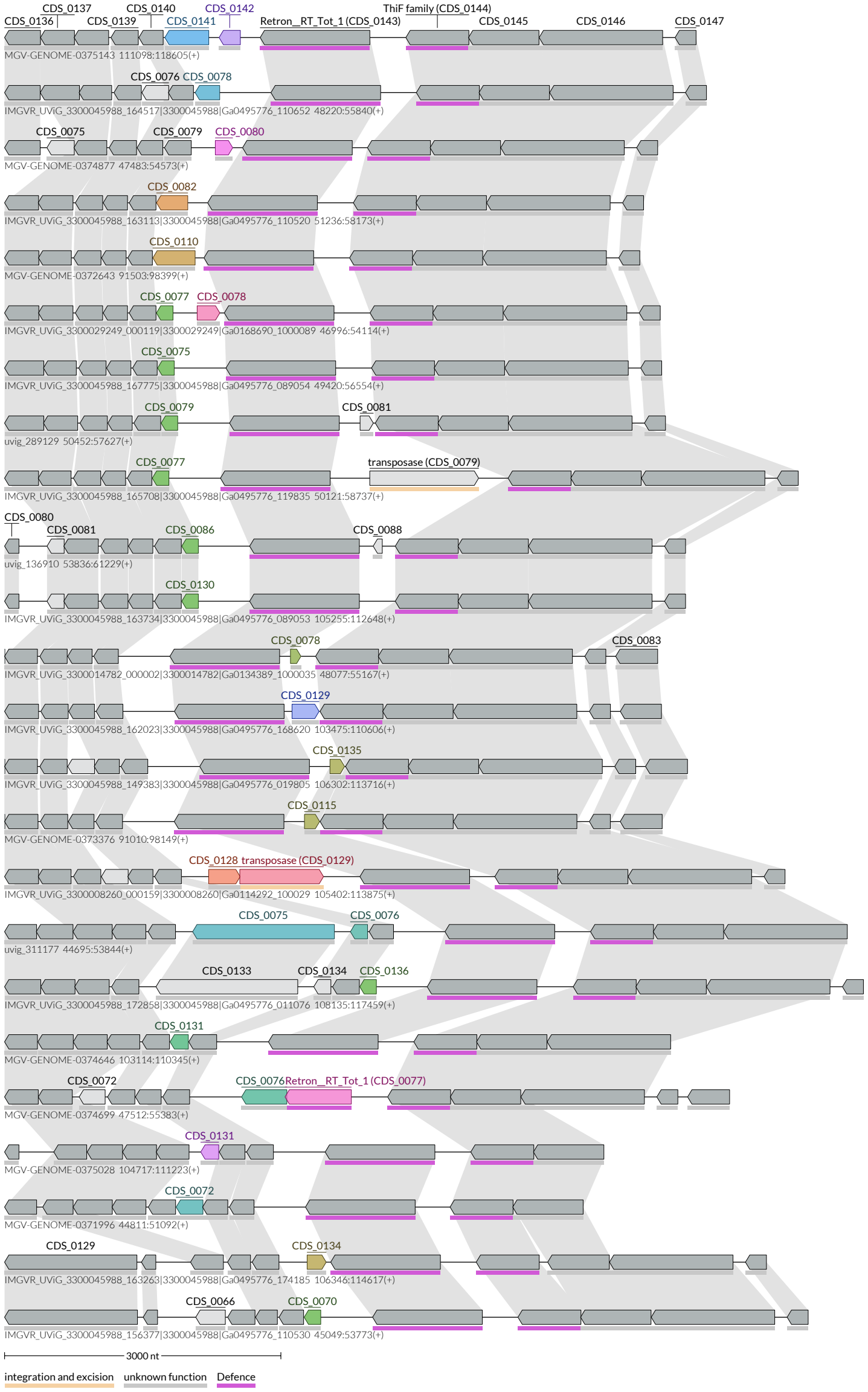
CDS_0137
CDS_0140
CDS_0142
ThiF family (CDS_0144)
CDS_0136
CDS_0139
CDS_0141
Retron__RT_Tot_1 (CDS_0143)
CDS_0145
CDS_0146
CDS_0147
MGV-GENOME-0375143 111098:118605(+)
CDS_0076
CDS_0078
IMGVR_UViG_3300045988_164517|3300045988|Ga0495776_110652 48220:55840(+)
CDS_0075
CDS_0079
CDS_0080
MGV-GENOME-0374877 47483:54573(+)
CDS_0082
IMGVR_UViG_3300045988_163113|3300045988|Ga0495776_110520 51236:58173(+)
CDS_0110
MGV-GENOME-0372643 91503:98399(+)
CDS_0077
CDS_0078
IMGVR_UViG_3300029249_000119|3300029249|Ga0168690_1000089 46996:54114(+)
CDS_0075
IMGVR_UViG_3300045988_167775|3300045988|Ga0495776_089054 49420:56554(+)
CDS_0079
CDS_0081
uvig_289129 50452:57627(+)
CDS_0077
transposase (CDS_0079)
IMGVR_UViG_3300045988_165708|3300045988|Ga0495776_119835 50121:58737(+)
CDS_0080
CDS_0081
CDS_0086
CDS_0088
uvig_136910 53836:61229(+)
CDS_0130
IMGVR_UViG_3300045988_163734|3300045988|Ga0495776_089053 105255:112648(+)
CDS_0078
CDS_0083
IMGVR_UViG_3300014782_000002|3300014782|Ga0134389_1000035 48077:55167(+)
CDS_0129
IMGVR_UViG_3300045988_162023|3300045988|Ga0495776_168620 103475:110606(+)
CDS_0135
IMGVR_UViG_3300045988_149383|3300045988|Ga0495776_019805 106302:113716(+)
CDS_0115
MGV-GENOME-0373376 91010:98149(+)
CDS_0128
transposase (CDS_0129)
IMGVR_UViG_3300008260_000159|3300008260|Ga0114292_100029 105402:113875(+)
CDS_0075
CDS_0076
uvig_311177 44695:53844(+)
CDS_0133
CDS_0134
CDS_0136
IMGVR_UViG_3300045988_172858|3300045988|Ga0495776_011076 108135:117459(+)
CDS_0131
MGV-GENOME-0374646 103114:110345(+)
CDS_0072
CDS_0076
Retron__RT_Tot_1 (CDS_0077)
MGV-GENOME-0374699 47512:55383(+)
CDS_0131
MGV-GENOME-0375028 104717:111223(+)
CDS_0072
MGV-GENOME-0371996 44811:51092(+)
CDS_0129
CDS_0134
IMGVR_UViG_3300045988_163263|3300045988|Ga0495776_174185 106346:114617(+)
CDS_0066
CDS_0070
IMGVR_UViG_3300045988_156377|3300045988|Ga0495776_110530 45049:53773(+)
3000 nt
integration and excision
unknown function
Defence