logo

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NTDB id 1191 AAC28468.1 484..945( )          MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAGNVIFDS    80
NTDB id 1002 ACIAD RS15100 WP 004923779.1    ..MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTAGSSC...    75
NTDB id 1119 NGFG RS11465 WP 017147169.1     .MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGKWPENNTSAGVASP...    76
NTDB id 328362 EGH18 RS12470 WP 124693335.1  .MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNNGKWPEDNASAGVASP...    76
NTDB id 1109 OK783 RS10670 WP 157147416.1    .MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGIWPKDNTSAGVASA...    76
consensus                                    ****!!!!!!!!!!!!!!!!*!!!!!*!*!!!*!!!*!!***!********************  !  **  *  ******
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logo

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NTDB id 1191 AAC28468.1 484..945( )          ANGLDGKYFAPGGVTV......TP.....DTGAINIAFDAGANAGKTMTITPTVIAASGQISTWTCAPGA..........    139
NTDB id 1002 ACIAD RS15100 WP 004923779.1    ......VGVQETSASNATTNVATATCGASSAGQIIVTMDTTKAKCANITLTPTY...ASGAVTWKCTTTS..........    136
NTDB id 1119 NGFG RS11465 WP 017147169.1     PTDIKGKYVKEVEVKN...GVVTA........TMLSSGVNNEIKGKKLSLWGRR...ENGSVKWFCGQPVTRAD..DDTV    140
NTDB id 328362 EGH18 RS12470 WP 124693335.1  .ADIKGKYVQKVEVNN...GVVTA........QMASSNVNKEIQGKKLSLWAKR...EDGSVKWFCGQPVQRADAKADTV    141
NTDB id 1109 OK783 RS10670 WP 157147416.1    .STIKGKYVQKVEVAK...GVVTA........QMASTGVNKEIQDKKLSLWAKR...QDGSVKWFCGQPVTRTGDNDDTV    141
consensus                                       *******  ***  *    ***!*          **  **  **  **  **********      ****!*!****  *      ***
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logo

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NTDB id 1191 AAC28468.1 484..945( )          .....ANGIETRRLPTSCQ ......    153
NTDB id 1002 ACIAD RS15100 WP 004923779.1    .........DKKYVPSECRG......    147
NTDB id 1119 NGFG RS11465 WP 017147169.1     ADAKDGKEIDTKHLPSTCRDKASDAK    166
NTDB id 328362 EGH18 RS12470 WP 124693335.1  ADDKNNNGIDTKHLPSTCRDNFDAS.    166
NTDB id 1109 OK783 RS10670 WP 157147416.1    ADA..NNAIDTKHLPSTCRDESSAT.    164
consensus                                    **     ********!**!**
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X non conserved
X similar
X ≥ 50% conserved