```
logo
                                                  ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA   77
NTDB id 1184 GTF74 RS02335 WP 000649326.1
NTDB id 1167 A1552VC RS11065 WP 000649326.1   ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA   77
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    ...MKAYKNKQQKGFTLIELMIVVAVIGVLAAIAIPQYQNYVKKSAIGVGLANITALKTNIEDYIATEGSFPATTAGTAA   77
NTDB id 1197 PSJM300 03940 AFN76866.1         ......MKAQMQKGFTLIELMIVVAIIGILAAIALPAYQDYTVRSNAAAALAEITPGKIGFE.QAINEGKTPSLTSTDEG   73
NTDB id 1204 PSJM300 03935 AFN76865.1         ....MKPVKSSCRGFSLIELMIVVAIIGILAAIALPAYQNYTVRSTAASALAEITPAKAAFE.QAISEGHTPSLVSSLPG   75
NTDB id 1191 AAC28468.1 484..945( )           ......MLKKVQKGFTLIELMIVIAIIGILAAIALPAYQDYTARAQATEGFKATAGLQTDLGAWRADRGSFPNAAALVAG   74
NTDB id 1424 RS RS02795 WP 011000517.1        MKSMRHLNKRVQKGFTLIELMIVVAIVGILAAIAIPAYQDYTVRARVTEGLSLAAQAKALVSENAANAQSDLSV....GS   76
NTDB id 1002 ACIAD RS15100 WP 004923779.1     ........MNAQKGFTLIELMIVIAIIGILAAIAIPAYTDYTVRARVSEGLTAASSMKTTVSENILNAGALVAGTPSTAG   72
NTDB id 1075953 ACJEIH RS00975 WP 015444381.1 .........MRMKGFTLIELMIVAAILGILVAIAIPAYQDYTIRARVVEGLEMASAAKLAVVETSLIDNTLPANQADTGY   71
NTDB id 1274 LPP RS09530 WP 011214168.1       .........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGY   71
NTDB id 1258 GC085 RS09745 WP 014844212.1     .........MRQKGFTLIELMIVVAIIGILAAIAIPAYQDYTIRARVTEGLTLADSAKLAVSETAITNNALPATQAATGY   71
consensus                                                **!!*!!!!!!!*!**!*!*!!!*!*!**!**** * ***   * **  * *  ***** **  * ***

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NTDB id 1184 GTF74 RS02335 WP 000649326.1     G........FTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFD...AGVVSSSK.............IQLARDANGLWTC  133
NTDB id 1167 A1552VC RS11065 WP 000649326.1   G........FTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFD...AGVVSSSK.............IQLARDANGLWTC  133
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    G........FTRLGTVEDMGDGKIVIAPTASGALGGTIKYTFD...AGVVSSSK.............IQLARDANGLWTC  133
NTDB id 1197 PSJM300 03940 AFN76866.1         .........YIGITDST...S.YCDVD..LDTAADGHIECTAKGGNAGKFDGKTITLNRT.............ADGEWSC  125
NTDB id 1204 PSJM300 03935 AFN76865.1         .........FIGIGAST...N.YCTVT..LIATATGSITCTTQNGMPDRFNGKTITIART............TDGLWDC  127
NTDB id 1191 AAC28468.1 484..945( )           NVIFDSANGLDGK.YFAPGGV.T.......VTPDTGAINIAFD...AGANAGKTMTITPTVI.......AASGQISTWTC  135
NTDB id 1424 RS RS02795 WP 011000517.1        S.............VFTPTKN.VANLTIAGTGTVPGQI.TVTY...TTAAGGTLALVPTAAGTALPVSSAPSGPIMWTC  138
NTDB id 1002 ACIAD RS15100 WP 004923779.1     S....SCVGVQEISASNATTN.VATATCGA..SSAGQIIVTMD...TTKAKGANITLTPTY..........ASGAVTWKC  132
NTDB id 1075953 ACJEIH RS00975 WP 015444381.1 T............SPAPTIN.VASIAIGN....QGII.TITY...TANAGNGTITMVPTI........DATNRILTWDC  121
NTDB id 1274 LPP RS09530 WP 011214168.1       V............SPAATPN.VASIAIGA....NGVI.TITY...TAAAGGGTIVMTPTL........QAN.GDVTWTC  120
NTDB id 1258 GC085 RS09745 WP 014844212.1     V............SPAATPN.VQSIAIGA....NGVI.TITY...TAAAGGGTIIMTPTL........QAN.GDVTWTC  120
consensus                                               * *     *    **  *     ! !  *      *   * ** * **          *  *   !*!

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NTDB id 1184 GTF74 RS02335 WP 000649326.1     STT................VTSEIAPKGCTAGATIN  153
NTDB id 1167 A1552VC RS11065 WP 000649326.1   STT................VTSEIAPKGCTAGATIN  153
NTDB id 1160 ABDM36 RS02330 WP 000649326.1    STT................VTSEIAPKGCTAGATIN  153
NTDB id 1197 PSJM300 03940 AFN76866.1         AST................LDAKYKPGKCS......  139
NTDB id 1204 PSJM300 03935 AFN76865.1         TSD................LDSRYKPGKCL......  141
NTDB id 1191 AAC28468.1 484..945( )           APGAANG............IETRRLPTSCQ*.....  153
NTDB id 1424 RS RS02795 WP 011000517.1        YAQNKAQAASSVAPSGTMSLAAKYAPAECR......  168
NTDB id 1002 ACIAD RS15100 WP 004923779.1     TTT................SDKKYVPSECRG.....  147
NTDB id 1075953 ACJEIH RS00975 WP 015444381.1 TGG...............TLISKYRPAYCRP.....  137
NTDB id 1274 LPP RS09530 WP 011214168.1       TGG...............TLLAKYRPASCRP.....  136
NTDB id 1258 GC085 RS09745 WP 014844212.1     TGG...............TLLAKYRPASCRP.....  136
consensus                                     **                * *** !* !
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X non conserved

X similar

X ≥ 50% conserved