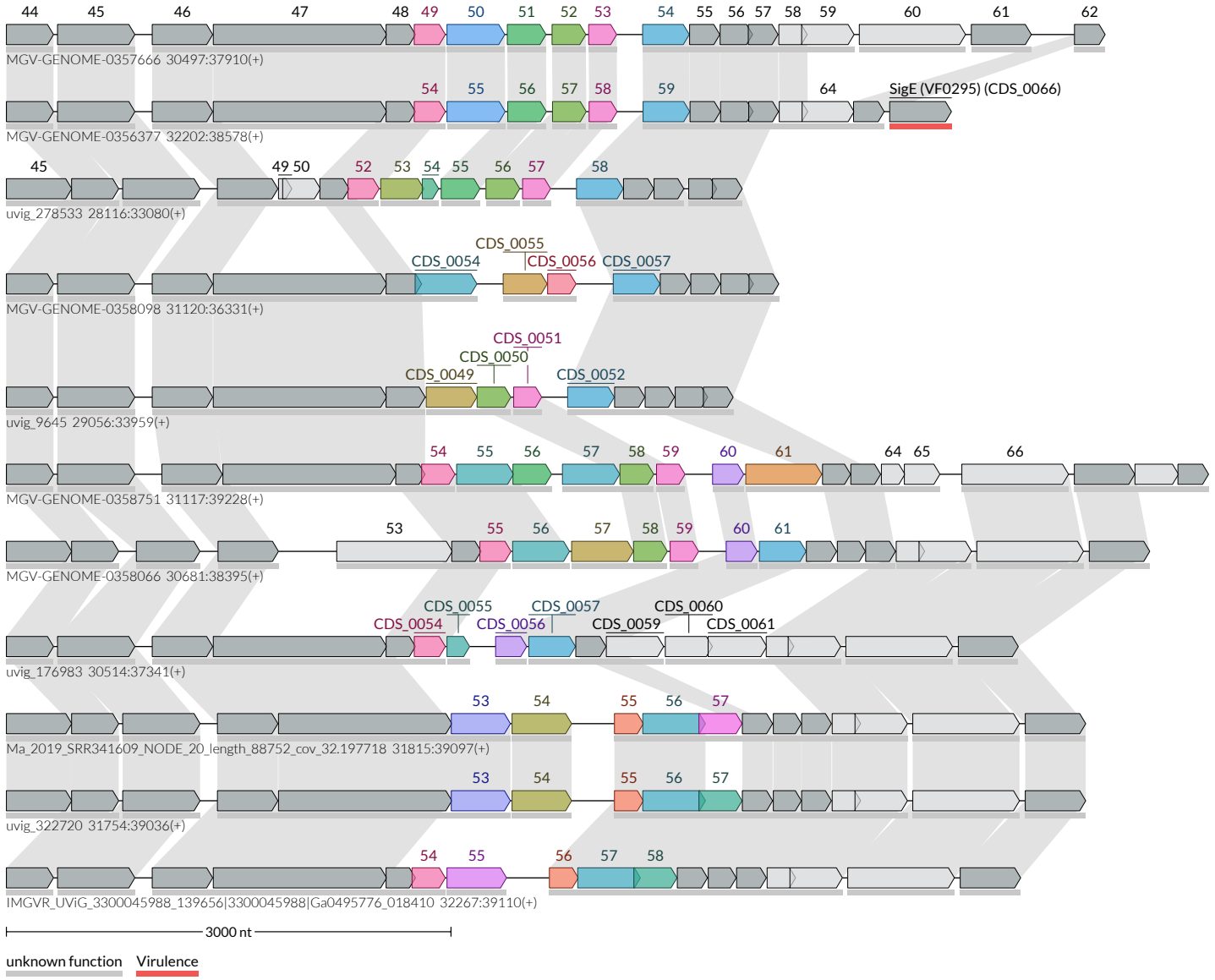
44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62
MGV-GENOME-0357666 30497:37910(+)
54 55 56 57 58 59 64
SigE (VF0295) (CDS_0066)
MGV-GENOME-0356377 32202:38578(+)
45 49 50 52 53 54 55 56 57 58
uvig_278533 28116:33080(+)
CDS_0055
CDS_0054 CDS_0056 CDS_0057
MGV-GENOME-0358098 31120:36331(+)
CDS_0051
CDS_0050
CDS_0049 CDS_0052
uvig_9645 29056:33959(+)
54 55 56 57 58 59 60 61 64 65 66
MGV-GENOME-0358751 31117:39228(+)
53 55 56 57 58 59 60 61
MGV-GENOME-0358066 30681:38395(+)
CDS_0055 CDS_0057 CDS_0060
CDS_0054 CDS_0056 CDS_0059 CDS_0061
uvig_176983 30514:37341(+)
53 54 55 56 57
Ma_2019_SRR341609_NODE_20_length_88752_cov_32.197718 31815:39097(+)
53 54 55 56 57
uvig_322720 31754:39036(+)
54 55 56 57 58
IMGVR_UViG_3300045988_139656|3300045988|Ga0495776_018410 32267:39110(+)
3000 nt
unknown function
Virulence