

Predicted Toxin (Red Pentagon) Domain ID: D59
-- RefSeq Protein Description: putative toxin-antitoxin system toxin component, PIN family
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: PIN_SF superfamily
-- HHpred(Pfam) Annotation: PIN_9 ; PIN like domain (96.80%)
-- HHpred(PDB) Annotation: Uncharacterized protein TTHA0540 (97.56%)

Predicted Antitoxin (Blue Pentagon) Domain ID: M121
-- RefSeq Protein Description: hypothetical protein
-- TADB Annotation: No Hit
-- NCBI-CDD Annotation: mnt superfamily
-- HHpred(Pfam) Annotation: ParD ; Antitoxin ParD (95.14%)
-- HHpred(PDB) Annotation: PIN domain, RHH protein, DNA (96.14%)

Botrimarina mediterranea (TA-Like pair: WP_197529382.1 & WP_197529383.1)
Lacipirellula parvula (TA-Like pair: WP_172992133.1 & WP_152099924.1)
Lacipirellula parvula (TA-Like pair: WP_152099887.1 & WP_172992124.1)
Fimbrioglobus ruber (TA-Like pair: WP_088259594.1 & WP_088259593.1)
Posidoniimonas polymericola (TA-Like pair: WP_146589318.1 & WP_146589316.1)
Lacipirellula parvula (TA-Like pair: WP_152099811.1 & WP_152099812.1)

