```
logo
T10122 AZV00182.1 MK291441:c9463-9056          ..MRYVAFLHTDEAGGFGISFPDFPGAISDGDTVEQAIQRGERALAFHVQGMREDG.LDVPAPRSVDDILADPDLAEWRE   77
T10168 WP 001132284.1 NC 003028:c1704351-1704172 MPMTQKEMVKLLTAHGWIK................TRGGKG...SHIKMEKQGE.RPITILH..........GELNKYTE   50
T6283 WP 003098540.1 NC 022808:c1463325-1463143 ..MKFSEFRRWLKAQGVTF................EAG.KG...SHFKITAPNGKQTTFADHGA........KEMPEPTR   50
T875 15965792 NC 003047:2199475-2199663        ..MKSGDIIAALQKDGWYE................VAT.KG...SHVQFKHPKKHGRVTVPHPK........RDLPIGTL   50
T6107 WP 009893919.1 NC 006351:535977-536156    ..MNSSKLIRMLEEDGWRL................VRV.TG...SHHHFKHPKKPGLVTVPHPK........KDLPIGTV   50
T3593 YP 001085049.1 NC 009085:c2346189-2346031 ..........MIEADGWYE................VRV.SG...SHHHFKHPTKKGLVTIPHPK........KDLPNGTV   42
consensus                                         *     *  * * !*                  *!   **     *   * *******        ****  *
```

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logo
T10122 AZV00182.1 MK291441:c9463-9056          GAQIAHVALILDRGSPKRVNISLDPGLLDAIDAEAARRGMTRSAFLSSAARAEIHAAH  135
T10168 WP 001132284.1 NC 003028:c1704351-1704172 RGIRKQAGL.....................................................   59
T6283 WP 003098540.1 NC 022808:c1463325-1463143 KAIIKQLGLK....................................................   60
T875 15965792 NC 003047:2199475-2199663         RSIEKQSGLKLR...................................................   62
T6107 WP 009893919.1 NC 006351:535977-536156    KSIQKSAGL......................................................   59
T3593 YP 001085049.1 NC 009085:c2346189-2346031 KSILKQAGLN.....................................................   52
consensus                                       * * ** *!
```

X non conserved

X similar

X ≥ 50% conserved