context_tree

2024-10-07

Genomic context AMP_138200_c2_g1_i1

Anaerotignum propionicum DSM 1682

CP014223.1

Anaerotignum sp. MB30–C6 chromosome

Gene

- 5,10–methylene–tetrahydropholate dehydrogenase
- AAA–like domain protein
- agnoprotein
- AMP
- Baseplate J like protein
- copper amine oxidase N–terminal domain–containing protein
- DUF1643 domain–containing protein
- helix–turn–helix transcriptional regulator
- Helix–turn–helix XRE–family like protein
- hypothetical protein
- Integrase
- L,D–transpeptidase
- Protein translocase subunit SecY, Protein, membrane protein, PROTEIN TRANSPORT–IMMUNE
- RDD family
- recombinase family protein
- ribonuclease R
- SMI1 / KNR4 family protein
- SmpB protein
- SprT–like domain–containing protein Spartan/DNA Complex repair, protease, DNA BINDING
- SsrA–binding protein
- SsrA–binding protein SmpB
- Sulfotransferase 4A1
- tail protein
- type I restriction–modification system methyltransferase

CP133078.1

Caudoviricetes sp. isolate ctzDR1

BK058526.1

Caudoviricetes sp. isolate ctJ1L4

BK026787.1