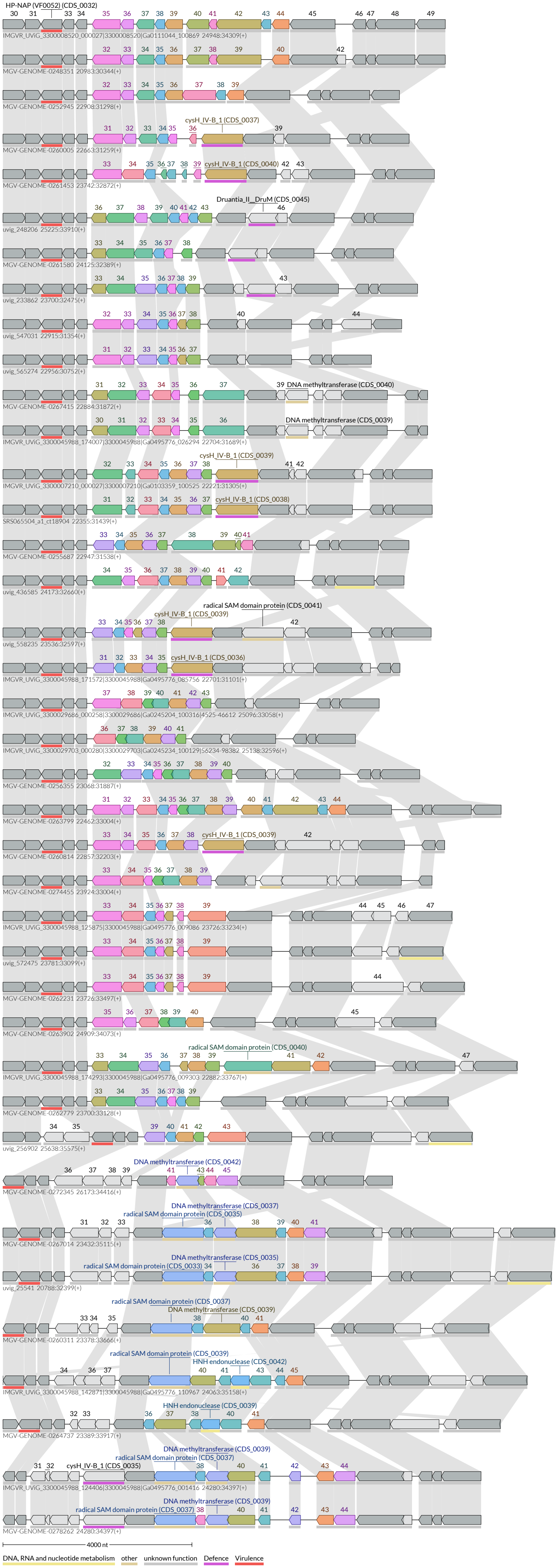
HP-NAP (VF0052) (CDS_0032)
IMGVR_UViG_3300008520_000027|3300008520|Ga0111044_100869 24948:34309(+)
MGV-GENOME-0248351 20983:30344(+)
MGV-GENOME-0252945 22908:31298(+)
cysH_IV-B_1 (CDS_0037)
MGV-GENOME-0260005 22663:31259(+)
cysH_IV-B_1 (CDS_0040)
MGV-GENOME-0261453 23742:32872(+)
Druantia_II__DruM (CDS_0045)
uvig_248206 25225:33910(+)
MGV-GENOME-0261580 24125:32389(+)
uvig_233862 23700:32475(+)
uvig_547031 22915:31354(+)
uvig_565274 22956:30752(+)
DNA methyltransferase (CDS_0040)
MGV-GENOME-0267415 22884:31872(+)
DNA methyltransferase (CDS_0039)
IMGVR_UViG_3300045988_174007|3300045988|Ga0495776_026294 22704:31689(+)
cysH_IV-B_1 (CDS_0039)
IMGVR_UViG_3300007210_000027|3300007210|Ga0103359_100525 22221:31305(+)
cysH_IV-B_1 (CDS_0038)
SRS065504_a1_ct18904 22355:31439(+)
MGV-GENOME-0255687 22947:31538(+)
uvig_436585 24173:32660(+)
radical SAM domain protein (CDS_0041)
cysH_IV-B_1 (CDS_0039)
uvig_558235 23536:32597(+)
cysH_IV-B_1 (CDS_0036)
IMGVR_UViG_3300045988_171572|3300045988|Ga0495776_085756 22701:31101(+)
IMGVR_UViG_3300029686_000258|3300029686|Ga0245204_100316|4525-46612 25096:33058(+)
IMGVR_UViG_3300029703_000280|3300029703|Ga0245234_100129|56234-98382 25138:32596(+)
MGV-GENOME-0256355 23068:31887(+)
MGV-GENOME-0263799 22462:33004(+)
MGV-GENOME-0260814 22857:32203(+)
MGV-GENOME-0274455 23924:33004(+)
IMGVR_UViG_3300045988_125875|3300045988|Ga0495776_009086 23726:33234(+)
uvig_572475 23781:33099(+)
MGV-GENOME-0262231 23726:33497(+)
MGV-GENOME-0263902 24909:34073(+)
radical SAM domain protein (CDS_0040)
IMGVR_UViG_3300045988_174293|3300045988|Ga0495776_009303 22882:33767(+)
MGV-GENOME-0262779 23700:33128(+)
uvig_256902 25638:35575(+)
DNA methyltransferase (CDS_0042)
MGV-GENOME-0272345 26173:34416(+)
DNA methyltransferase (CDS_0037)
radical SAM domain protein (CDS_0035)
MGV-GENOME-0267014 23432:35115(+)
DNA methyltransferase (CDS_0035)
radical SAM domain protein (CDS_0033)
uvig_25541 20788:32399(+)
radical SAM domain protein (CDS_0037)
DNA methyltransferase (CDS_0039)
MGV-GENOME-0260311 23378:33666(+)
radical SAM domain protein (CDS_0039)
HNH endonuclease (CDS_0042)
IMGVR_UViG_3300045988_142871|3300045988|Ga0495776_110967 24063:35158(+)
HNH endonuclease (CDS_0039)
MGV-GENOME-0264737 23389:33917(+)
DNA methyltransferase (CDS_0039)
radical SAM domain protein (CDS_0037)
cysH_IV-B_1 (CDS_0035)
IMGVR_UViG_3300045988_124406|3300045988|Ga0495776_001416 24280:34397(+)
DNA methyltransferase (CDS_0039)
radical SAM domain protein (CDS_0037)
MGV-GENOME-0278262 24280:34397(+)
4000 nt
DNA, RNA and nucleotide metabolism
other
unknown function
Defence
Virulence