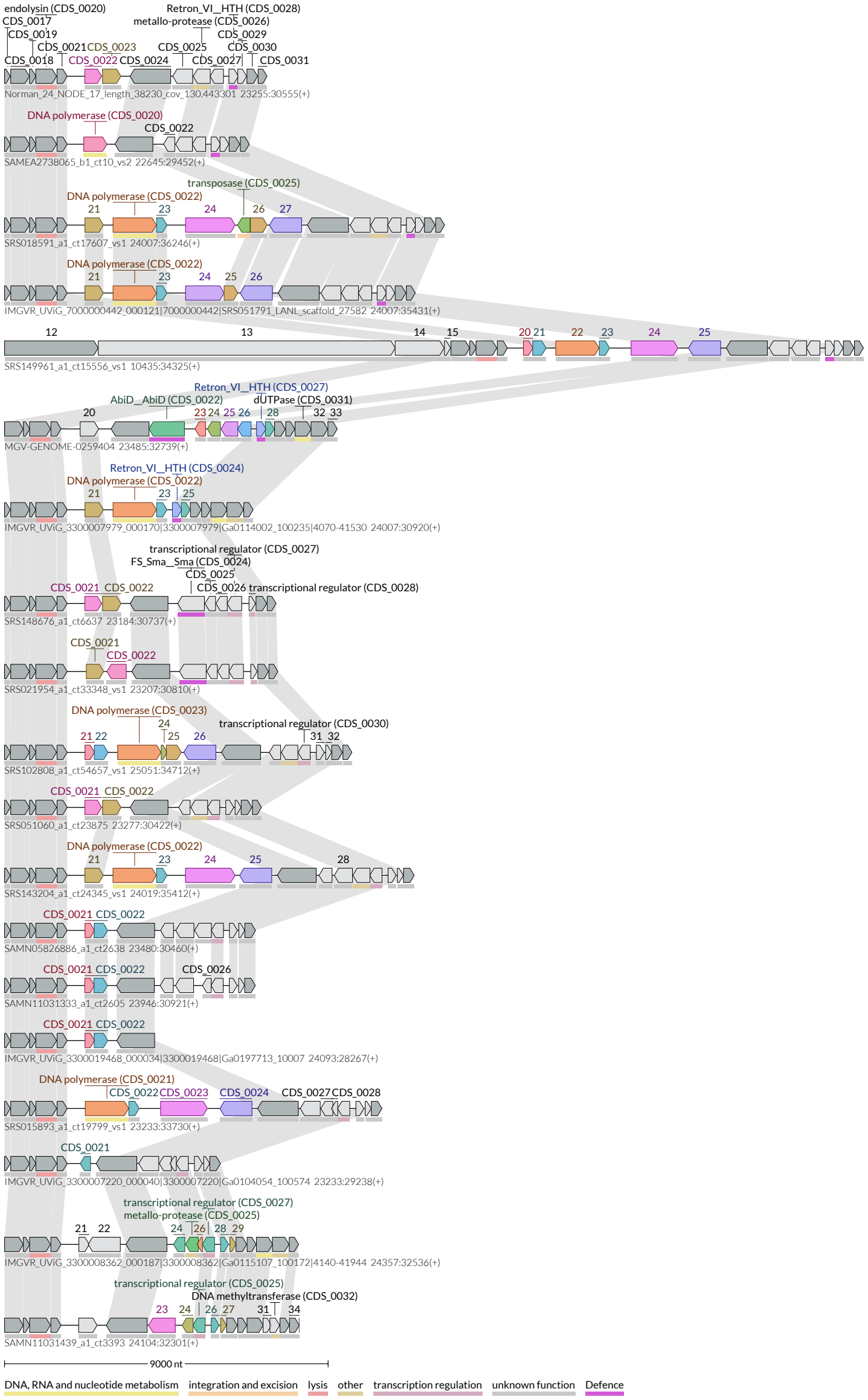
endolysin (CDS_0020)
Retron_VI__HTH (CDS_0028)
metallo-protease (CDS_0026)
CDS_0017
CDS_0019
CDS_0029
CDS_0021
CDS_0023
CDS_0025
CDS_0030
CDS_0018
CDS_0022
CDS_0024
CDS_0027
CDS_0031
Norman_24_NODE_17_length_38230_cov_130.443301 23255:30555(+)
DNA polymerase (CDS_0020)
CDS_0022
SAMEA2738065_b1_ct10_vs2 22645:29452(+)
transposase (CDS_0025)
DNA polymerase (CDS_0022)
21
23
24
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27
SRS018591_a1_ct17607_vs1 24007:36246(+)
DNA polymerase (CDS_0022)
21
23
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26
IMGVR_UViG_7000000442_000121|7000000442|SRS051791_LANL_scaffold_27582 24007:35431(+)
12
13
14
15
20
21
22
23
24
25
SRS149961_a1_ct15556_vs1 10435:34325(+)
Retron_VI__HTH (CDS_0027)
AbiD__AbiD (CDS_0022)
dUTPase (CDS_0031)
20
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32
33
MGV-GENOME-0259404 23485:32739(+)
Retron_VI__HTH (CDS_0024)
DNA polymerase (CDS_0022)
21
23
25
IMGVR_UViG_3300007979_000170|3300007979|Ga0114002_100235|4070-41530 24007:30920(+)
transcriptional regulator (CDS_0027)
FS_Sma__Sma (CDS_0024)
CDS_0025
CDS_0021
CDS_0022
CDS_0026
transcriptional regulator (CDS_0028)
SRS148676_a1_ct6637 23184:30737(+)
CDS_0021
CDS_0022
SRS021954_a1_ct33348_vs1 23207:30810(+)
DNA polymerase (CDS_0023)
transcriptional regulator (CDS_0030)
21
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32
SRS102808_a1_ct54657_vs1 25051:34712(+)
CDS_0021
CDS_0022
SRS051060_a1_ct23875 23277:30422(+)
DNA polymerase (CDS_0022)
21
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SRS143204_a1_ct24345_vs1 24019:35412(+)
CDS_0021
CDS_0022
SAMN05826886_a1_ct2638 23480:30460(+)
CDS_0021
CDS_0022
CDS_0026
SAMN11031333_a1_ct2605 23946:30921(+)
CDS_0021
CDS_0022
IMGVR_UViG_3300019468_000034|3300019468|Ga0197713_10007 24093:28267(+)
DNA polymerase (CDS_0021)
CDS_0022
CDS_0023
CDS_0024
CDS_0027
CDS_0028
SRS015893_a1_ct19799_vs1 23233:33730(+)
CDS_0021
IMGVR_UViG_3300007220_000040|3300007220|Ga0104054_100574 23233:29238(+)
transcriptional regulator (CDS_0027)
metallo-protease (CDS_0025)
21
22
24
26
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29
IMGVR_UViG_3300008362_000187|3300008362|Ga0115107_100172|4140-41944 24357:32536(+)
transcriptional regulator (CDS_0025)
DNA methyltransferase (CDS_0032)
23
24
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34
SAMN11031439_a1_ct3393 24104:32301(+)
9000 nt
DNA, RNA and nucleotide metabolism
integration and excision
lysis
other
transcription regulation
unknown function
Defence