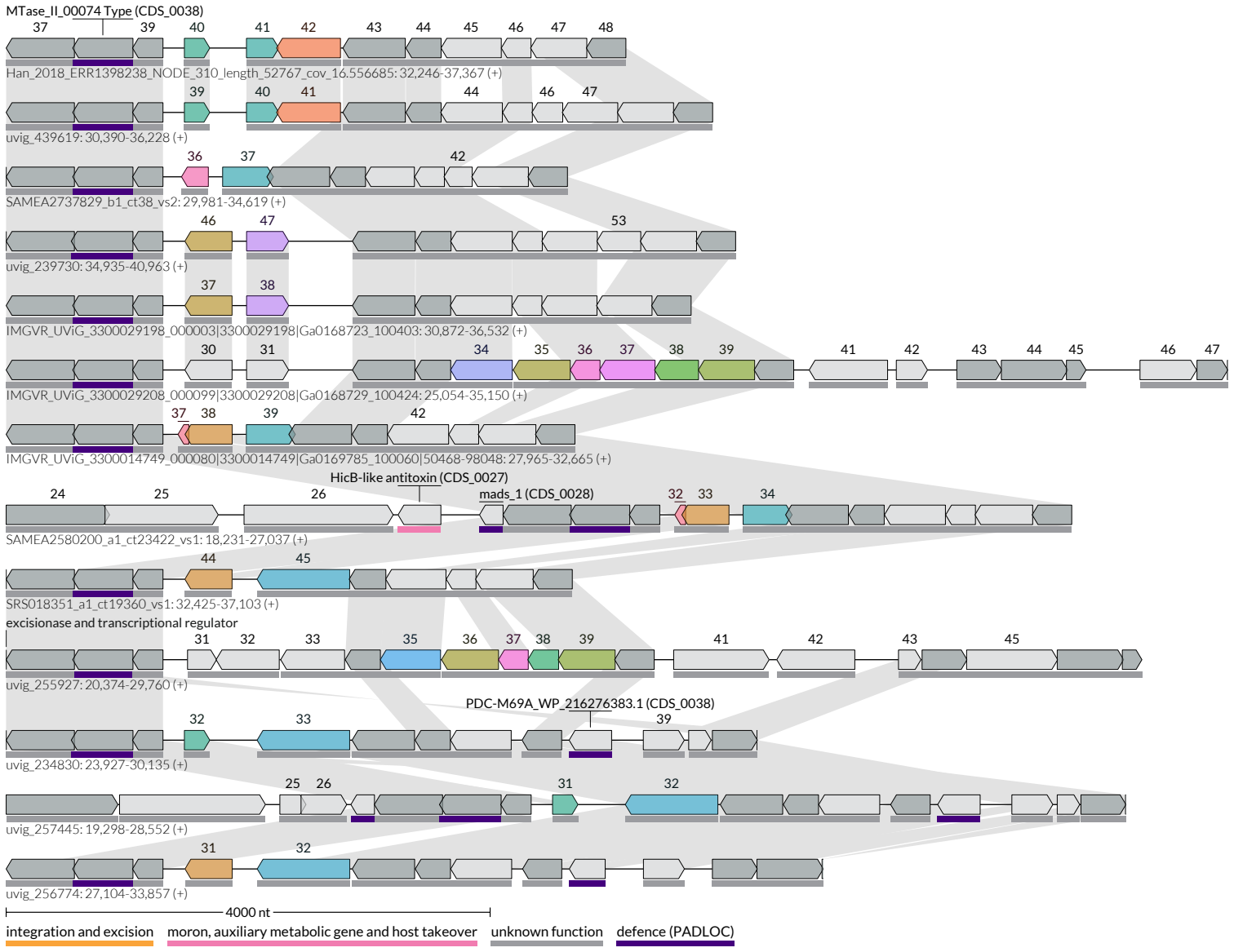
MTase_II_00074 Type (CDS_0038)
37
39
40
41
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44
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46
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48
Han_2018_ERR1398238_NODE_310_length_52767_cov_16.556685: 32,246-37,367 (+)
39
40
41
44
46
47
uvig_439619: 30,390-36,228 (+)
36
37
42
SAMEA2737829_b1_ct38_vs2: 29,981-34,619 (+)
46
47
53
uvig_239730: 34,935-40,963 (+)
37
38
IMGVR_UViG_3300029198_000003|3300029198|Ga0168723_100403: 30,872-36,532 (+)
30
31
34
35
36
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41
42
43
44
45
46
47
IMGVR_UViG_3300029208_000099|3300029208|Ga0168729_100424: 25,054-35,150 (+)
37
38
39
42
IMGVR_UViG_3300014749_000080|3300014749|Ga0169785_100060|50468-98048: 27,965-32,665 (+)
HicB-like antitoxin (CDS_0027)
24
25
26
mads_1 (CDS_0028)
32
33
34
SAMEA2580200_a1_ct23422_vs1: 18,231-27,037 (+)
44
45
SRS018351_a1_ct19360_vs1: 32,425-37,103 (+)
excisionase and transcriptional regulator
31
32
33
35
36
37
38
39
41
42
43
45
uvig_255927: 20,374-29,760 (+)
PDC-M69A_WP_216276383.1 (CDS_0038)
32
33
39
uvig_234830: 23,927-30,135 (+)
25
26
31
32
uvig_257445: 19,298-28,552 (+)
31
32
uvig_256774: 27,104-33,857 (+)
4000 nt
integration and excision
moron, auxiliary metabolic gene and host takeover
unknown function
defence (PADLOC)