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logo
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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 987825 AAGU62 RS15750 WP 015872740.1  MASRGVNKVILIGNLGQDPEVRYMPNGGAVANLTLATSDSWRDKQTGEQKERTEWHRVALYGKLAEIAGEYLRKGSQVYI  80
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI  80
consensus                                        **!!!!**!*!!*!!**!*!!*!*!!*!!*****!!!**!*!* *!***!*!!!!*********!!*!!*!!*!!*  !!*
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logo
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NTDB id 1103 NMB RS07590 WP 002212976.1       EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAP........YEEGYGQSQEAY..QRP......AQQS  136
NTDB id 1131 NGFG RS05740 WP 003695064.1      EGRIQSRKYQGKDGIERTAYDIVAN....EMKMLGGRNENSGGAP........YDEGYGQSQEAY..QRP......AQQS  136
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFGGN.......QGSGWNQAPAQTNYNQGGYSDNYAQNN  147
NTDB id 987825 AAGU62 RS15750 WP 015872740.1  EGQLRTRKWQDQSGQDRYTTEVVVD.ISGSMQMLGGRQGGAP......MGNQGQQGGWGQPQQPQ.............GA  140
NTDB id 1166 A1552VC RS00795 WP 000168289.1   EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPA.............KQ  147
consensus                                     !!*****!!*!** !**!********     **!!**!* *** *         *  ***! *     *         ***
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logo
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NTDB id 1103 NMB RS07590 WP 002212976.1       RQPASDA.PS..HPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF  174
NTDB id 1131 NGFG RS05740 WP 003695064.1      RQPAPDA.PS..HPQEAPAAPRRQ.PVPAAAPVEDIDDDIPF  174
NTDB id 1390 A4U84 RS00055 WP 021115958.1     NFNGGNAT.........RPQPAQKPAAQAEPPMDNFDDDIPF  180
NTDB id 987825 AAGU62 RS15750 WP 015872740.1  AFSGGQAAPRQQSQQGGYSAPAQPAAPQSSEPPMDFDDDIPF  182
NTDB id 1166 A1552VC RS00795 WP 000168289.1   HQP............MQQSAPQQYSQPQYNEPPMDFDDDIPF  177
consensus                                      ***  * *     * *  *! *    **  !****!!!!!!
```

X non conserved

X similar

X ≥ 50% conserved