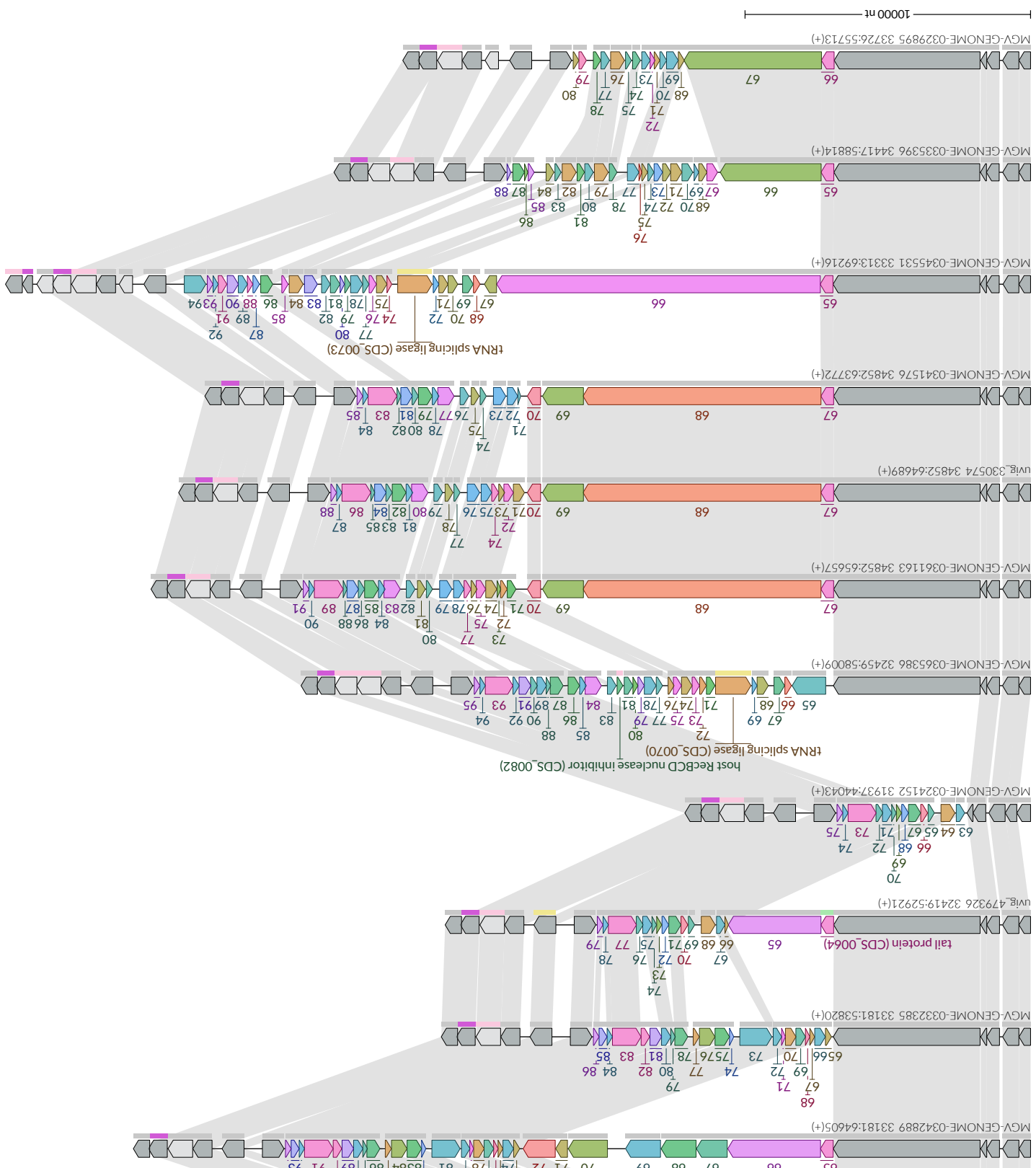
uvig_225646 31399:51812(+)
queuine tRNA-ribosyltransferase (CDS_0074)
CBASS_QueC (CDS_0075)
CDS_0067
CDS_0068
CDS_0066
CDS_0065
CDS_0071
MGV-GENOME-031690B 32875:47467(+)
CDS_0065
CDS_0061
CDS_0062
MGV-GENOME-031014I 31499:44305(+)
CDS_0061
MGV-GENOME-030944A 31499:44030(+)
CDS_0065
CDS_0063
CDS_0064
CDS_0066
MGV-GENOME-032404J 32220:52878(+)
CDS_0063
CDS_0064
CDS_0065
uvig_225235 31709:49101(+)
64
MGV-GENOME-031516I 32432:47515(+)
DpdL_QueD (CDS_0072) QueE-like radical SAM domain (CDS_0073)
MGV-GENOME-031975G 31753:50778(+)
CDS_0056
CDS_0063
CDS_0062
CDS_0061
CDS_0064
MGV-GENOME-029144S 31271:41321(+)
tail protein (CDS_0064)
MGV-GENOME-033568Z 32505:58994(+)
MGV-GENOME-032786I 32225:57116(+)
MGV-GENOME-033038J 32220:55974(+)
MGV-GENOME-032370J 33125:55161(+)
DpdL_FolE (CDS_0078)
uvig_579859 32027:54095(+)
tail protein (CDS_0064)
uvig_97782 31680:41075(+)
CDS_0062
CDS_0063
CDS_0067
MGV-GENOME-031912I 33214:48450(+)
HNH endonuclease (CDS_0068)
uvig_329538 33214:51409(+)
GTP cyclohydrolase (CDS_0077)
HNH endonuclease (CDS_0069)
uvig_452545 33497:58606(+)
tail protein (CDS_0063)
MGV-GENOME-032081S 35501:53210(+)
CDS_0065
CDS_0066
queuine tRNA-ribosyltransferase (CDS_0069)
CDS_0070
uvig_216805 32459:47032(+)
CDS_0065
CDS_0066
MGV-GENOME-031342O 31443:46495(+)
CDS_0066
CDS_0065
CDS_0067
uvig_378557 31443:48941(+)
MGV-GENOME-031702J 33181:46568(+)
CDS_0062
uvig_340447 33181:50791(+)
CDS_0062
CDS_0063
CDS_0064
CDS_0065
CDS_0067
MGV-GENOME-032765I 32225:54463(+)
62
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66
MGV-GENOME-032694S 34425:50307(+)
71
72
CDS_0070
uvig_225646 31399:51812(+)
58
59
60
61
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CBASS_QueC (CDS_0075)
