

context_tree

2024-10-07

Genomic context AMP_143020_c3_g1_i1

Anaerotignum propionicum DSM 1682



Anaerotignum sp. MB30-C6 chromosome



Caudoviricetes sp. isolate ctzDR1



Caudoviricetes sp. isolate ctJ1L4



| Gene | |
|------|---|
| | AMP |
| | bifunctional methylenetetrahydrofolate dehydrogenase/methenyltetrahydrofolate cyclohydrolase FOLD |
| | c("L", "D-transpeptidase") |
| | c("phage/plasmid primase", " P4 family") |
| | Copper amine oxidase N-terminal domain-containing protein |
| | DUF1643 domain-containing protein |
| | DUF6273 domain-containing protein |
| | helix-turn-helix transcriptional regulator |
| | Helix-turn-helix transcriptional regulator |
| | hypothetical protein |
| | N-6 DNA methylase |
| | Phage tail protein |
| | Protein-export membrane protein SecG |
| | RDD domain-containing protein |
| | Recombinase family protein |
| | Ribonuclease R |
| | SIR2-like domain-containing protein |
| | site-specific integrase |
| | SMI1 / KNR4 family (SUKH-1) |
| | SprT-like domain-containing protein |
| | SsrA-binding protein |
| | tyrosine-type recombinase/integrase |
| | Uncharacterized protein |