```
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 332840 EKN56 RS16525 WP 130592791.1 MASRGINKVILVGNLGQDPEIRYMPNGGAVANITIATSESWRDKQTGEQKEKTEWHRVVIFGKLAEIAGEYLRKGSQVYI   80
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                       **!!!!**!*!!*!!**!*!!*!*!!*****!!!!*!*!* *!***!*!!!!*****!!*!!*!!*!!* !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPY........EEGYGQSQEA..YQRPAQQSRQPAS.   141
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIV....ANEMKMLGGRNENSGGAPY........DEGYGQSQEA..YQRPAQQSRQPAP.   141
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQ....GDVLQMLDSRSSGGDFGGNQG.......SGWNQAPAQTNYNQGGYSDNYAQNN   147
NTDB id 332840 EKN56 RS16525 WP 130592791.1 EGSLQTRKWQDQSGQDRYTTEVVV.NIGGVMQMLGGRQGG........GEGQSAQGSWGQPQQP................   135
NTDB id 1166 A1552VC RS00795 WP 000168289.1 EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQP................   144
consensus                                   !!*****!!*!** !**!******    * ***!!**!* * * *            ***! *     **  *

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    NFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 332840 EKN56 RS16525 WP 130592791.1 QQSQAQASRPAQQQSAPMQQNEPPMDFDDDIPF   168
NTDB id 1166 A1552VC RS00795 WP 000168289.1 AKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                         *   **  ** ****  !****!!!!!!
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X non conserved

X similar

X $\geq$ 50% conserved