

## context tree

2024-10-07

Genomic context AMP\_138200\_c2\_g1\_i1

Caudoviricetes sp. isolate ctiL05

BK033981.1 

Caudoviricetes sp. isolate ctvmC44

BK025808.1  holin PGRP protein hypothetical protein

Caudoviricetes sp. isolate ctlnE22

BK026979.1  PGRP protein GDSL like Lipase Acylhydrolase

Caudoviricetes sp. isolate ctUyu4

BK027835.1 
























diales bacterium isolate KR001\_HIC

CP107216.1

Caudoviricetes sp. isolate ctIN07

BK038133.1 

## Gene

- |   |  |
|---|--|
|    | adaptor protein MecA   |
|    | DNA-packaging protein small subunit  |
|    | DnaJ-like protein  |
|    | FeoB-associated Cys-rich membrane protein  |
|    | GDSL like Lipase Acylhydrolase   |
|    | GROUP II INTRON-ENCODED PROTEIN LTRA/RNA, GROUP II INTRONS, RIBONUCLEOPROTEIN.8A |
|   | Histone chaperone  |
|  | holin  |
|  | hypothetical protein   |
|  | Integrase  |
|  | MBL fold metallo-hydrolase   |
|  | MBOAT family protein   |
|  | N-acetylmuramoyl-L-alanine amidase   |
|  | NUDIX hydrolase  |
|  | peptidyl-prolyl cis-transisomerase   |
|  | peptidylprolyl isomerase   |
|  | PGRP protein   |
|  | Regulatory protein   |
|  | Short C-terminal domain  |
|  | SmpB protein   |
|  | SsrA-binding protein SmpB  |
|  | UDP-N-acetylglucosamine 1-carboxyvinyltransferase                                |
|  | vacuolar protein sorting-associated protein                                      |