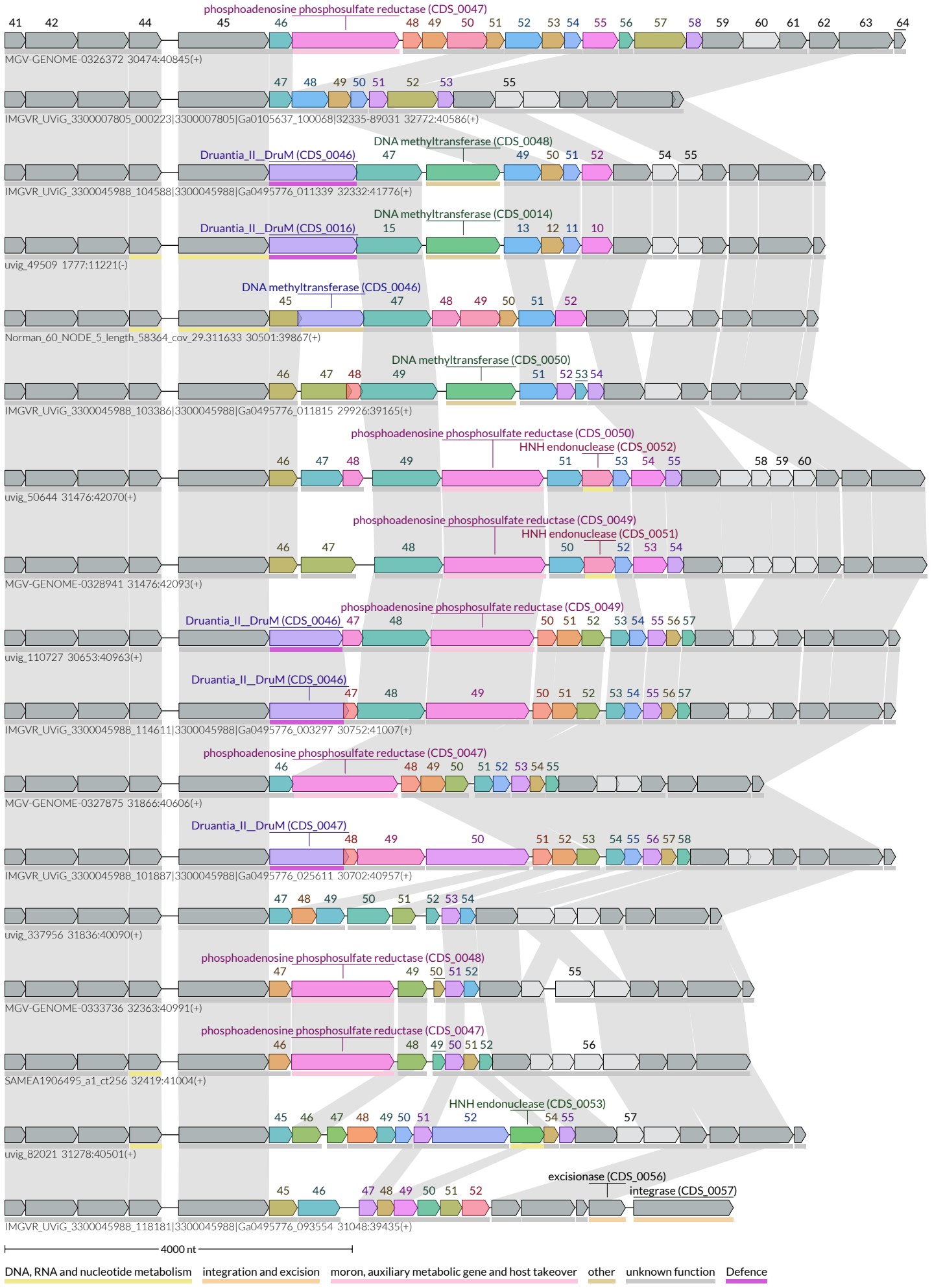
phosphoadenosine phosphosulfate reductase (CDS_0047)
MGV-GENOME-0326372 30474:40845(+)
IMGVR_UViG_3300007805_000223|3300007805|Ga0105637_100068|32335-89031 32772:40586(+)
DNA methyltransferase (CDS_0048)
Druantia_II__DruM (CDS_0046)
IMGVR_UViG_3300045988_104588|3300045988|Ga0495776_011339 32332:41776(+)
DNA methyltransferase (CDS_0014)
Druantia_II__DruM (CDS_0016)
uvig_49509 1777:11221(-)
DNA methyltransferase (CDS_0046)
Norman_60_NODE_5_length_58364_cov_29.311633 30501:39867(+)
DNA methyltransferase (CDS_0050)
IMGVR_UViG_3300045988_103386|3300045988|Ga0495776_011815 29926:39165(+)
phosphoadenosine phosphosulfate reductase (CDS_0050)
HNH endonuclease (CDS_0052)
uvig_50644 31476:42070(+)
phosphoadenosine phosphosulfate reductase (CDS_0049)
HNH endonuclease (CDS_0051)
MGV-GENOME-0328941 31476:42093(+)
phosphoadenosine phosphosulfate reductase (CDS_0049)
Druantia_II__DruM (CDS_0046)
uvig_110727 30653:40963(+)
Druantia_II__DruM (CDS_0046)
IMGVR_UViG_3300045988_114611|3300045988|Ga0495776_003297 30752:41007(+)
phosphoadenosine phosphosulfate reductase (CDS_0047)
MGV-GENOME-0327875 31866:40606(+)
Druantia_II__DruM (CDS_0047)
IMGVR_UViG_3300045988_101887|3300045988|Ga0495776_025611 30702:40957(+)
uvig_337956 31836:40090(+)
phosphoadenosine phosphosulfate reductase (CDS_0048)
MGV-GENOME-0333736 32363:40991(+)
phosphoadenosine phosphosulfate reductase (CDS_0047)
SAMEA1906495_a1_ct256 32419:41004(+)
HNH endonuclease (CDS_0053)
uvig_82021 31278:40501(+)
excisionase (CDS_0056)
integrase (CDS_0057)
IMGVR_UViG_3300045988_118181|3300045988|Ga0495776_093554 31048:39435(+)
4000 nt
DNA, RNA and nucleotide metabolism
integration and excision
moron, auxiliary metabolic gene and host takeover
other
unknown function
Defence