Predicted Antitoxin (Blue Pentagon) Domain ID: D25

- -- RefSeq Protein Description: ImmA/IrrE family metallo-endopeptidase
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: Peptidase\_M78
- -- HHpred(Pfam) Annotation: Peptidase M78; IrrE N-terminal-like domain (99.11%)
- -- HHpred(PDB) Annotation: IRRE PROTEIN (99.28%)

Predicted Toxin (Red Pentagon) Domain ID: M56

- -- RefSeg Protein Description: hypothetical protein
- -- TADB Annotation: No Hit
- -- NCBI-CDD Annotation: No Hit
- -- HHpred(Pfam) Annotation: PIN 12; PIN domain (57.29%)
- -- HHpred(PDB) Annotation: hypothetical protein PH0500 (70.74%)

Chamaesiphon minutus PCC 6605 (TA-Like pair: WP\_015158718.1 & WP\_015158719.1)

Thioflexothrix psekupsii (TA-Like pair: WP\_086487771.1 & WP\_086487770.1)

<sup>1 237 10 10</sup> 

<sup>\*\*</sup> For the description of flanking genes, please check this link: http://130.235.240.53/NetFlax/Mini\_pdfs/M56.txt