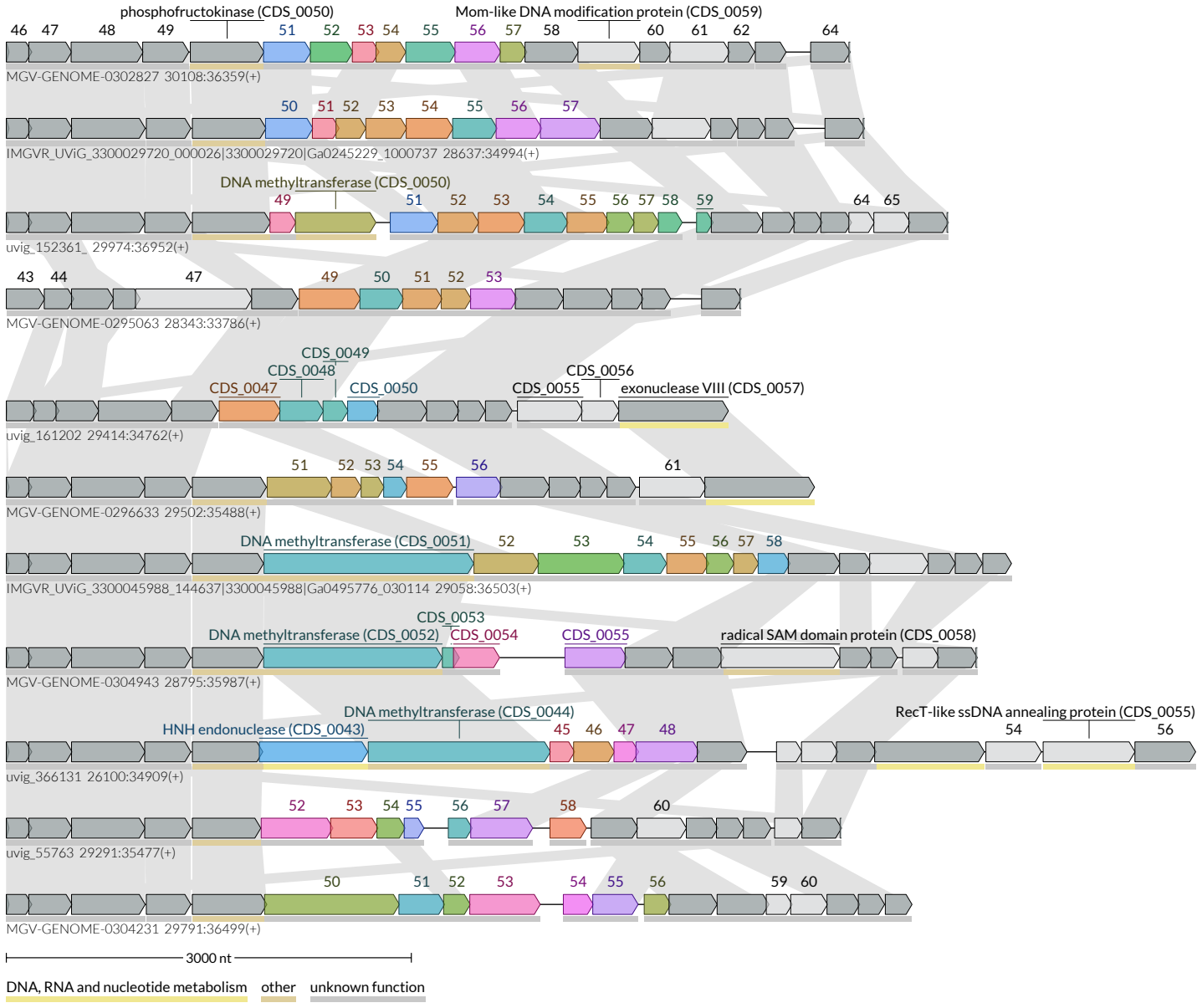
phosphofructokinase (CDS_0050)
Mom-like DNA modification protein (CDS_0059)
46 47 48 49 51 52 53 54 55 56 57 58 60 61 62 64
MGV-GENOME-0302827 30108:36359(+)
50 51 52 53 54 55 56 57
IMGVR_UViG_3300029720_000026|3300029720|Ga0245229_1000737 28637:34994(+)
DNA methyltransferase (CDS_0050)
49 51 52 53 54 55 56 57 58 59 64 65
uvig_152361_ 29974:36952(+)
43 44 47 49 50 51 52 53
MGV-GENOME-0295063 28343:33786(+)
CDS_0049
CDS_0048
CDS_0056
CDS_0047
CDS_0050
CDS_0055
exonuclease VIII (CDS_0057)
uvig_161202 29414:34762(+)
51 52 53 54 55 56 61
MGV-GENOME-0296633 29502:35488(+)
DNA methyltransferase (CDS_0051)
52 53 54 55 56 57 58
IMGVR_UViG_3300045988_144637|3300045988|Ga0495776_030114 29058:36503(+)
CDS_0053
DNA methyltransferase (CDS_0052)
CDS_0054
CDS_0055
radical SAM domain protein (CDS_0058)
MGV-GENOME-0304943 28795:35987(+)
DNA methyltransferase (CDS_0044)
RecT-like ssDNA annealing protein (CDS_0055)
HNH endonuclease (CDS_0043)
45 46 47 48 54 56
uvig_366131 26100:34909(+)
52 53 54 55 56 57 58 60
uvig_55763 29291:35477(+)
50 51 52 53 54 55 56 59 60
MGV-GENOME-0304231 29791:36499(+)
3000 nt
DNA, RNA and nucleotide metabolism
other
unknown function