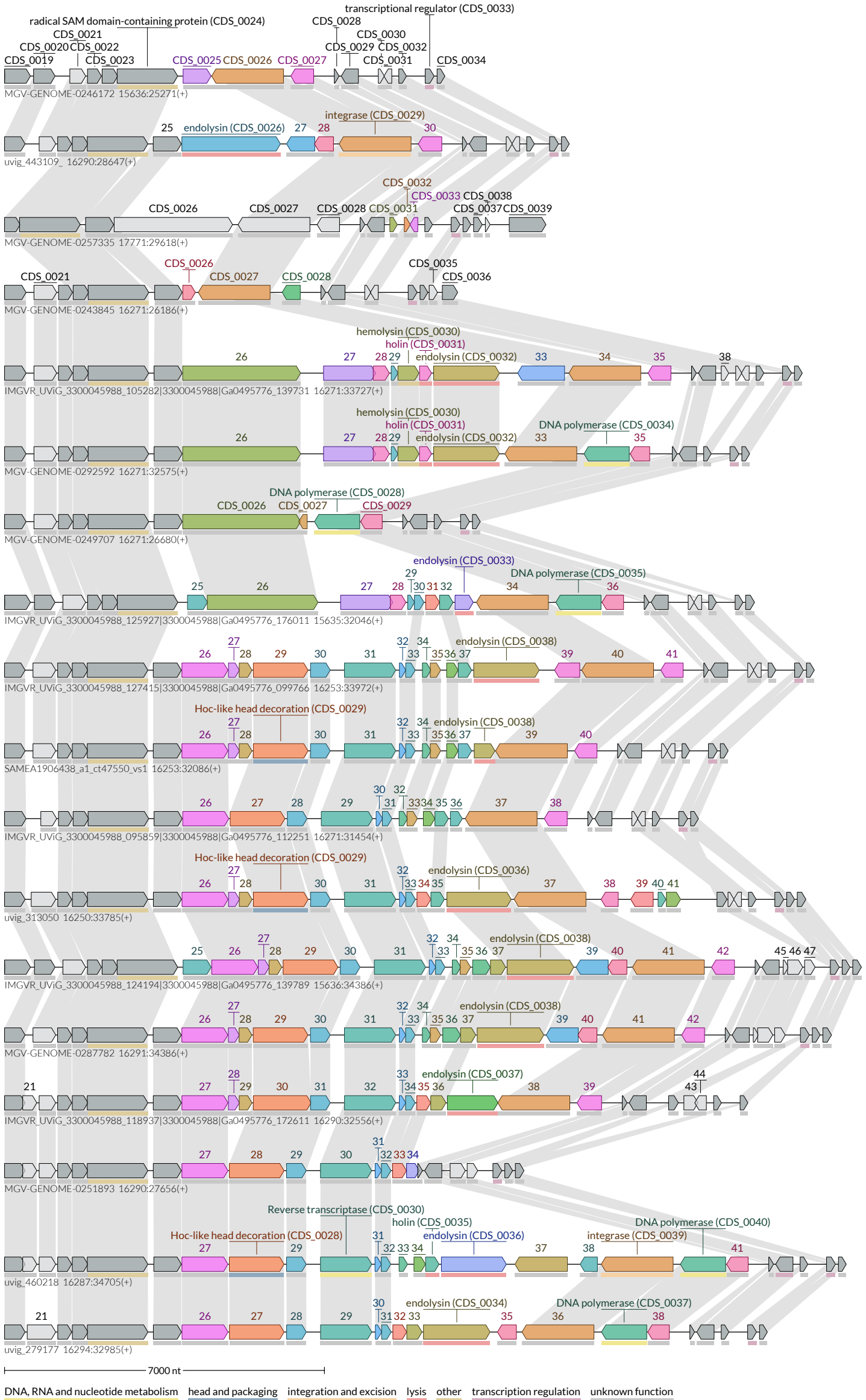
transcriptional regulator (CDS_0033)
radical SAM domain-containing protein (CDS_0024)
CDS_0028
CDS_0021
CDS_0030
CDS_0020 CDS_0022
CDS_0029 CDS_0032
CDS_0019
CDS_0023
CDS_0025 CDS_0026 CDS_0027
CDS_0031
CDS_0034
MGV-GENOME-0246172 15636:25271(+)
integrase (CDS_0029)
25
endolysin (CDS_0026)
27
28
30
uvig_443109_ 16290:28647(+)
CDS_0032
CDS_0033 CDS_0038
CDS_0026
CDS_0027
CDS_0028 CDS_0031
CDS_0037 CDS_0039
MGV-GENOME-0257335 17771:29618(+)
CDS_0026
CDS_0035
CDS_0021
CDS_0027
CDS_0028
CDS_0036
MGV-GENOME-0243845 16271:26186(+)
hemolysin (CDS_0030)
holin (CDS_0031)
26
27
28 29
endolysin (CDS_0032)
33
34
35
38
IMGVR_UViG_3300045988_105282|3300045988|Ga0495776_139731 16271:33727(+)
hemolysin (CDS_0030)
holin (CDS_0031)
DNA polymerase (CDS_0034)
26
27
28 29
endolysin (CDS_0032)
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35
MGV-GENOME-0292592 16271:32575(+)
DNA polymerase (CDS_0028)
CDS_0026
CDS_0027
CDS_0029
MGV-GENOME-0249707 16271:26680(+)
endolysin (CDS_0033)
29
DNA polymerase (CDS_0035)
25
26
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30 31 32
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36
IMGVR_UViG_3300045988_125927|3300045988|Ga0495776_176011 15635:32046(+)
27
32
34
endolysin (CDS_0038)
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35 36 37
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41
IMGVR_UViG_3300045988_127415|3300045988|Ga0495776_099766 16253:33972(+)
Hoc-like head decoration (CDS_0029)
27
32
34
endolysin (CDS_0038)
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35 36 37
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SAMEA1906438_a1_ct47550_vs1 16253:32086(+)
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33 34 35
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38
IMGVR_UViG_3300045988_095859|3300045988|Ga0495776_112251 16271:31454(+)
Hoc-like head decoration (CDS_0029)
27
32
endolysin (CDS_0036)
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33 34 35
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39 40 41
uvig_313050 16250:33785(+)
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endolysin (CDS_0038)
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35 36 37
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45 46 47
IMGVR_UViG_3300045988_124194|3300045988|Ga0495776_139789 15636:34386(+)
27
32
34
endolysin (CDS_0038)
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35 36 37
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MGV-GENOME-0287782 16291:34386(+)
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33
endolysin (CDS_0037)
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34 35 36
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IMGVR_UViG_3300045988_118937|3300045988|Ga0495776_172611 16290:32556(+)
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32 33 34
MGV-GENOME-0251893 16290:27656(+)
Reverse transcriptase (CDS_0030)
holin (CDS_0035)
DNA polymerase (CDS_0040)
Hoc-like head decoration (CDS_0028)
31
endolysin (CDS_0036)
integrase (CDS_0039)
27
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32 33 34
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41
uvig_460218 16287:34705(+)
endolysin (CDS_0034)
DNA polymerase (CDS_0037)
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21
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31 32 33
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uvig_279177 16294:32985(+)
7000 nt
DNA, RNA and nucleotide metabolism
head and packaging
integration and excision
lysis
other
transcription regulation
unknown function