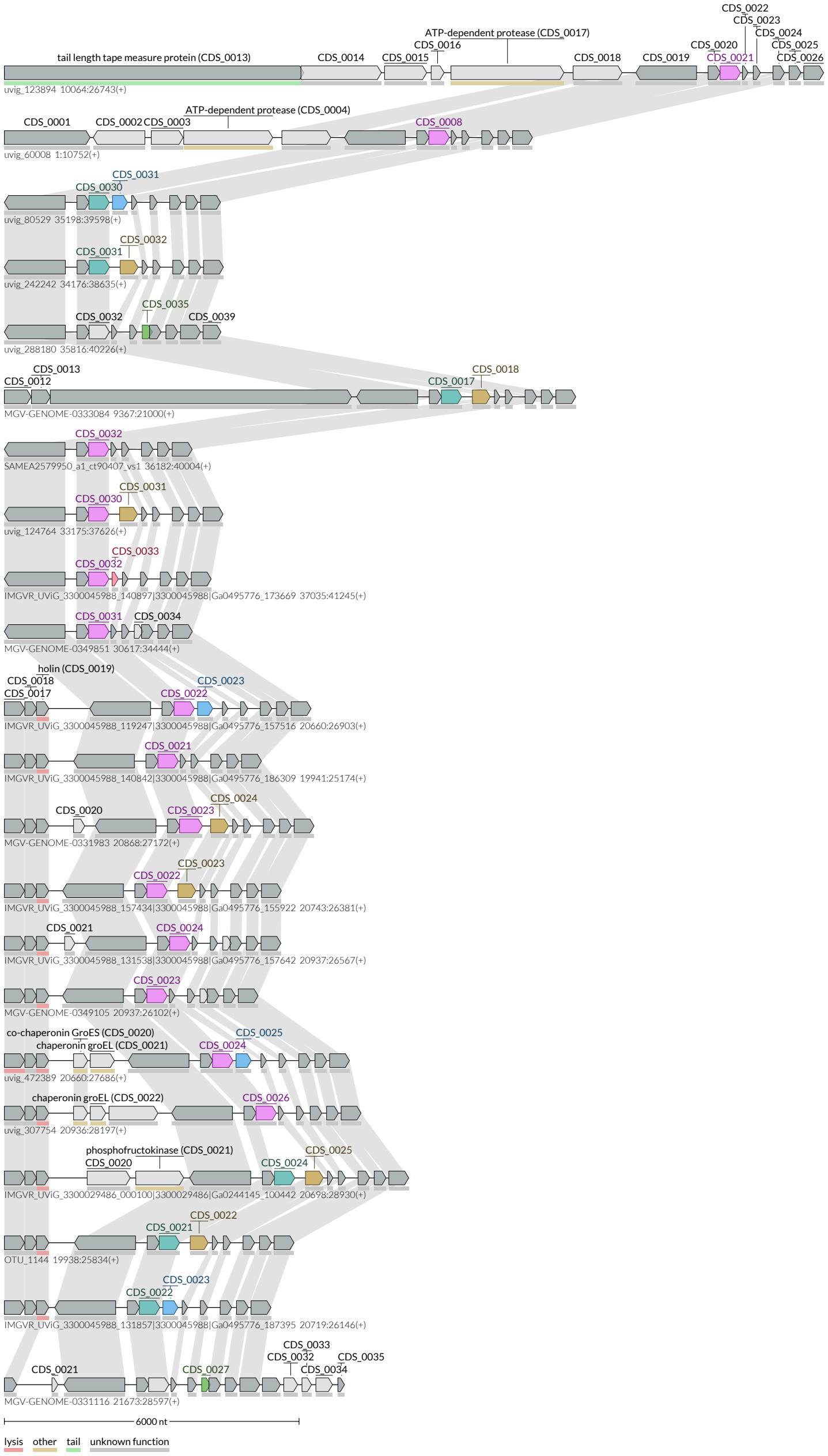
CDS_0022
CDS_0023
ATP-dependent protease (CDS_0017)
CDS_0024
CDS_0016
CDS_0020
CDS_0025
tail length tape measure protein (CDS_0013)
CDS_0014
CDS_0015
CDS_0018
CDS_0019
CDS_0021
CDS_0026
uvig_123894 10064:26743(+)
ATP-dependent protease (CDS_0004)
CDS_0001
CDS_0002
CDS_0003
CDS_0008
uvig_60008 1:10752(+)
CDS_0031
CDS_0030
uvig_80529 35198:39598(+)
CDS_0032
CDS_0031
uvig_242242 34176:38635(+)
CDS_0035
CDS_0032
CDS_0039
uvig_288180 35816:40226(+)
CDS_0013
CDS_0012
CDS_0018
CDS_0017
MGV-GENOME-0333084 9367:21000(+)
CDS_0032
SAMEA2579950_a1_ct90407_vs1 36182:40004(+)
CDS_0031
CDS_0030
uvig_124764 33175:37626(+)
CDS_0033
CDS_0032
IMGVR_UViG_3300045988_140897|3300045988|Ga0495776_173669 37035:41245(+)
CDS_0031
CDS_0034
MGV-GENOME-0349851 30617:34444(+)
holin (CDS_0019)
CDS_0018
CDS_0017
CDS_0023
CDS_0022
IMGVR_UViG_3300045988_119247|3300045988|Ga0495776_157516 20660:26903(+)
CDS_0021
IMGVR_UViG_3300045988_140842|3300045988|Ga0495776_186309 19941:25174(+)
CDS_0024
CDS_0020
CDS_0023
MGV-GENOME-0331983 20868:27172(+)
CDS_0023
CDS_0022
IMGVR_UViG_3300045988_157434|3300045988|Ga0495776_155922 20743:26381(+)
CDS_0021
CDS_0024
IMGVR_UViG_3300045988_131538|3300045988|Ga0495776_157642 20937:26567(+)
CDS_0023
MGV-GENOME-0349105 20937:26102(+)
co-chaperonin GroES (CDS_0020)
chaperonin groEL (CDS_0021)
CDS_0025
CDS_0024
uvig_472389 20660:27686(+)
chaperonin groEL (CDS_0022)
CDS_0026
uvig_307754 20936:28197(+)
phosphofructokinase (CDS_0021)
CDS_0020
CDS_0025
CDS_0024
IMGVR_UViG_3300029486_000100|3300029486|Ga0244145_100442 20698:28930(+)
CDS_0022
CDS_0021
OTU_1144 19938:25834(+)
CDS_0023
CDS_0022
IMGVR_UViG_3300045988_131857|3300045988|Ga0495776_187395 20719:26146(+)
CDS_0033
CDS_0032
CDS_0035
CDS_0021
CDS_0027
CDS_0034
MGV-GENOME-0331116 21673:28597(+)
6000 nt
lysis
other
tail
unknown function