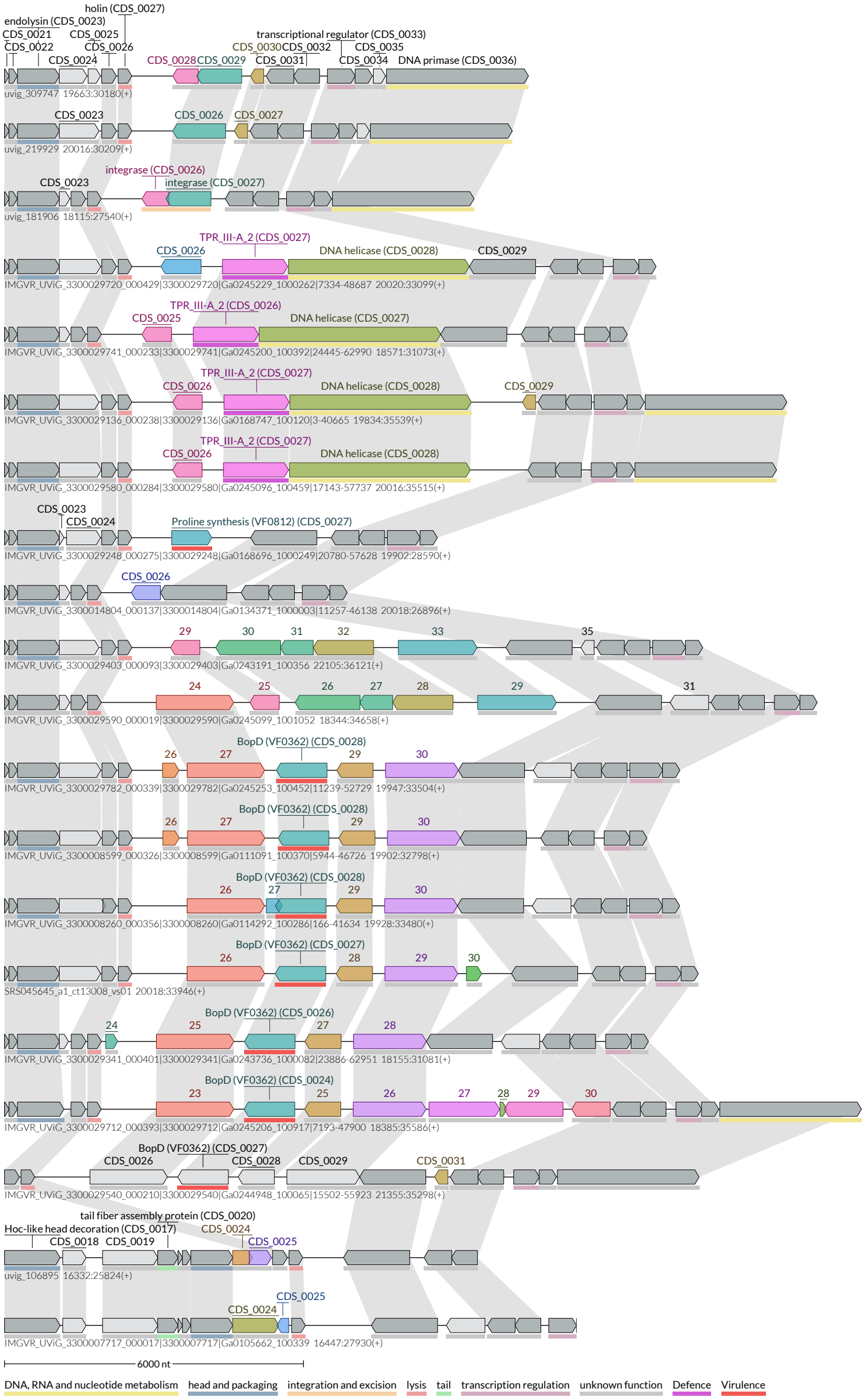
holin (CDS_0027)
endolysin (CDS_0023)
CDS_0021
CDS_0025
CDS_0022
CDS_0026
transcriptional regulator (CDS_0033)
CDS_0030
CDS_0032
CDS_0035
CDS_0024
CDS_0028
CDS_0029
CDS_0031
CDS_0034
DNA primase (CDS_0036)
uvig_309747 19663:30180(+)
CDS_0023
CDS_0026
CDS_0027
uvig_219929 20016:30209(+)
integrase (CDS_0026)
CDS_0023
integrase (CDS_0027)
uvig_181906 18115:27540(+)
TPR_III-A_2 (CDS_0027)
CDS_0026
DNA helicase (CDS_0028)
CDS_0029
IMGVR_UViG_3300029720_000429|3300029720|Ga0245229_1000262|7334-48687 20020:33099(+)
TPR_III-A_2 (CDS_0026)
CDS_0025
DNA helicase (CDS_0027)
IMGVR_UViG_3300029741_000233|3300029741|Ga0245200_100392|24445-62990 18571:31073(+)
TPR_III-A_2 (CDS_0027)
CDS_0026
DNA helicase (CDS_0028)
CDS_0029
IMGVR_UViG_3300029136_000238|3300029136|Ga0168747_100120|3-40665 19834:35539(+)
TPR_III-A_2 (CDS_0027)
CDS_0026
DNA helicase (CDS_0028)
IMGVR_UViG_3300029580_000284|3300029580|Ga0245096_100459|17143-57737 20016:35515(+)
CDS_0023
CDS_0024
Proline synthesis (VF0812) (CDS_0027)
IMGVR_UViG_3300029248_000275|3300029248|Ga0168696_1000249|20780-57628 19902:28590(+)
CDS_0026
IMGVR_UViG_3300014804_000137|3300014804|Ga0134371_1000003|11257-46138 20018:26896(+)
29
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31
32
33
35
IMGVR_UViG_3300029403_000093|3300029403|Ga0243191_100356 22105:36121(+)
24
25
26
27
28
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31
