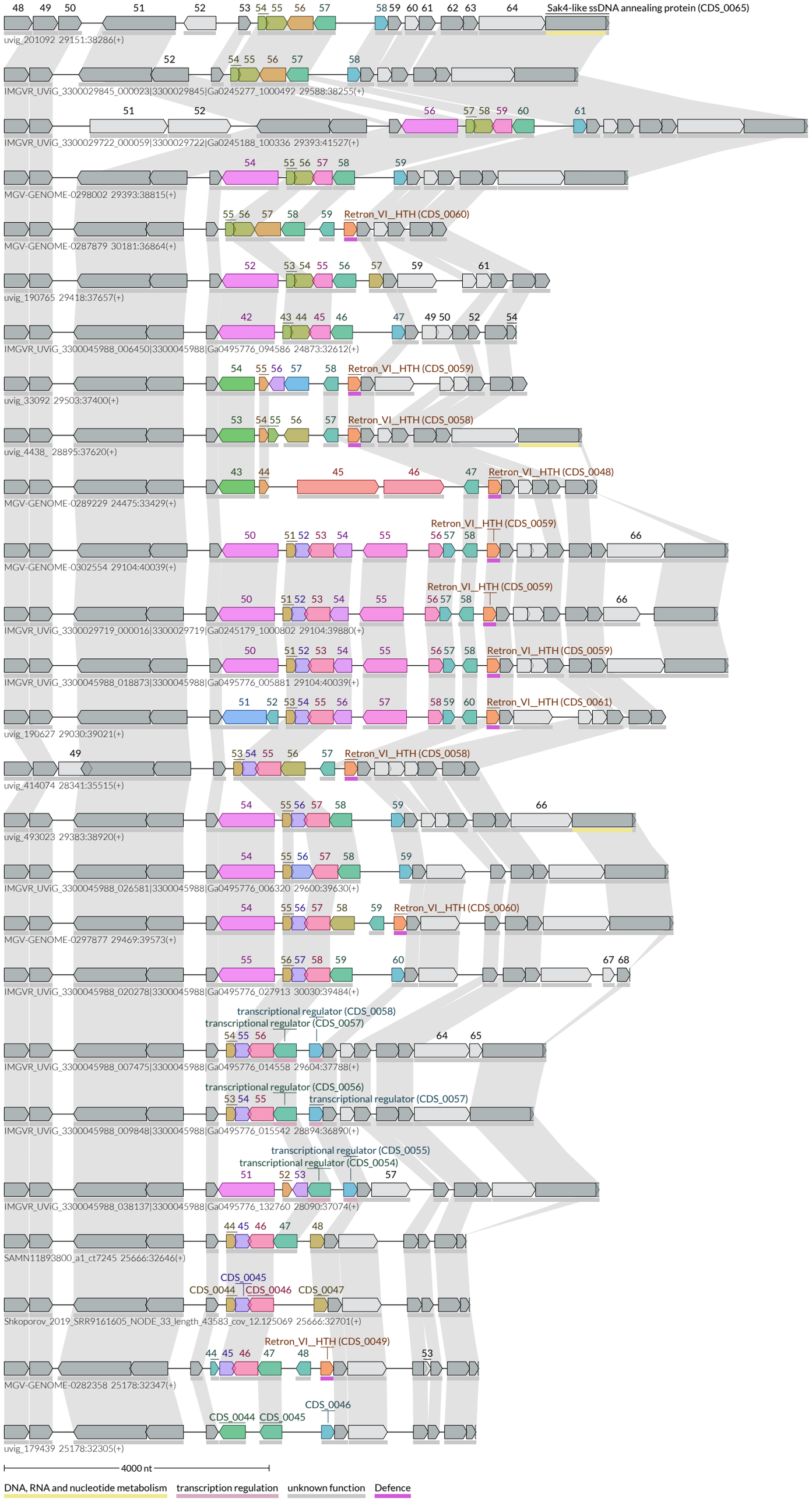
Sak4-like ssDNA annealing protein (CDS_0065)
uvig_201092 29151:38286(+)
IMGVR_UViG_3300029845_000023|3300029845|Ga0245277_1000492 29588:38255(+)
IMGVR_UViG_3300029722_000059|3300029722|Ga0245188_100336 29393:41527(+)
MGV-GENOME-0298002 29393:38815(+)
Retron_VI_HTH (CDS_0060)
MGV-GENOME-0287879 30181:36864(+)
uvig_190765 29418:37657(+)
IMGVR_UViG_3300045988_006450|3300045988|Ga0495776_094586 24873:32612(+)
Retron_VI_HTH (CDS_0059)
uvig_33092 29503:37400(+)
Retron_VI_HTH (CDS_0058)
uvig_4438_ 28895:37620(+)
Retron_VI_HTH (CDS_0048)
MGV-GENOME-0289229 24475:33429(+)
MGV-GENOME-0302554 29104:40039(+)
IMGVR_UViG_3300029719_000016|3300029719|Ga0245179_1000802 29104:39880(+)
IMGVR_UViG_3300045988_018873|3300045988|Ga0495776_005881 29104:40039(+)
Retron_VI_HTH (CDS_0061)
uvig_190627 29030:39021(+)
uvig_414074 28341:35515(+)
uvig_493023 29383:38920(+)
IMGVR_UViG_3300045988_026581|3300045988|Ga0495776_006320 29600:39630(+)
MGV-GENOME-0297877 29469:39573(+)
IMGVR_UViG_3300045988_020278|3300045988|Ga0495776_027913 30030:39484(+)
transcriptional regulator (CDS_0058)
transcriptional regulator (CDS_0057)
IMGVR_UViG_3300045988_007475|3300045988|Ga0495776_014558 29604:37788(+)
transcriptional regulator (CDS_0056)
IMGVR_UViG_3300045988_009848|3300045988|Ga0495776_015542 28894:36890(+)
transcriptional regulator (CDS_0055)
transcriptional regulator (CDS_0054)
IMGVR_UViG_3300045988_038137|3300045988|Ga0495776_132760 28090:37074(+)
SAMN11893800_a1_ct7245 25666:32646(+)
CDS_0045
CDS_0044
CDS_0046
CDS_0047
Shkoporov_2019_SRR9161605_NODE_33_length_43583_cov_12.125069 25666:32701(+)
Retron_VI_HTH (CDS_0049)
MGV-GENOME-0282358 25178:32347(+)
CDS_0046
CDS_0044
CDS_0045
uvig_179439 25178:32305(+)
4000 nt
DNA, RNA and nucleotide metabolism
transcription regulation
unknown function
Defence