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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1   ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1103 NMB RS07590 WP 002212976.1    ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY   75
NTDB id 1390 A4U84 RS00055 WP 021115958.1  ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY   77
NTDB id 1216449 SMB554 RS08480 WP 003529546.1 .MAGSVNKVILIGNVGADPEIRRTQDGRPIANLRIATSETWRDRNSGERREKTEWHTVVVFNEGLCKVVEQYVKKGAKLY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY   79
consensus                                      **!!!!**!**!  !!**!****!********!!!!*!*!***!***!*!!!!  *****  ********!**!!* *!
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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1   LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY.DEGYGQ.......SQEAYQR...PAQQSRQPA   140
NTDB id 1103 NMB RS07590 WP 002212976.1    LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY.EEGYGQ.......SQEAYQR...PAQQSRQPA   140
NTDB id 1390 A4U84 RS00055 WP 021115958.1  VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQGSGWNQAPAQTNYNQGGYSD...NYAQNNNFN   150
NTDB id 1216449 SMB554 RS08480 WP 003529546.1 IEGQLQTRKWQDQTGNDRYSTEVVLQGFNSTLTMLDGRGGEGGGAGRGGSDYGG.......G..GYEEYDQ....SRPSS   146
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPA.......QQGSWGQPQQPAKQHQPMQ   152
consensus                                  *!!****!!*!**  !**!*******       * !!***!  ******  * ****         * **      ** ****
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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1   PDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1    SDAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1  GGNATR....PQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1216449 SMB554 RS08480 WP 003529546.1 GGARSG....GQS..NQPNQGGNFSRDLDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 ...QSA....PQQ..YSQPQYNEPPMDFDDDIPF   177
consensus                                    * *      **    *** *** *****!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved