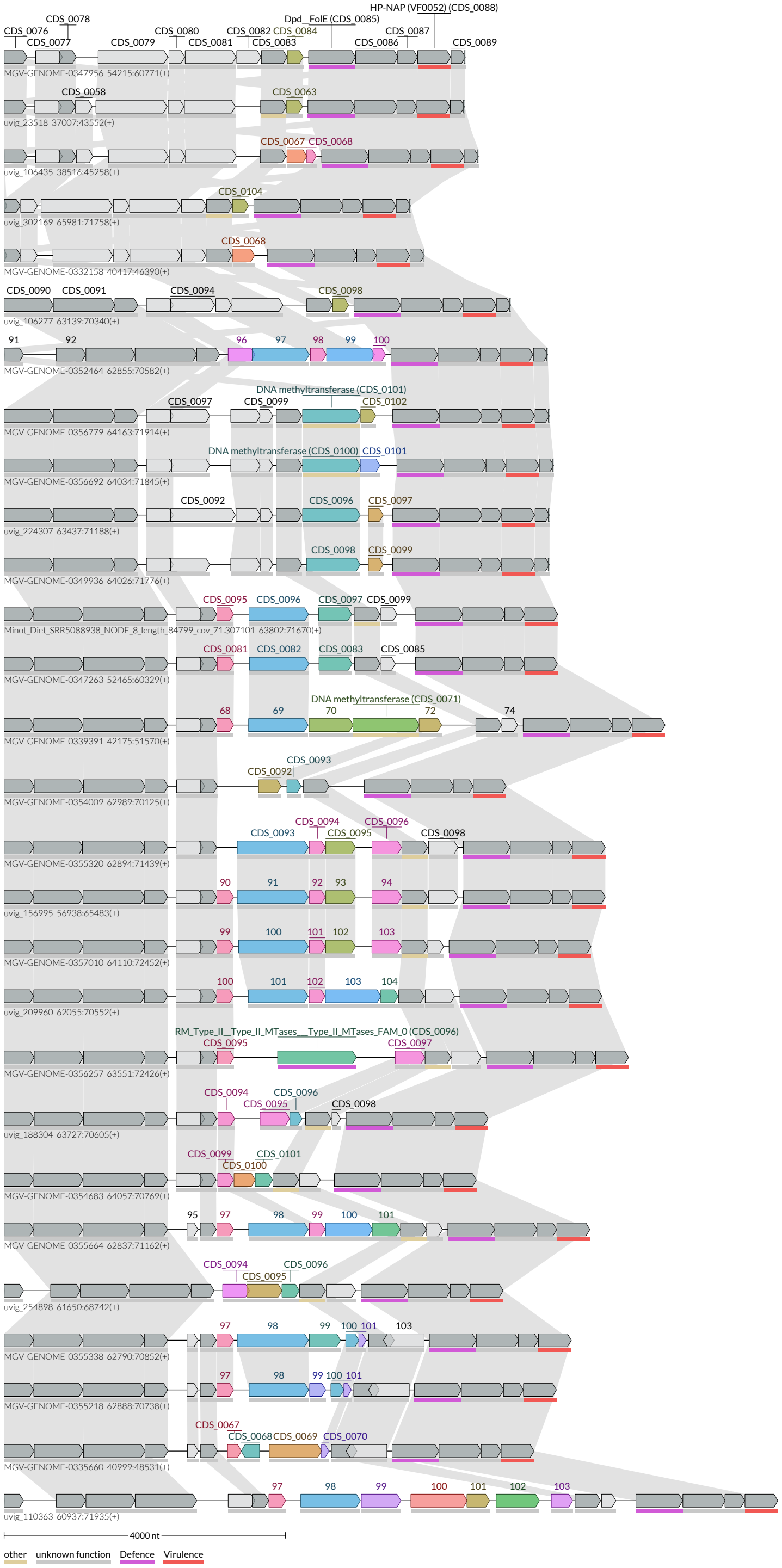
HP-NAP (VF0052) (CDS_0088)
CDS_0078
Dpd__FolE (CDS_0085)
CDS_0076
CDS_0080
CDS_0082
CDS_0084
CDS_0087
CDS_0077
CDS_0079
CDS_0081
CDS_0083
CDS_0086
CDS_0089
MGV-GENOME-0347956 54215:60771(+)
CDS_0058
CDS_0063
uvig_23518 37007:43552(+)
CDS_0067
CDS_0068
uvig_106435 38516:45258(+)
CDS_0104
uvig_302169 65981:71758(+)
CDS_0068
MGV-GENOME-0332158 40417:46390(+)
CDS_0090
CDS_0091
CDS_0094
CDS_0098
uvig_106277 63139:70340(+)
91
92
96
97
98
99
100
MGV-GENOME-0352464 62855:70582(+)
DNA methyltransferase (CDS_0101)
CDS_0097
CDS_0099
CDS_0102
MGV-GENOME-0356779 64163:71914(+)
DNA methyltransferase (CDS_0100)
CDS_0101
MGV-GENOME-0356692 64034:71845(+)
CDS_0092
CDS_0096
CDS_0097
uvig_224307 63437:71188(+)
CDS_0098
CDS_0099
MGV-GENOME-0349936 64026:71776(+)
CDS_0095
CDS_0096
CDS_0097
CDS_0099
Minot_Diet_SRR5088938_NODE_8_length_84799_cov_71.307101 63802:71670(+)
CDS_0081
CDS_0082
CDS_0083
CDS_0085
MGV-GENOME-0347263 52465:60329(+)
DNA methyltransferase (CDS_0071)
68
69
70
72
74
MGV-GENOME-0339391 42175:51570(+)
CDS_0093
CDS_0092
MGV-GENOME-0354009 62989:70125(+)
CDS_0094
CDS_0096
CDS_0093
CDS_0095
CDS_0098
MGV-GENOME-0355320 62894:71439(+)
90
91
92
93
94
uvig_156995 56938:65483(+)
99
100
101
102
103
MGV-GENOME-0357010 64110:72452(+)
100
101
102
103
104
uvig_209960 62055:70552(+)
RM_Type_II__Type_II_MTases___Type_II_MTases_FAM_0 (CDS_0096)
CDS_0095
CDS_0097
MGV-GENOME-0356257 63551:72426(+)
CDS_0094
CDS_0096
CDS_0095
CDS_0098
uvig_188304 63727:70605(+)
CDS_0099
CDS_0101
CDS_0100
MGV-GENOME-0354683 64057:70769(+)
95
97
98
99
100
101
MGV-GENOME-0355664 62837:71162(+)
CDS_0094
CDS_0096
CDS_0095
uvig_254898 61650:68742(+)
97
98
99
100
101
103
MGV-GENOME-0355338 62790:70852(+)
97
98
99
100
101
MGV-GENOME-0355218 62888:70738(+)
CDS_0067
CDS_0068
CDS_0069
CDS_0070
MGV-GENOME-0335660 40999:48531(+)
97
98
99
100
101
102
103
uvig_110363 60937:71935(+)
4000 nt
other
unknown function
Defence
Virulence