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logo
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```
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 178036 I596 RS03815 WP 067644428.1   .MARGINKVILVGNLGADPETRYTASGAAITNIRLATSESWRDKQTGENQERTEWHRVVLFGKLGEIAGEYLKKGRQVYI   79
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                        **!!!!**!*!! !!**!****!*!* ****!!!!*!*!* *!** !*!!!!*********!*!!*!!*!!  !!*
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logo
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NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA   143
NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVANE....MKMLGGRNENSGGAPYEEGYGQS.........QEAYQRPAQQSRQPASDA   143
NTDB id 1390 A4U84 RS00055 WP 021115958.1    EGKLKTRKWQDQNGQDRYTTEIQGDV....LQMLDSRSSGGDFGGNQ.GSGWNQAPAQTNYNQGGYSDNYAQNNNFNGGN   153
NTDB id 178036 I596 RS03815 WP 067644428.1   EGSLRTNKYTDKDGVERYTTDIVANE....MQMLGGGGAGGGGEGGYGGGGS.....GGGGGGGYGRERGPARGAAP..   148
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPA...QGGMN.....VPAQQGSWGQPQQPAKQHQP..   150
consensus                                    !!*****!*****!**!*********    **!!***  *****    * ! *         **** ** ** ** **
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logo
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```
NTDB id 1131 NGFG RS05740 WP 003695064.1     PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1103 NMB RS07590 WP 002212976.1      PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1390 A4U84 RS00055 WP 021115958.1    ...ATRPQPAQ.KPAAQAEPPMDNFDDDIPF   180
NTDB id 178036 I596 RS03815 WP 067644428.1   ...ARSAPPAS.....SNQPPDTFEDDDIPF   171
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ...MQQSAPQQYSQPQYNEPPMDF.DDDIPF   177
consensus                                        * ****  ***  * *!**  !!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved