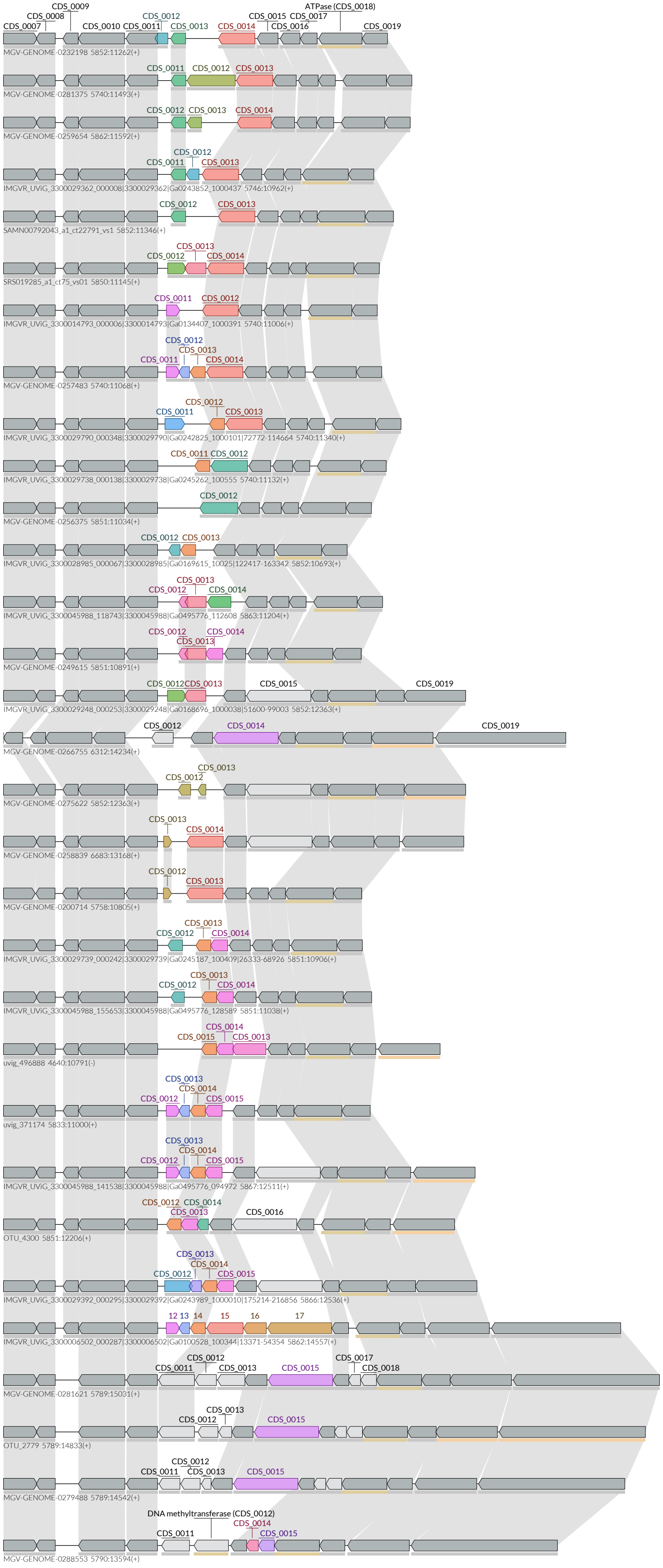
CDS_0009
CDS_0008
CDS_0007
CDS_0010
CDS_0011
CDS_0012
CDS_0013
CDS_0014
CDS_0015
CDS_0016
CDS_0017
ATPase (CDS_0018)
CDS_0019
MGV-GENOME-0232198 5852:11262(+)
MGV-GENOME-0281375 5740:11493(+)
MGV-GENOME-0259654 5862:11592(+)
IMGVR_UViG_3300029362_000008|3300029362|Ga0243852_1000437 5746:10962(+)
SAMN00792043_a1_ct22791_vs1 5852:11346(+)
SRS019285_a1_ct75_vs01 5850:11145(+)
IMGVR_UViG_3300014793_000006|3300014793|Ga0134407_1000391 5740:11006(+)
MGV-GENOME-0257483 5740:11068(+)
IMGVR_UViG_3300029790_000348|3300029790|Ga0242825_1000101|72772-114664 5740:11340(+)
IMGVR_UViG_3300029738_000138|3300029738|Ga0245262_100555 5740:11132(+)
MGV-GENOME-0256375 5851:11034(+)
IMGVR_UViG_3300028985_000067|3300028985|Ga0169615_10025|122417-163342 5852:10693(+)
IMGVR_UViG_3300045988_118743|3300045988|Ga0495776_112608 5863:11204(+)
MGV-GENOME-0249615 5851:10891(+)
IMGVR_UViG_3300029248_000253|3300029248|Ga0168696_1000038|51600-99003 5852:12363(+)
MGV-GENOME-0266755 6312:14234(+)
MGV-GENOME-0275622 5852:12363(+)
MGV-GENOME-0258839 6683:13168(+)
MGV-GENOME-0200714 5758:10805(+)
IMGVR_UViG_3300029739_000242|3300029739|Ga0245187_100409|26333-68926 5851:10906(+)
IMGVR_UViG_3300045988_155653|3300045988|Ga0495776_128589 5851:11038(+)
uvig_496888 4640:10791(-)
uvig_371174 5833:11000(+)
IMGVR_UViG_3300045988_141538|3300045988|Ga0495776_094972 5867:12511(+)
OTU_4300 5851:12206(+)
IMGVR_UViG_3300029392_000295|3300029392|Ga0243989_1000010|175214-216856 5866:12536(+)
IMGVR_UViG_3300006502_000287|3300006502|Ga0100528_100344|13371-54354 5862:14557(+)
MGV-GENOME-0281621 5789:15031(+)
OTU_2779 5789:14833(+)
MGV-GENOME-0279488 5789:14542(+)
DNA methyltransferase (CDS_0012)
MGV-GENOME-0288553 5790:13594(+)
3000 nt
integration and excision
other
unknown function