```
logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV    78
NTDB id 354262 E4T21 RS02050 WP 149283061.1  .MARGINKVILIGNLGQDPEVRFTPSGTAVANLNLATTDNWMDRQSGQRQERTEWHRVVLFNKLAEIAQQYVKKGAKLYI    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI    80
consensus                                       *!*!!!!**!!!!*!!**!**!*! !!!****!!***! !* *!***!*!!!!!**** **!!*!**!**!!***!*

logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1     EGKLKTRKWQDQNGQDRYTTEIQGD....VLQMLDSRSSGGDFGGN.....................QGSGWNQAPAQTN.   133
NTDB id 354262 E4T21 RS02050 WP 149283061.1  EGRLQTRKWQDQNGQDRYSTEIVANDM....QMLDGRGDGGQQAGGFGGGQQGGYGAQQGGGFGGGQQAAYGAAAPQQAP   155
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQG............GMPAQGGMN.VPAQQGSWG........   139
consensus                                    !!*!*!!!!!!!*!!!!!*!!****    **!!!**!* *!  **        *  ** *     *!***** *  *

logo
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..........YNQGGYSDNYAQNNNFNGGNATRPQPAQKPAA................QAEPPMDNFDDDIPF   180
NTDB id 354262 E4T21 RS02050 WP 149283061.1  APQQQSNFAGQQSGGFSG.......QQGGGYGGGQPAQQPQQPQQPSSGGQKGGFGAPD.PGSFDDFDDEIPF   220
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ................................QPQ...QPAKQHQPMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                              * ****          ***    *!****!*   **        *  **   * **!!!*!!!
```

X non conserved

X similar

X $\geq$ 50% conserved