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logo

NTDB id 1103 NMB RS07590 WP 002212976.1      ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1131 NGFG RS05740 WP 003695064.1     ...MSLNKVILIGRLGRDPEVRYMPNGEAVCNFSVATSETWNDRN.GQRVERTEWHNITMYRKLAEIAGQYLKKGGLVYL   76
NTDB id 1166 A1552VC RS00795 WP 000168289.1  MASRGVNKVILIGNLGQDPEVRYMPSGGAVANITIATSETWRDKATGEQKEKTEWHRVTLYGKLAEVAGEYLRKGSQVYI   80
NTDB id 107489 RGQ30 RS14015 WP 130557627.1 ..MASVNKVILVGNLGKDPDVRYLPDGGAVVNLALATSSNWKDKNTGEKREETEWHRVVIYGRLAEIAGEYCKKGRSVYI   78
NTDB id 1390 A4U84 RS00055 WP 021115958.1   ..MAGVNKVIIVGNLGNDPDMRTMPNGDAVATLSVATSESWNDKMTGERREVTEWHRIVFFRRQAEVAGQYLRKGSKVYV   78
consensus                                       **!!!!**!*!!*!!**!**!*!*!! ****!!!**!*!***!***!*!!!!*****!!*!!*!**!!  !!*

logo

NTDB id 1103 NMB RS07590 WP 002212976.1      EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYEEGYGQ.SQEAYQRPAQQSRQPAS..........   141
NTDB id 1131 NGFG RS05740 WP 003695064.1     EGRIQSRKYQGKDGIERTAYDIVA....NEMKMLGGRNENSGGAPYDEGYGQ.SQEAYQRPAQQSRQPAP..........   141
NTDB id 1166 A1552VC RS00795 WP 000168289.1  EGQLQTRKWQDQSGQDRYSTEVVVQGYNGIMQMLGGRAQQGGMPAQGGMNVPAQQGSWGQPQQPAKQH............   148
NTDB id 107489 RGQ30 RS14015 WP 130557627.1 EGRLKTRKWQDQSGVDKYTTEIIA....DQMQLLGGRDGGGGGSSMGGG......DDFDQSPAPSRSSGG..........   138
NTDB id 1390 A4U84 RS00055 WP 021115958.1   EGKLKTRKWQDQNGQDRYTTEIQG....DVLQMLDSRSSGGDFGG.NQG......SGWNQAPAQTNYNQGGYSDNYAQNN   147
consensus                                   !!*****!!*!** !**********       ***!**!  *****    *      ******* ***** *

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NTDB id 1103 NMB RS07590 WP 002212976.1      ....DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1131 NGFG RS05740 WP 003695064.1     ....DAPSHPQEAPAAPRRQPVPAAAPVEDIDDDIPF   174
NTDB id 1166 A1552VC RS00795 WP 000168289.1  ......QPMQQSAPQ..QYSQPQYNEPPMDFDDDIPF   177
NTDB id 107489 RGQ30 RS14015 WP 130557627.1 NS.GGGSSMRS..SA..PAPQRSAPQSVADLDDDIPF   170
NTDB id 1390 A4U84 RS00055 WP 021115958.1   NFNGGNATRPQ..PA..QKPAAQAEPPMDNFDDDIPF   180
consensus                                        * * ** ***    *     *  *****!!!!!!
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X non conserved

X similar

X ≥ 50% conserved