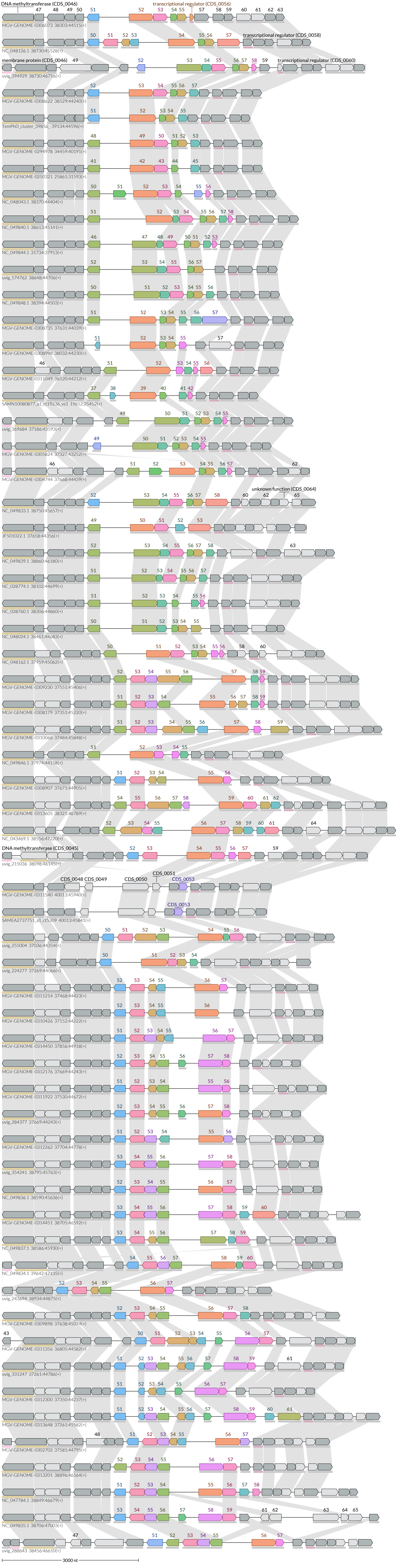
DNA methyltransferase (CDS_0046)
transcriptional regulator (CDS_0056)
MGV-GENOME-0306073 38303:44515(+)
transcriptional regulator (CDS_0058)
NC_048126.1 38730:45528(+)
membrane protein (CDS_0046)
transcriptional regulator (CDS_0060)
uvig_394929 38730:46716(+)
MGV-GENOME-0308622 38529:44240(+)
TemPhD_cluster_39856_ 39134:44596(+)
MGV-GENOME-0294978 34459:40191(+)
MGV-GENOME-0250321 25861:31593(+)
NC_048043.1 38370:44404(+)
NC_049840.1 38613:45141(+)
NC_049844.1 31734:37913(+)
uvig_574762 38648:44706(+)
NC_049848.1 38394:44503(+)
MGV-GENOME-0308735 37631:44039(+)
MGV-GENOME-0308998 38032:44230(+)
MGV-GENOME-0311049 38120:44212(+)
SAMN10080877_a1_ct19236_vs1 19812:25452(+)
uvig_369684 37186:43593(+)
MGV-GENOME-0305824 37327:43252(+)
MGV-GENOME-0304744 37668:44439(+)
unknown function (CDS_0064)
NC_049833.1 38750:45657(+)
JF501022.1 37658:44356(+)
NC_049839.1 38860:46180(+)
NC_028774.1 38102:44699(+)
NC_028760.1 38306:44860(+)
NC_048024.1 38461:44683(+)
NC_048162.1 37959:45062(+)
MGV-GENOME-0309230 37551:45406(+)
MGV-GENOME-0308179 37351:45220(+)
MGV-GENOME-0310068 37484:45848(+)
NC_049846.1 37974:44158(+)
MGV-GENOME-0308907 37671:44905(+)
MGV-GENOME-0313605 38321:46789(+)
NC_043469.1 38556:47220(+)
DNA methyltransferase (CDS_0045)
uvig_215036 38098:46145(+)
CDS_0048
CDS_0049
CDS_0050
CDS_0051
CDS_0053
MGV-GENOME-0311540 40013:45940(+)
SAMEA2737751_a1_ct5309 40013:45841(+)
uvig_255004 37036:44354(+)
uvig_224277 37269:44066(+)
MGV-GENOME-0311214 37468:44423(+)
MGV-GENOME-0310426 37152:44222(+)
MGV-GENOME-0314450 37856:44918(+)
MGV-GENOME-0312176 37669:44243(+)
MGV-GENOME-0311922 37530:44672(+)
uvig_284377 37669:44243(+)
MGV-GENOME-0312262 37704:44778(+)
uvig_354241 38795:45763(+)
NC_049836.1 38590:45636(+)
MGV-GENOME-0314451 38705:46592(+)
NC_049837.1 38586:45930(+)
NC_049834.1 39642:47135(+)
uvig_243694 38934:44875(+)
MGV-GENOME-0309898 37638:45074(+)
MGV-GENOME-0311356 36805:44582(+)
uvig_331247 37261:44786(+)
MGV-GENOME-0312300 37350:44237(+)
MGV-GENOME-0313648 37261:45582(+)
MGV-GENOME-0302702 37581:44795(+)
MGV-GENOME-0313201 38896:46164(+)
NC_047784.1 38849:46679(+)
NC_049835.1 38706:47003(+)
uvig_288643 38456:46610(+)
3000 nt
moron, auxiliary metabolic gene and host takeover
other
transcription regulation
unknown function