

logo

NTDB id 1103 NMB RS07590 WP 002212976.1	...MSLNKVKILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL	76
NTDB id 1131 NGFG RS05740 WP 003695064.1	...MSLNKVKILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNR.NGQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL	76
NTDB id 1390 A4U84 RS00055 WP 021115958.1	..MAGVNKVIIVGNLGNPDMDRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLKKGSKVYV	78
NTDB id 1003104 ABFR12 RS17220 WP 000168282.1	MASRGVNKVIILGRVGGDPEVRYSPSGTAFANLTVATSEQWRDKQTGEQKEQTEWHRVAVVGKLAEVVGRYVKKGDQVYF	80
NTDB id 1166 A1552VC RS00795 WP 000168289.1	MASRGVNKVIILGNLGDPEVRYMPSGGAVANITITATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYT	80
consensus	**!!!!!!**!*!!!!!!**!*!!!!!!**!*!!!!!!**!*!!!!!!**!*!!!!!!**!*!!!!!!**!*!!!!!!**!*!!!!!!	

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NTDB id 1103 NMB RS07590 WP 002212976.1	EGRIQSRKYQKGDGIERTAYDIVA....NEMKMLGGRNENSSGGAPYEEGYGQSSE.....AYQRPAAQSSRQPASDA	143
NTDB id 1131 NGFG RS05740 WP 003695064.1	EGRIQSRKYQKGDGIERTAYDIVA....NEMKMLGGRNENSSGGAPYEEGYGQSSE.....AYQRPAAQSSRQPAPDA	143
NTDB id 1390 A4U84 RS00055 WP 021115958.1	EGKLKTRKWQDQNGQDRYTTEIQQ....DVLQMLDSRSSGGD.FGGNQGSWNQAPAQTNYNQGGYSDNYAQNN....NF	149
NTDB id 1003104 ABFR12 RS17220 WP 000168282.1	EGMLRTRKWQDQGTGQDRYTTEIN.VGINGVMQMLGGTGDQSKQ.QAADR.....QS.....	128
NTDB id 1166 A1552VC RS00795 WP 000168289.1	EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQGGG.MPAQGGM..N.VPAQ....QGSWGQPQQPAK....QH	148
consensus	!!*****!!**!*!!!!*****	

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NTDB id 1103 NMB RS07590 WP 002212976.1	PSHPQEAAPAPRRQPVPAAPVEDI	174
NTDB id 1131 NGFG RS05740 WP 003695064.1	PSHPQEAAPAPRRQPVPAAPVEDI	174
NTDB id 1390 A4U84 RS00055 WP 021115958.1	NGGNATRPQPAQKPAQAEPMDNFDDDDIPF	180
NTDB id 1003104 ABFR12 RS17220 WP 000168282.1	QKPPQQQPSPTQHN	168
NTDB id 1166 A1552VC RS00795 WP 000168289.1	.QPMQQSAPQQYSQPQYN	177
consensus	* * *** ***!*****!!!!	

- non conserved
- similar
- ≥ 50% conserved