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 1390 A4U84 RS00055 WP 021115958.1         ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 111988 LOA RS01875 WP 025384900.1         .MARGINKVILIGNVGVDPEVRYMPNGNAVTTLSIATSETWKDKQSGDKQERTEWHRVVCFNRLGEIAGEYVRKGSKLYV   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1       MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                             **!!!!**!**! !!**!*!!*!*!! ****!!!!*!*!* *!** !*!!!!*********!*!!*!**!!* *!*
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NTDB id 1103 NMB RS07590 WP 002212976.1           EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYEEGYGQSQEAYQR.....PAQQSRQPASDAPSHP   147
NTDB id 1131 NGFG RS05740 WP 003695064.1          EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAPYDEGYGQSQEAYQR.....PAQQSRQPAPDAPSHP   147
NTDB id 1390 A4U84 RS00055 WP 021115958.1         EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGG.DFGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN   153
NTDB id 111988 LOA RS01875 WP 025384900.1         EGSLRTRKWQDQQGQERYTTEIVASDI....QMLDGKGQA..........NY...PEE....QSMSSQNTKTA.....KP   133
NTDB id 1166 A1552VC RS00795 WP 000168289.1       EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG.GMPAQGGMNV...PAQ....QGSWGQPQQPAKQ..HQP   150
consensus                                         !!****!!*!** !**!********     **!!***  *  * *   * *  * * *    *    *  ***      *
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logo

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NTDB id 1103 NMB RS07590 WP 002212976.1           QEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1          QEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1         ATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 111988 LOA RS01875 WP 025384900.1         .........SQGVAQDAFEELDDDIPF   151
NTDB id 1166 A1552VC RS00795 WP 000168289.1       MQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                            **   *    *  *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved