```
logo
                                                  ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1
NTDB id 1103 NMB RS07590 WP 002212976.1          ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1        ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 297350 SM212138 RS22295 WP 000157067.1   MASKGVNKVILVGNLGQDPEVRYMPNGGAVVNLSLATSDTWTDKQTGDKKERTEWHRVVLYGKLAEIASEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1      MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                            **!!!!**!*!!*!!**!*!!*!*!!  ****!!!**!*!* *!***!*!!!!********!!*!**!!*!!*  !!*

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1         EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYDEGY..GQSQEAYQRPAQQSR........QPAPD   142
NTDB id 1103 NMB RS07590 WP 002212976.1          EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPYEEGY..GQSQEAYQRPAQQSR........QPASD   142
NTDB id 1390 A4U84 RS00055 WP 021115958.1        EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDF.GG.......NQGSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 297350 SM212138 RS22295 WP 000157067.1   EGALRTRKWTDQSGVEKYTTEVVVS.QSGTMQMLGGRNSAGS..........GQQQGGWGQPQQPAA.............   136
NTDB id 1166 A1552VC RS00795 WP 000168289.1      EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGM.PAQGGMNVPAQQGSWGQPQQPAK.............   146
consensus                                        !!*****!!**** !*********     *  **!!**!* ***  *   *   **** ***** ****        * *

logo

NTDB id 1131 NGFG RS05740 WP 003695064.1         APSHP..QEAPAAPRRQPVPAAAPVEDIDDDIPF................   174
NTDB id 1103 NMB RS07590 WP 002212976.1          APSHP..QEAPAAPRRQPVPAAAPVEDIDDDIPF................   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1        NNFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF................   180
NTDB id 297350 SM212138 RS22295 WP 000157067.1   .PSHSG...MP....PQQHPANEPPMDFDDDIPFAPFGHSVARHALYVLS   178
NTDB id 1166 A1552VC RS00795 WP 000168289.1      .QHQPMQQSAPQ..QYSQPQYNEPPMDFDDDIPF................   177
consensus                                         ****  * *!    **  **  !****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved