logo



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NTDB id 1131 NGFG RS05740 WP 003695064.1        ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 356365 Psal025 RS02720 WP 016212110.1   ..MRGVNKVIIVGRLGQDPDIRRTQSGESVATLSVATSEQWQDQ.HGVNHESTEWHRIVLFSGLANIAKDYLRKGAQIYV   77
NTDB id 1390 A4U84 RS00055 WP 021115958.1       ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
NTDB id 1166 A1552VC RS00795 WP 000168289.1     MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
consensus                                         ****!!!!**! !!*!!**!*** !**!* ***!!!!*!*!* *!***!*!!!!** *****!**!**!!*!!***!*
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NTDB id 1131 NGFG RS05740 WP 003695064.1        EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQS.........QEAYQRPAQQSRQPAPDA   143
NTDB id 356365 Psal025 RS02720 WP 016212110.1   EGKIKTRKWQDNQGKDRYTTEIVG....KNMQMLGGGERVQT.AK.......PPPPPPPSSLGQQAF..............   131
NTDB id 1390 A4U84 RS00055 WP 021115958.1       EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGD.FGGNQGSGWNQAPAQTNYNQGGYSDNYAQNNN....F   149
NTDB id 1166 A1552VC RS00795 WP 000168289.1     EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGG.MPAQGGM...NVPAQ....QGSWGQPQQPAKQ....H   148
consensus                                       !!** *!!*!***!**!*******      ***!!***  *** **** * *  * ****   !***  * ** **
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logo



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NTDB id 1131 NGFG RS05740 WP 003695064.1        PSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 356365 Psal025 RS02720 WP 016212110.1   ......KAKQAQLHAQNSSAEMEKIGEEIPF   156
NTDB id 1390 A4U84 RS00055 WP 021115958.1       NGGNATRPQPAQKPAAQAEPPMDNFDDDIPF   180
NTDB id 1166 A1552VC RS00795 WP 000168289.1     Q..PMQQSAPQQYSQPQYNEPPMDFDDDIPF   177
consensus                                       *  *  *******  * ** ***** ***!!!
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X non conserved

X similar

X ≥ 50% conserved