```
logo
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYR.KLAEIAGQYLKKGGLVY    75
NTDB id 1148333 HDIA RS08120 WP 099555712.1 .MAGSVNKVILVGNLGADPEIRRTQDGRPIANLRVATSESWRDRNSGERREKTEWHRVVIFNEGLCKVAEQYLRKGAKVY    79
NTDB id 1166 A1552VC RS00795 WP 000168289.1 MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYG.KLAEVAGEYLRKGSQVY    79
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFR.RQAEVAGQYLRKGSKVY    77
consensus                                       **!!!!**!*!! !!**!****!********!!!!*!*!***!***!*!!!!*****  *****!**!!*!!*  !!

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY.EEGYG....QSQE.....AYQRPAQQSRQPAS   141
NTDB id 1131 NGFG RS05740 WP 003695064.1     LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY.DEGYG....QSQE.....AYQRPAQQSRQPAP   141
NTDB id 1148333 HDIA RS08120 WP 099555712.1 IEGQLQTREWDDQSGQKRYSTEIVLQGFNSTLTMLDGRGERGEGDFGGGSGGGSDFGRSSPM...................   141
NTDB id 1166 A1552VC RS00795 WP 000168289.1 IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG..MPAQGGM...NVPAQ....QGSWGQPQQPAK....   146
NTDB id 1390 A4U84 RS00055 WP 021115958.1   VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD..FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNN....   147
consensus                                   *!!****!***** !**!*******       * !!**! *****   *  ! *     **      **  * ** *

logo
NTDB id 1103 NMB RS07590 WP 002212976.1      DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1148333 HDIA RS08120 WP 099555712.1 ERSGGSSGRAASSTGGGGSRGGYSSDMDDEIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1 QHQ..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NFNGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
consensus                                   *    *   **   *    ********!!*!!!
```

X non conserved
X similar
X ≥ 50% conserved