```
logo
                                                  MKSSEMIRWLADGWYLVR GKGSHYKFKHPKKRGIVTYPHPKKELPIGTYRSLRKQAGLKLR
                                                  MPNTQKKIVKLKKQGWIRTIAYVGKGSHLQMTKNEPGTLTPPKKRDLKYFKGAIESLKLR
                                                  NFGDFRAALEEHVTEEAGTTLFHTEAQGEHQLTFADLGAGDMNEPLEKAIESLKLR
T10168 WP 001132284.1 NC 003028:c1704351-1704172  MPMTQKEMVKLLTAHGWIKTRGGKGSHIKMEKQGER.PITILH..GELNKYTERGIRKQAGL...   59
T6283 WP 003098540.1 NC 022808:c1463325-1463143   ..MKFSEFRRWLKAQGVTFEA.GKGSHFKITAPNGKQTTFADHGAKEMPEPTRKAIIKQLGLK..   60
T875 15965792 NC 003047:2199475-2199663           ..MKSGDIIAALQKDGWYEVA.TKGSHVQFKHPKKHGRVTVPHPKRDLPIGTLRSIEKQSGLKLR   62
T6107 WP 009893919.1 NC 006351:535977-536156      ..MNSSKLIRMLEEDGWRLVR.VTGSHHHFKHPKKPGLVTVPHPKKDLPIGTVKSIQKSAGL...   59
consensus                                          !********! **!*  *  **!!!********** ****!********!***! !**!!*
```

X non conserved

X similar

X $\geq$ 50% conserved