

Predicted Antitoxin (Blue Pentagon) Domain ID: D29  
RefSeq Protein Description: ATP-binding protein  
TADB Annotation: No Hit  
NCBI-CDD Annotation: AAA\_21  
HHpred(Pfam) Annotation: AAA\_21 ; AAA domain, putative AbiEii toxin, Type IV TA system (98.89%)  
HHpred(PDB) Annotation: DNA double-strand break repair rad50 (95.83%)

Predicted Toxin (Red Pentagon) Domain ID: M64  
-- RefSeq Protein Description: hypothetical protein  
-- TADB Annotation: No Hit  
-- NCBI-CDD Annotation: No Hit  
-- HHpred(Pfam) Annotation: No Hit  
-- HHpred(PDB) Annotation: No Hit

Acholeplasma multilocale ATCC 49900 (TA-Like pair: WP\_162479492.1 & WP\_162479493.1)  
Mesoplasma florum W37 (TA-Like pair: WP\_023025660.1 & WP\_023025662.1)



for the description of flanking genes, please check this link: [http://130.235.240.53/NetFlax/Mini\\_pdfs/M64.txt](http://130.235.240.53/NetFlax/Mini_pdfs/M64.txt)