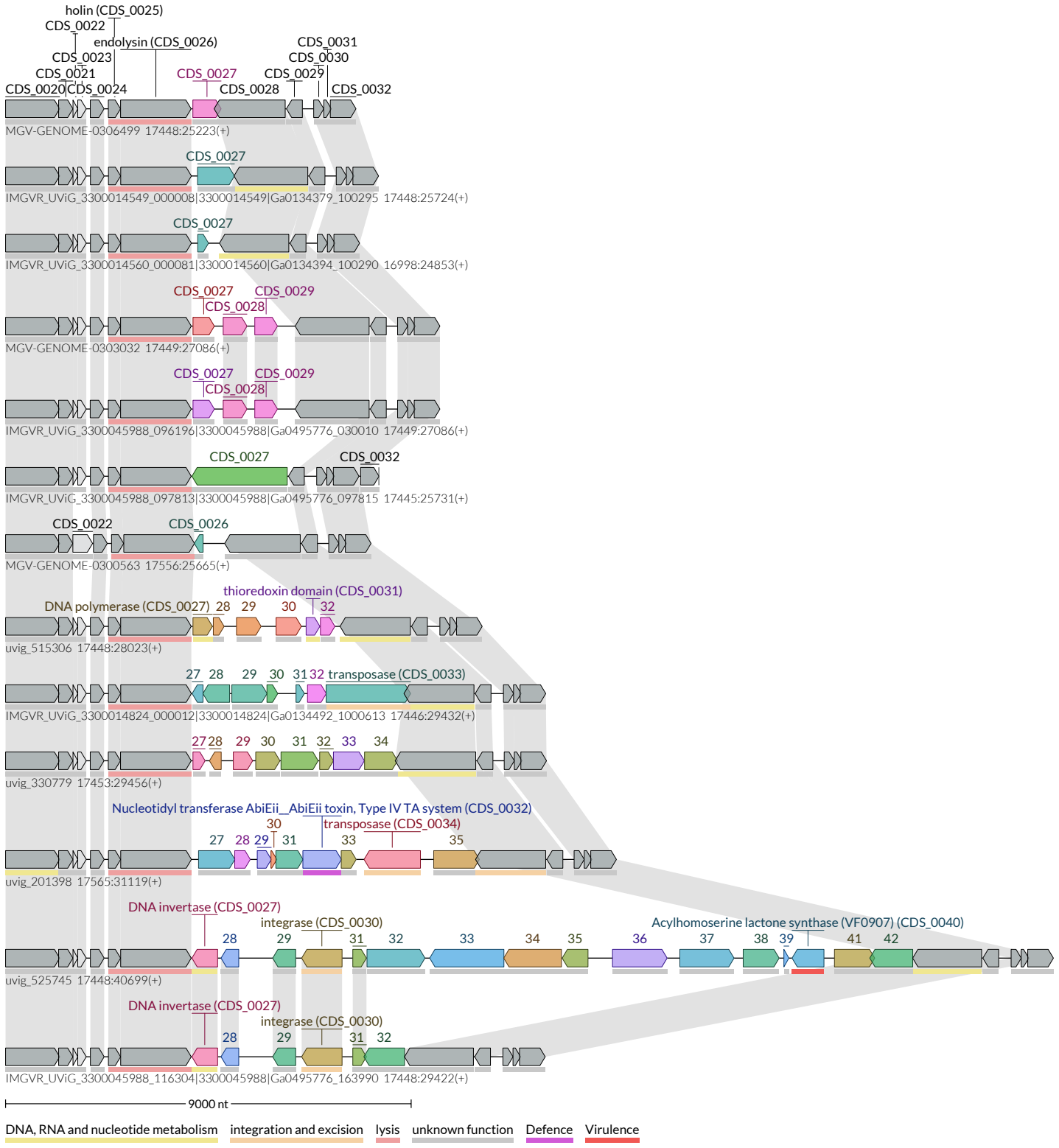
holin (CDS_0025)
CDS_0022
endolysin (CDS_0026)
CDS_0031
CDS_0023
CDS_0030
CDS_0021
CDS_0027
CDS_0029
CDS_0020
CDS_0024
CDS_0028
CDS_0032
MGV-GENOME-0306499 17448:25223(+)
CDS_0027
IMGVR_UViG_3300014549_000008|3300014549|Ga0134379_100295 17448:25724(+)
CDS_0027
IMGVR_UViG_3300014560_000081|3300014560|Ga0134394_100290 16998:24853(+)
CDS_0027
CDS_0029
CDS_0028
MGV-GENOME-0303032 17449:27086(+)
CDS_0027
CDS_0029
CDS_0028
IMGVR_UViG_3300045988_096196|3300045988|Ga0495776_030010 17449:27086(+)
CDS_0027
CDS_0032
IMGVR_UViG_3300045988_097813|3300045988|Ga0495776_097815 17445:25731(+)
CDS_0022
CDS_0026
MGV-GENOME-0300563 17556:25665(+)
thioredoxin domain (CDS_0031)
DNA polymerase (CDS_0027)
28
29
30
32
uvig_515306 17448:28023(+)
27
28
29
30
31
32
transposase (CDS_0033)
IMGVR_UViG_3300014824_000012|3300014824|Ga0134492_1000613 17446:29432(+)
27
28
29
30
31
32
33
34
uvig_330779 17453:29456(+)
Nucleotidyl transferase AbiEii__AbiEii toxin, Type IV TA system (CDS_0032)
30
transposase (CDS_0034)
27
28
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31
33
35
uvig_201398 17565:31119(+)
DNA invertase (CDS_0027)
integrase (CDS_0030)
Acylhomoserine lactone synthase (VF0907) (CDS_0040)
28
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33
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35
36
37
38
39
41
42
uvig_525745 17448:40699(+)
DNA invertase (CDS_0027)
integrase (CDS_0030)
28
29
31
32
IMGVR_UViG_3300045988_116304|3300045988|Ga0495776_163990 17448:29422(+)
9000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
unknown function
Defence
Virulence