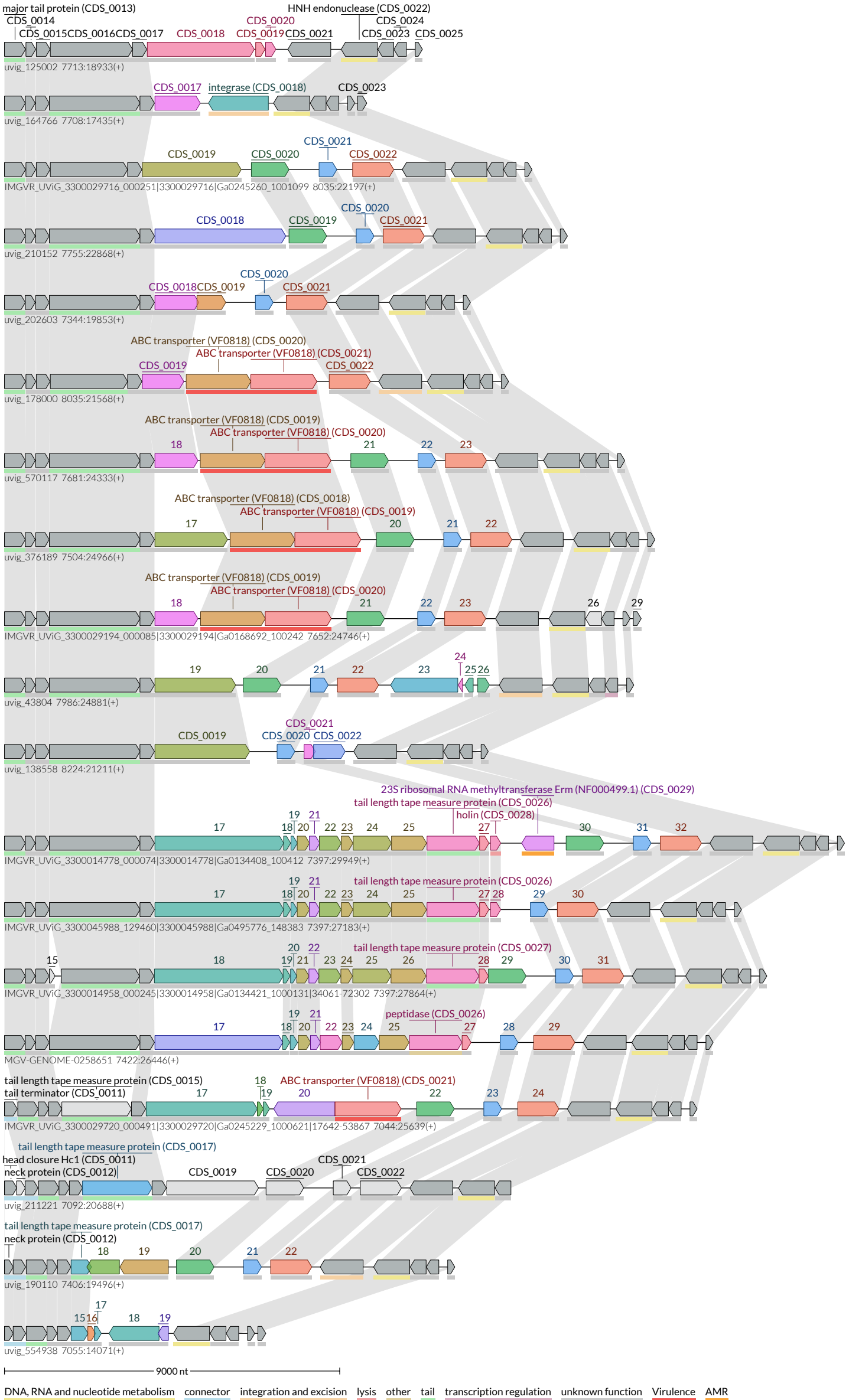
major tail protein (CDS_0013)
CDS_0014
CDS_0015
CDS_0016
CDS_0017
CDS_0018
CDS_0020
CDS_0019
CDS_0021
HNH endonuclease (CDS_0022)
CDS_0024
CDS_0023
CDS_0025
uvig_125002 7713:18933(+)
CDS_0017
integrase (CDS_0018)
CDS_0023
uvig_164766 7708:17435(+)
CDS_0021
CDS_0019
CDS_0020
CDS_0022
IMGVR_UViG_3300029716_000251|3300029716|Ga0245260_1001099 8035:22197(+)
CDS_0020
CDS_0018
CDS_0019
CDS_0021
uvig_210152 7755:22868(+)
CDS_0020
CDS_0018
CDS_0019
CDS_0021
uvig_202603 7344:19853(+)
ABC transporter (VF0818) (CDS_0020)
ABC transporter (VF0818) (CDS_0021)
CDS_0019
CDS_0022
uvig_178000 8035:21568(+)
ABC transporter (VF0818) (CDS_0019)
ABC transporter (VF0818) (CDS_0020)
18
21
22
23
uvig_570117 7681:24333(+)
ABC transporter (VF0818) (CDS_0018)
ABC transporter (VF0818) (CDS_0019)
17
20
21
22
uvig_376189 7504:24966(+)
ABC transporter (VF0818) (CDS_0019)
ABC transporter (VF0818) (CDS_0020)
18
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23
26
29
IMGVR_UViG_3300029194_000085|3300029194|Ga0168692_100242 7652:24746(+)
19
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22
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24
25
26
uvig_43804 7986:24881(+)
CDS_0021
CDS_0019
CDS_0020
CDS_0022
uvig_138558 8224:21211(+)
23S ribosomal RNA methyltransferase Erm (NF000499.1) (CDS_0029)
tail length tape measure protein (CDS_0026)
holin (CDS_0028)
17
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32
IMGVR_UViG_3300014778_000074|3300014778|Ga0134408_100412 7397:29949(+)
tail length tape measure protein (CDS_0026)
17
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30
IMGVR_UViG_3300045988_129460|3300045988|Ga0495776_148383 7397:27183(+)
tail length tape measure protein (CDS_0027)
15
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31
IMGVR_UViG_3300014958_000245|3300014958|Ga0134421_1000131|34061-72302 7397:27864(+)
peptidase (CDS_0026)
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MGV-GENOME-0258651 7422:26446(+)
tail length tape measure protein (CDS_0015)
tail terminator (CDS_0011)
ABC transporter (VF0818) (CDS_0021)
17
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24
IMGVR_UViG_3300029720_000491|3300029720|Ga0245229_1000621|17642-53867 7044:25639(+)
tail length tape measure protein (CDS_0017)
head closure Hc1 (CDS_0011)
neck protein (CDS_0012)
CDS_0019
CDS_0020
CDS_0021
CDS_0022
uvig_211221 7092:20688(+)
tail length tape measure protein (CDS_0017)
neck protein (CDS_0012)
18
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uvig_190110 7406:19496(+)
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16
18
19
uvig_554938 7055:14071(+)
9000 nt
DNA, RNA and nucleotide metabolism
connector
integration and excision
lysis
other
tail
transcription regulation
unknown function
Virulence
AMR