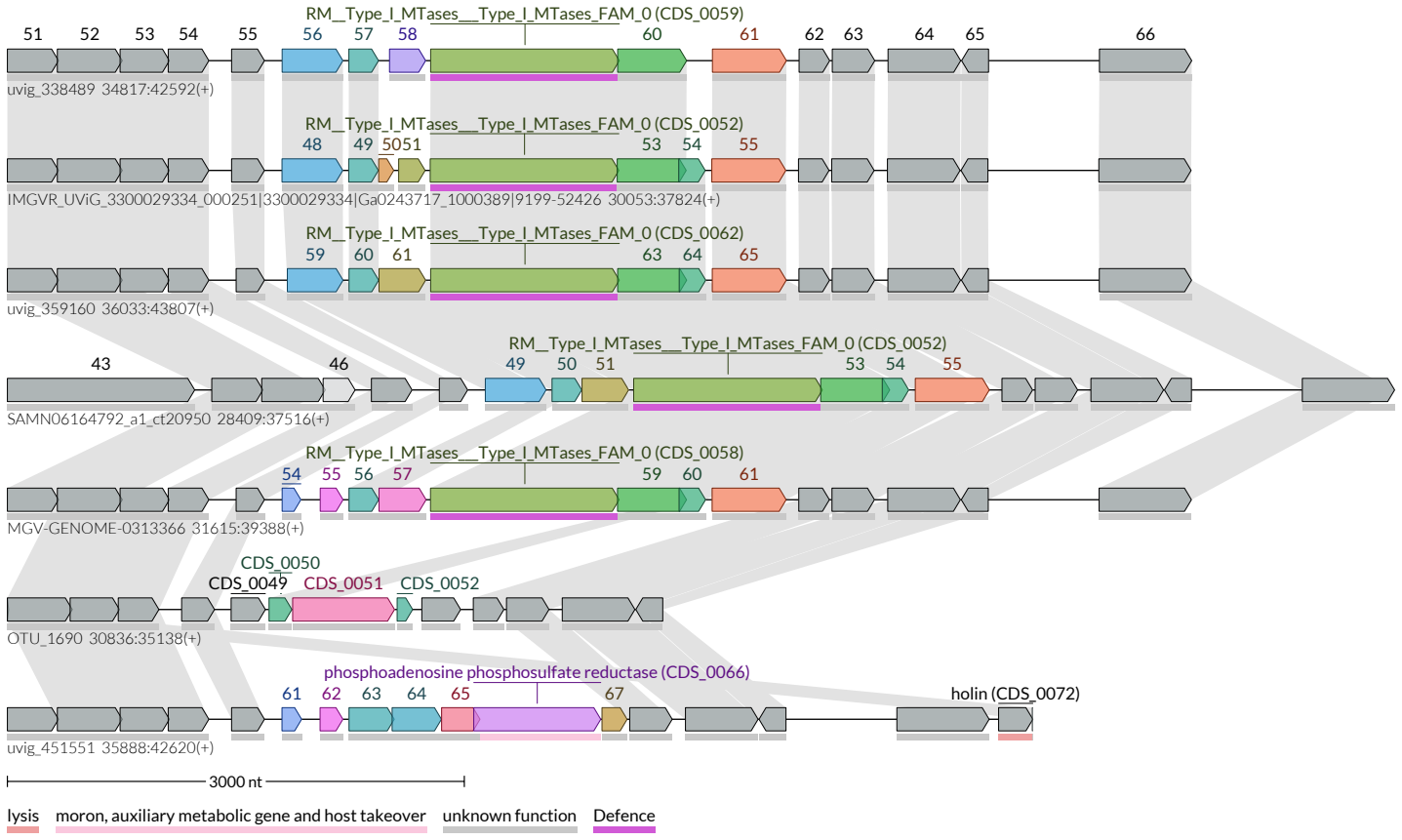
RM_Type_I_MTases___Type_I_MTases_FAM_0 (CDS_0059)
51 52 53 54 55 56 57 58 60 61 62 63 64 65 66
uvig_338489 34817:42592(+)
RM_Type_I_MTases___Type_I_MTases_FAM_0 (CDS_0052)
48 49 5051 53 54 55
IMGVR_UViG_3300029334_000251|3300029334|Ga0243717_1000389|9199-52426 30053:37824(+)
RM_Type_I_MTases___Type_I_MTases_FAM_0 (CDS_0062)
59 60 61 63 64 65
uvig_359160 36033:43807(+)
RM_Type_I_MTases___Type_I_MTases_FAM_0 (CDS_0052)
43 46 49 50 51 53 54 55
SAMN06164792_a1_ct20950 28409:37516(+)
RM_Type_I_MTases___Type_I_MTases_FAM_0 (CDS_0058)
54 55 56 57 59 60 61
MGV-GENOME-0313366 31615:39388(+)
CDS_0050
CDS_0049 CDS_0051 CDS_0052
OTU_1690 30836:35138(+)
phosphoadenosine phosphosulfate reductase (CDS_0066)
61 62 63 64 65 67
holin (CDS_0072)
uvig_451551 35888:42620(+)
3000 nt
lysis
moron, auxiliary metabolic gene and host takeover
unknown function
Defence