```
logo

NTDB id 1103 NMB RS07590 WP 002212976.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1       ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 1105166 ACNVED RS01675 WP 419420031.1 .MARGINKVILIGNVGSDPEVRYMPNGNAVATLSIATSETWKDKQSGDKQERTEWHRVVCFNRLGEIAGEFVRKGSKLYI   79
NTDB id 1390 A4U84 RS00055 WP 021115958.1      ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                          **!!!!**!**! !!**!*!!*!*!!*****!!!!*!*!* *!** !*!!!!***********!*!!****!!* *!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1        EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGYG..QSQEAYQRPAQQSRQPASDA....PSH   146
NTDB id 1131 NGFG RS05740 WP 003695064.1       EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGYG..QSQEAYQRPAQQSRQPAPDA....PSH   146
NTDB id 1166 A1552VC RS00795 WP 000168289.1    EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM...NVPAQQGSWGQP.QQPAKQH......Q   149
NTDB id 1105166 ACNVED RS01675 WP 419420031.1 EGSLRTRKWQDQQGQDRYTTEIIAT....DVQMLDSKGGSS........VGYDDAQSAA.AP.QHQQQPAYQ........   137
NTDB id 1390 A4U84 RS00055 WP 021115958.1      EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGD.FGGNQGSGWNQAPAQT.NYNQGGYSDNYAQNNNFNGG   152
consensus                                      !!*****!!*!** !**!*******    ***!!***  ***  *   * *  * * *   **!    ***

logo

NTDB id 1103 NMB RS07590 WP 002212976.1        PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1       PQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1    PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1105166 ACNVED RS01675 WP 419420031.1 .....QPT.NRKGTADDVFAELDDDIPF   159
NTDB id 1390 A4U84 RS00055 WP 021115958.1      NATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                      *   ***  *    *  *****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved