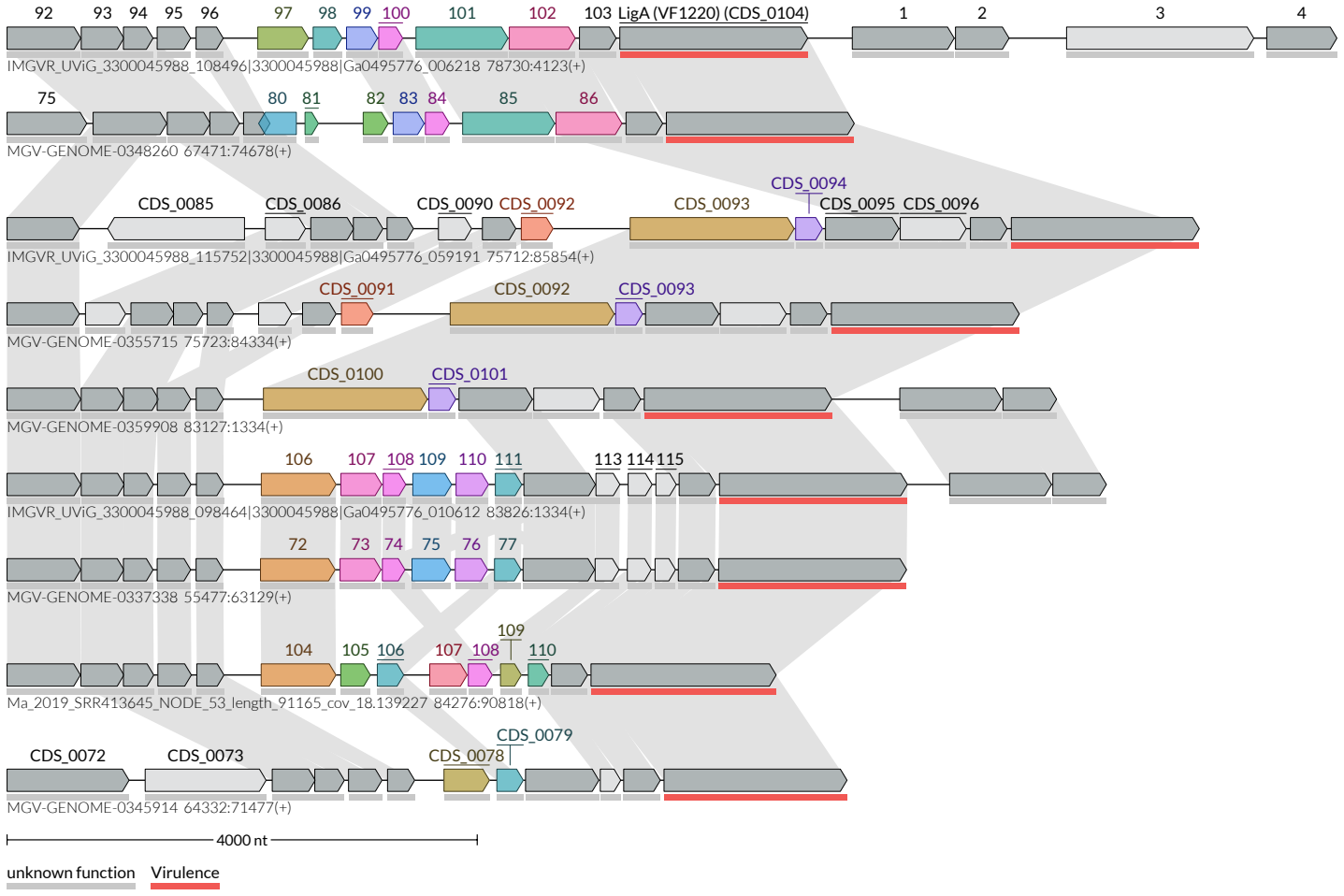
92
93
94
95
96
97
98
99
100
101
102
103
LigA (VF1220) (CDS_0104)
1
2
3
4
IMGVR_UViG_3300045988_108496|3300045988|Ga0495776_006218 78730:4123(+)
75
80
81
82
83
84
85
86
MGV-GENOME-0348260 67471:74678(+)
CDS_0085
CDS_0086
CDS_0090
CDS_0092
CDS_0093
CDS_0094
CDS_0095
CDS_0096
IMGVR_UViG_3300045988_115752|3300045988|Ga0495776_059191 75712:85854(+)
CDS_0091
CDS_0092
CDS_0093
MGV-GENOME-0355715 75723:84334(+)
CDS_0100
CDS_0101
MGV-GENOME-0359908 83127:1334(+)
106
107
108
109
110
111
113
114
115
IMGVR_UViG_3300045988_098464|3300045988|Ga0495776_010612 83826:1334(+)
72
73
74
75
76
77
MGV-GENOME-0337338 55477:63129(+)
104
105
106
107
108
109
110
Ma_2019_SRR413645_NODE_53_length_91165_cov_18.139227 84276:90818(+)
CDS_0072
CDS_0073
CDS_0078
CDS_0079
MGV-GENOME-0345914 64332:71477(+)
4000 nt
unknown function
Virulence