queuine tRNA-ribosyltransferase (CDS_0074)
MGV-GENOME-033035S 32525:55948(+)
uvig_452545 33497:58606(+)
tail protein (CDS_0063)
HNH endonuclease (CDS_0069)
GTP cyclohydrolase (CDS_0077)
uvig_329538 33214:51409(+)
HNH endonuclease (CDS_0068)
MGV-GENOME-031912I 33214:48450(+)
uvig_216805 32459:47032(+)
queuine tRNA-ribosyltransferase (CDS_0069)
CDS_0065
CDS_0066
CDS_0070
MGV-GENOME-032081S 35501:53210(+)
MGV-GENOME-032444G 32563:51119(+)
SAM-dependent methyltransferase (CDS_0069)
CDS_0066
MGV-GENOME-032404J 32220:52878(+)
CDS_0063
CDS_0064
CDS_0065
uvig_225235 31709:49101(+)
CDS_0063
CDS_0064
CDS_0065
MGV-GENOME-031690B 32875:47467(+)
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CDS_0065
CDS_0071
MGV-GENOME-031014I 31499:44305(+)
CDS_0061
CDS_0062
MGV-GENOME-030944A 31499:44030(+)
CDS_0061
MGV-GENOME-031702J 33181:46568(+)
CDS_0062
uvig_378557 31443:48941(+)
MGV-GENOME-031342O 31443:46495(+)
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CDS_0065
uvig_216805 32459:47032(+)
CDS_0065
CDS_0066
uvig_41662 31499:46697(+)
DpdL_QueD (CDS_0072) QueE-like radical SAM domain (CDS_0073)
MGV-GENOME-031516I 32432:47515(+)
MGV-GENOME-031975G 31753:50778(+)
CDS_0056
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CDS_0062
CDS_0061
CDS_0064
MGV-GENOME-029144S 31271:41321(+)
uvig_97782 31680:41075(+)
CDS_0062
CDS_0063
CDS_0067
uvig_579859 32027:54095(+)
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MGV-GENOME-032370J 33125:55161(+)
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MGV-GENOME-033038J 32220:55974(+)
MGV-GENOME-033568Z 32505:58994(+)
MGV-GENOME-032786I 32225:57116(+)
MGV-GENOME-033035S 32525:55948(+)
MGV-GENOME-032444G 32563:51119(+)
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CDS_0065
uvig_225235 31709:49101(+)