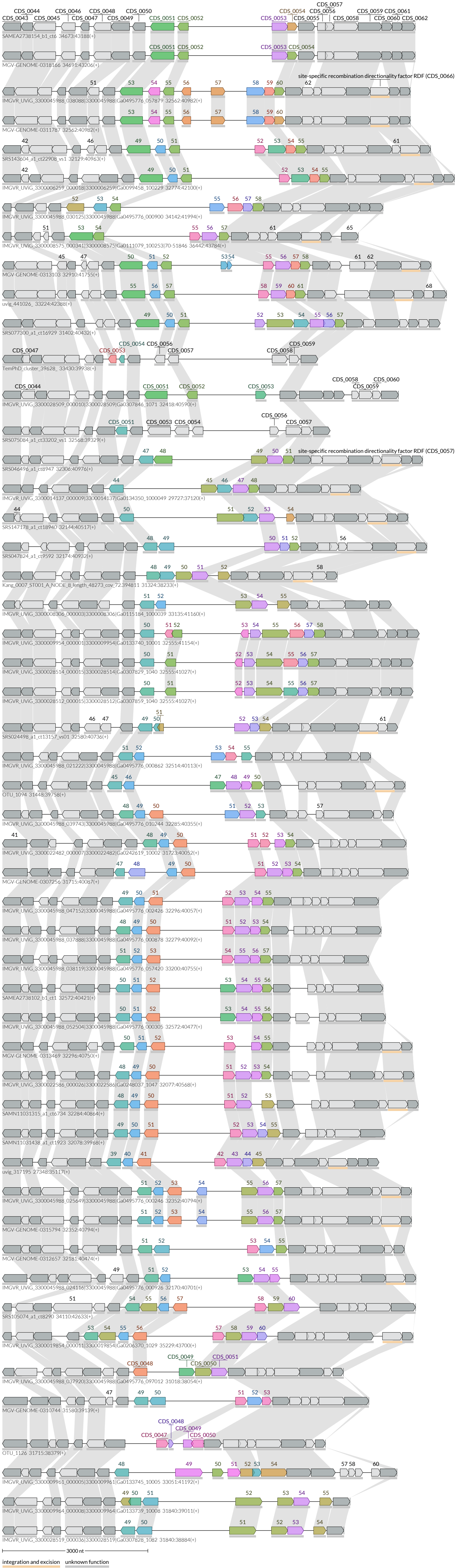
CDS_0044
CDS_0043
CDS_0045
CDS_0046
CDS_0047
CDS_0048
CDS_0049
CDS_0050
CDS_0051
CDS_0052
CDS_0053
CDS_0054
CDS_0055
CDS_0056
CDS_0057
CDS_0058
CDS_0059
CDS_0060
CDS_0061
CDS_0062
SAMEA2738154_b1_ct6 34673:43188(+)
MGV-GENOME-0318166 34691:43206(+)
site-specific recombination directionality factor RDF (CDS_0066)
IMGVR_UViG_3300045988_038088|3300045988|Ga0495776_057879 32562:40982(+)
MGV-GENOME-0311787 32562:40982(+)
SRS143604_a1_ct22908_vs1 32129:40963(+)
IMGVR_UViG_3300006259_000018|3300006259|Ga0099458_100229 32774:42100(+)
IMGVR_UViG_3300045988_030125|3300045988|Ga0495776_000900 34142:41994(+)
IMGVR_UViG_3300008575_000341|3300008575|Ga0111079_100253|70-51846 36442:43784(+)
MGV-GENOME-0313103 32910:41755(+)
uvig_441026_ 33224:42388(+)
SRS077300_a1_ct16929 31402:40432(+)
TemPhD_cluster_39628_ 33430:39938(+)
IMGVR_UViG_3300028509_000010|3300028509|Ga0307846_1071 32418:40590(+)
SRS075084_a1_ct33202_vs1 32568:39329(+)
site-specific recombination directionality factor RDF (CDS_0057)
SRS046496_a1_ct8947 32306:40976(+)
IMGVR_UViG_3300014137_000009|3300014137|Ga0134350_1000049 29727:37120(+)
SRS147178_a1_ct18940 32144:40517(+)
SRS047824_a1_ct9592 32174:40932(+)
Kang_0007_ST001_A_NODE_8_length_48273_cov_72.394811 31324:38233(+)
IMGVR_UViG_3300008306_000003|3300008306|Ga0115184_1000039 33135:41160(+)
IMGVR_UViG_3300009954_000001|3300009954|Ga0133740_10001 32555:41154(+)
IMGVR_UViG_3300028514_000015|3300028514|Ga0307829_1040 32555:41027(+)
IMGVR_UViG_3300028512_000015|3300028512|Ga0307859_1040 32555:41027(+)
SRS024498_a1_ct13157_vs01 32580:40736(+)
IMGVR_UViG_3300045988_021222|3300045988|Ga0495776_000862 32514:40113(+)
OTU_1094 31448:39758(+)
IMGVR_UViG_3300045988_039743|3300045988|Ga0495776_010244 32285:40355(+)
IMGVR_UViG_3300022482_000007|3300022482|Ga0242619_10002 31723:40052(+)
MGV-GENOME-0307256 31715:40087(+)
IMGVR_UViG_3300045988_047152|3300045988|Ga0495776_002426 32296:40057(+)
IMGVR_UViG_3300045988_037888|3300045988|Ga0495776_000878 32279:40092(+)
IMGVR_UViG_3300045988_038119|3300045988|Ga0495776_057420 33200:40755(+)
SAMEA2738102_b1_ct1 32572:40421(+)
IMGVR_UViG_3300045988_052504|3300045988|Ga0495776_000305 32572:40477(+)
MGV-GENOME-0313469 32296:40750(+)
IMGVR_UViG_3300022586_000026|3300022586|Ga0248037_1047 32077:40568(+)
SAMN11031315_a1_ct6734 32284:40864(+)
SAMN11031438_a1_ct1923 32078:39968(+)
uvig_317195 27348:35117(+)
IMGVR_UViG_3300045988_025649|3300045988|Ga0495776_000246 32352:40794(+)
MGV-GENOME-0315794 32352:40794(+)
MGV-GENOME-0312657 32181:40474(+)
IMGVR_UViG_3300045988_024116|3300045988|Ga0495776_000926 32170:40701(+)
SRS105074_a1_ct8290 34110:42633(+)
IMGVR_UViG_3300019854_000011|3300019854|Ga0206370_1029 35229:43700(+)
CDS_0048
CDS_0049
CDS_0050
CDS_0051
IMGVR_UViG_3300045988_037920|3300045988|Ga0495776_097012 31018:38054(+)
MGV-GENOME-0310744 31580:39139(+)
CDS_0047
OTU_1126 31715:38379(+)
IMGVR_UViG_3300009961_000005|3300009961|Ga0133745_10005 33051:41192(+)
IMGVR_UViG_3300009964_000008|3300009964|Ga0133739_10008 31840:39011(+)
IMGVR_UViG_3300028519_000036|3300028519|Ga0307828_1082 31840:38884(+)
3000 nt
integration and excision
unknown function