```
logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1103 NMB RS07590 WP 002212976.1         ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
NTDB id 1099852 ACLNGM RS15820 WP 090959229.1  .MAGSVNKVILVGNLGADPEIKRLNSGDAIANLRIATSETWRDKASGERKERTEWHSVVIFNENLVKVAEQYLKKGSTVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1    MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
consensus                                          **!!!!**!*!! !!*******!*!******!!!!*!*!* *!***!*!!!! ***** *****!**!!*!!* !!

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGY.GQ............SQEAYQRPAQQS..   136
NTDB id 1103 NMB RS07590 WP 002212976.1         LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGY.GQ............SQEAYQRPAQQS..   136
NTDB id 1390 A4U84 RS00055 WP 021115958.1       VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQT.NYNQGGYSDNYAQNNNFNGG   152
NTDB id 1099852 ACLNGM RS15820 WP 090959229.1  IEGQLQTRSWEDQQGQKRYTTEVVLQRFRGELQMLGGRGGDGGGPSYDGGG..E.RSGGRSGGGGD....YGRGG....G   148
NTDB id 1166 A1552VC RS00795 WP 000168289.1    IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMN..V.PAQ.....QGS....WGQPQ....Q   143
consensus                                      *!!****!***** !**!******     ***!!**!  ***** ****   *           *   * ******

logo
NTDB id 1131 NGFG RS05740 WP 003695064.1        ..RQPAPDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1         ..RQPASDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NATRPQPAQ............KPAAQAEPPMDNFDDDIPF   180
NTDB id 1099852 ACLNGM RS15820 WP 090959229.1  GAGGR...SSGGGGGY.....GGGGGGGRPSDDMDDEIPF   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1    PAKQHQPMQQSAPQQY.....S.QPQYNEPPMDFDDDIPF   177
consensus                                       **** *   * **       *    ** !****!!*!!!
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X non conserved

X similar

X $\geq$ 50% conserved