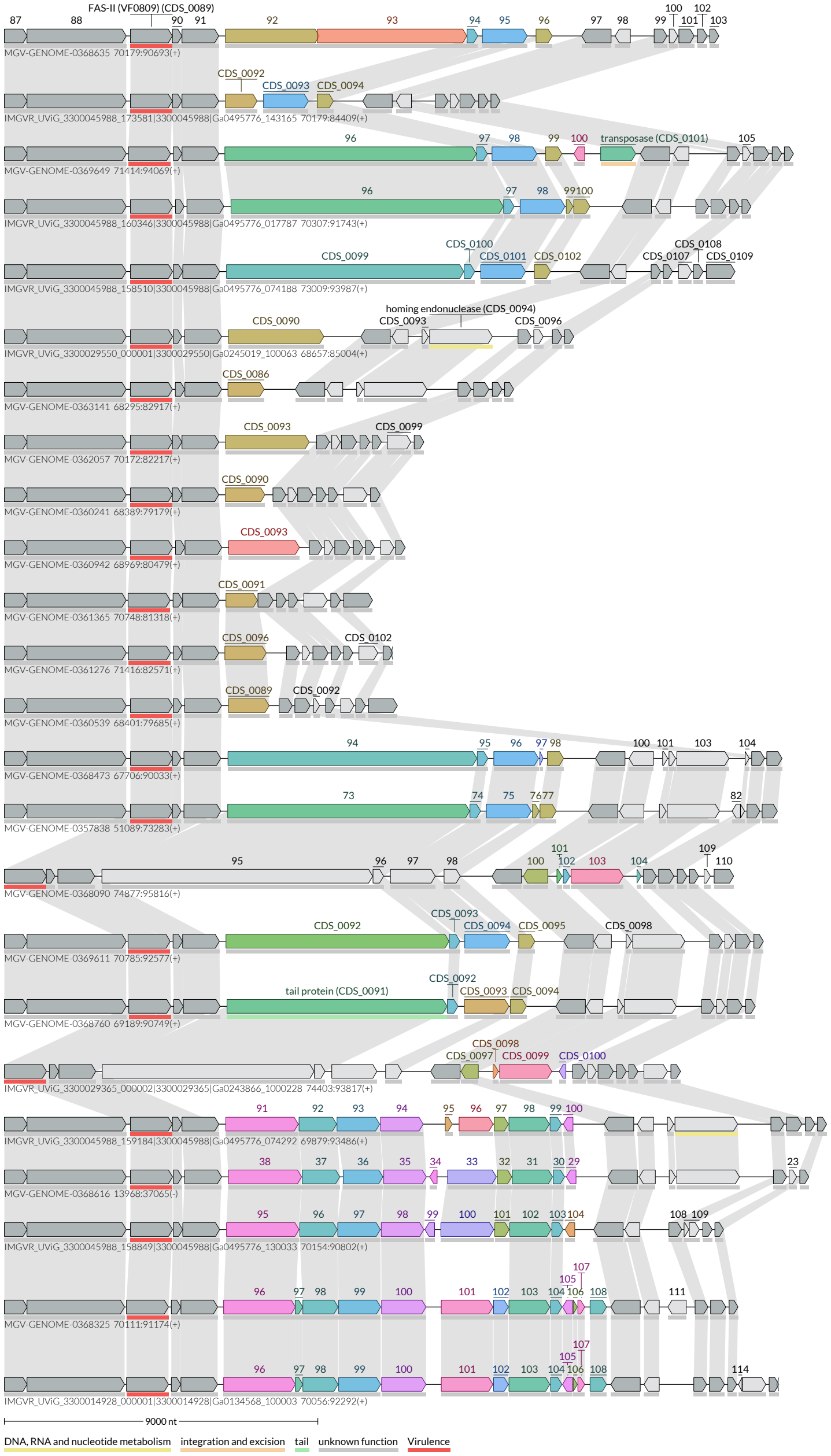
FAS-II (VF0809) (CDS_0089)
MGV-GENOME-0368635 70179:90693(+)
IMGVR_UViG_3300045988_173581|3300045988|Ga0495776_143165 70179:84409(+)
transposase (CDS_0101)
MGV-GENOME-0369649 71414:94069(+)
IMGVR_UViG_3300045988_160346|3300045988|Ga0495776_017787 70307:91743(+)
IMGVR_UViG_3300045988_158510|3300045988|Ga0495776_074188 73009:93987(+)
homing endonuclease (CDS_0094)
IMGVR_UViG_3300029550_000001|3300029550|Ga0245019_100063 68657:85004(+)
MGV-GENOME-0363141 68295:82917(+)
MGV-GENOME-0362057 70172:82217(+)
MGV-GENOME-0360241 68389:79179(+)
MGV-GENOME-0360942 68969:80479(+)
MGV-GENOME-0361365 70748:81318(+)
MGV-GENOME-0361276 71416:82571(+)
MGV-GENOME-0360539 68401:79685(+)
MGV-GENOME-0368473 67706:90033(+)
MGV-GENOME-0357838 51089:73283(+)
MGV-GENOME-0368090 74877:95816(+)
MGV-GENOME-0369611 70785:92577(+)
tail protein (CDS_0091)
MGV-GENOME-0368760 69189:90749(+)
IMGVR_UViG_3300029365_000002|3300029365|Ga0243866_1000228 74403:93817(+)
IMGVR_UViG_3300045988_159184|3300045988|Ga0495776_074292 69879:93486(+)
MGV-GENOME-0368616 13968:37065(-)
IMGVR_UViG_3300045988_158849|3300045988|Ga0495776_130033 70154:90802(+)
MGV-GENOME-0368325 70111:91174(+)
IMGVR_UViG_3300014928_000001|3300014928|Ga0134568_100003 70056:92292(+)
9000 nt
DNA, RNA and nucleotide metabolism
integration and excision
tail
unknown function
Virulence