



- [K] ENOG410825M - repressor LexA
- [L] ENOG4105C68 - recombinase RecA
- [L] ENOG4105DCH - radical SAM protein
- [L] ENOG4105ZS2 - RecA domain protein
- [L] ENOG4107YKW - nucleotidyltransferase/DNA polymerase involved in DNA repair
- [S] ENOG4105ES7 - putative DNA modification/repair radical SAM protein
- [L] ENOG411DNGJ - error-prone DNA polymerase
- [L] ENOG4105T12 - phage SPO1 DNA polymerase-related protein
- [L] ENOG4107SFH - DNA-directed DNA polymerase
- [L] ENOG410QKP1 - helicase
- [O] ENOG4108Z18 - hemin transporter
- [O] ENOG4108HW8 - peptide-methionine (S)-S-oxide reductase
- hypothetical protein
- [S] ENOG4108I35 - hypothetical protein
- [E] ENOG4107S55 - aminopeptidase
- hypothetical protein
- [V] ENOG41061TG - HNH endonuclease
- [P] ENOG4107EER - sulfite reductase flavoprotein subunit
- [L] ENOG411DGUH - excinuclease ABC subunit A
- [L] ENOG4107QS5 - recombinase RecQ
- [S] ENOG4107SRQ - hypothetical protein
- [K] ENOG410GBSD - GntR family transcriptional regulator
- [L] ENOG4105CCW - excinuclease ABC subunit B
- [L] ENOG4105CQ3 - DNA polymerase IV
- hypothetical protein