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logo

NTDB id 1119 NGFG RS11465 WP 017147169.1      MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGKWPENNTS..........   70
NTDB id 222517 A6J44 RS01730 WP 082295363.1  MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGEWPKDNDS..........   70
NTDB id 1109 OK783 RS10670 WP 157147416.1    MNTLQKGFTLIELMIVIAIVGILAAVALPAYQDYTARAQVSEAILLAEGQKSAVTEYYLNHGIWPKDNTS..........   70
NTDB id 1188 NELON RS01755 WP 003772458.1    MKAMQKGFTLIELMIVIAIIGILAAIALPAYQDYTVRSKVSEALIAASSPKSALSEAFQTDGITGMTAAAKAFNKTAAAG   80
NTDB id 1002 ACIAD RS15100 WP 004923779.1    .MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTP.......STA.   71
consensus                                    ****!!!!!!!!!!!!!!!*!!!!!*!*!!!*!!!!*!**!!!****!***!****!*****! **   * *

logo

NTDB id 1119 NGFG RS11465 WP 017147169.1      ........AGVASPPTDIKGKYVKEVEVKNG...VVTATM...LSSGVNNEIKGKKLS........LWGRRENGSVKWFC  128
NTDB id 222517 A6J44 RS01730 WP 082295363.1  ........AGVASS.SSIKGKYVKEVKVENG...VVTATM...NSSNVNKEIKDKRLS........LWGRRENGSVKWFC  127
NTDB id 1109 OK783 RS10670 WP 157147416.1    ........AGVASA.STIKGKYVQKVEVAKG...VVTAQM...ASTGVNKEIQDKKLS........LWAKRQDGSVKWFC  127
NTDB id 1188 NELON RS01755 WP 003772458.1    GGAGGAAAAGTQHASK.....YVKSVKVTEASPWDTTVAISATEANGIPTGLDGNTIRLFPLVNKAVPAATSTGAIDWAC  155
NTDB id 1002 ACIAD RS15100 WP 004923779.1    ....GSSCVGVQEISASNATTNVATATCGASSAGQTIVTMDTTKA.......KGANI........TLTPTYASGAVTWKC  132
consensus                                            *!***  *******!* * *  *   ******    ***** ********        *****  !***!*!

logo

NTDB id 1119 NGFG RS11465 WP 017147169.1      GQPVTRADDDTVAD...AKDGKEIDTKHLPSTCRDKASDAK   166
NTDB id 222517 A6J44 RS01730 WP 082295363.1  GQPVTRNANDDTVTADGTGNDGKIDTKHLPSTCRDTSSAGT   168
NTDB id 1109 OK783 RS10670 WP 157147416.1    GQPVTRTGDNDDTVAD...ANNAIDTKHLPSTCRDESSAT.   164
NTDB id 1188 NELON RS01755 WP 003772458.1    ASAAKETADKRFGTGSGV.AVGTLPAKYAPSECK.......   188
NTDB id 1002 ACIAD RS15100 WP 004923779.1    TTTS....................DKKYVPSECRG......   147
consensus                                    ****** **     *         ***!**!!*!**  *
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X non conserved

X similar

X ≥ 50% conserved