

ed Toxin (Red Pentagon) Domain ID: D59

eq Protein Description: putative toxin-antitoxin system toxin component

B Annotation: No Hit

l-CDD Annotation: PIN_SF superfamily

red(Pfam) Annotation: PIN_9 ; PIN like domain (96.80%)

red(PDB) Annotation: Uncharacterized protein TTHA0540 (97.56%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M112

erisePIN family Description: AAA family ATPase

-- TADB Annotation: No Hit

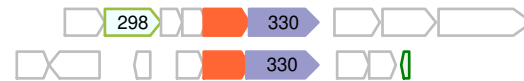
-- NCBI-CDD Annotation: P-loop_NTPase superfamily_AAA

-- HHpred(Pfam) Annotation: Pox_A32 ; Poxvirus A32 protein (98.98%)

HHpred(PDB) Annotation: Genome packaging NTPase B204 (99.25%)

thermus ruber DSM 1279 (TA-Like pair: WP_013013699.1 & WP_013013700.1)

mus hydrothermalis DSM 14884 (TA-Like pair: WP_013703332.1 & WP_013703333.1)



ion of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini_pdfs/M112.txt