IMGVR_UViG_3300029590_000019|3300029590|Ga0245099_1001052 18344:34658(+)
BopD (VF0362) (CDS_0028)
26
27
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IMGVR_UViG_3300029782_000339|3300029782|Ga0245253_100452|11239-52729 19947:33504(+)
BopD (VF0362) (CDS_0028)
26
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IMGVR_UViG_3300008599_000326|3300008599|Ga0111091_100370|5944-46726 19902:32798(+)
BopD (VF0362) (CDS_0028)
26
27
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30
IMGVR_UViG_3300008260_000356|3300008260|Ga0114292_100286|166-41634 19928:33480(+)
BopD (VF0362) (CDS_0027)
26
28
29
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SRS045645_a1_ct13008_vs01 20018:33946(+)
BopD (VF0362) (CDS_0026)
24
25
27
28
IMGVR_UViG_3300029341_000401|3300029341|Ga0243736_1000082|23886-62951 18155:31081(+)
BopD (VF0362) (CDS_0024)
23
25
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30
IMGVR_UViG_3300029712_000393|3300029712|Ga0245206_100917|7193-47900 18385:35586(+)
BopD (VF0362) (CDS_0027)
CDS_0026
CDS_0028
CDS_0029
CDS_0031
IMGVR_UViG_3300029540_000210|3300029540|Ga0244948_100065|15502-55923 21355:35298(+)
tail fiber assembly protein (CDS_0020)
Hoc-like head decoration (CDS_0017)
CDS_0024
CDS_0018
CDS_0019
CDS_0025
uvig_106895 16332:25824(+)
CDS_0025
CDS_0024
IMGVR_UViG_3300007717_000017|3300007717|Ga0105662_100339 16447:27930(+)
6000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
lysis
tail
transcription regulation
unknown function
Defence
Virulence