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logo
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```
NTDB id 1131 NGFG RS05740 WP 003695064.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRK.LAEIAGQYLKKGGLVY   75
NTDB id 1390 A4U84 RS00055 WP 021115958.1     ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRR.QAEVAGQYLRKGSKVY   77
NTDB id 186740 KC8 RS12355 WP 029624217.1     .MAGSVNKVILVGNLGRDPESRSFQNGGKVVNLRIATSETWKDRATGERREKTEWHSVAIFNEGLARTAEQYLRKGSKVY   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1   MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGK.LAEVAGEYLRKGSQVY   79
consensus                                        * *!!!!**!*!!*!!**!****!**!*****!!!!*!*!***!***!*!!!!******* *!**!**!!*!!**!!
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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1      LEGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPY........DEGYGQSQEAYQRPAQQSRQPAPDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1     VEGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGGNQ.......GSGWNQAPAQTNYNQGGYSDNYAQN   146
NTDB id 186740 KC8 RS12355 WP 029624217.1     IEGQLQTRKWQDQSGADKYSTEIVLQGFNSQLVMLDGPGGGSGGGGRGGSGAGG.SSGWDEGDSGFGG..SGS.SGMSSG   155
NTDB id 1166 A1552VC RS00795 WP 000168289.1   IEGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPAKQ..HQ...PMQ..   152
consensus                                     *!!****!!*!***!********* **** ***!! *** * *******    *  ******* ***     *  **  *
```

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logo
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```
NTDB id 1131 NGFG RS05740 WP 003695064.1      PS....HPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1     NNFNG.GNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 186740 KC8 RS12355 WP 029624217.1     GGFGGGGGGGRSGGTGGGARPSNPFDSDLDDDVPF   190
NTDB id 1166 A1552VC RS00795 WP 000168289.1   ..........QSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                       * * * * **** ** * *********!!!*!!
```

X non conserved

X similar

X $\geq$ 50% conserved