```
logo
NTDB id 79303 HRS36 RS11270 WP 173237386.1  .......................MNKINLSELTLDNIGQWPIPVKIGVILGVSILIVALGYWLIIKANFEQYANLQTKEVSLKS   61
NTDB id 1255 GCO85 RS04845 WP 011213355.1   .......................MDTVNLNELTLENVGQWPMPVKIGVILGISILIIGLGYWLIVKPNFEQLEIQEAQELALKT   61
NTDB id 1012 ACIAD RS15185 WP 004923719.1   MKLDQLDKTPKESGQPKKKRMNVEKFFQQFNMLDMSNYGSWPLSVKITCWIFIFFLVCALGYFVVIRPQLDDIQTAQAQEKNLLN   85
NTDB id 1056 ABD1 RS15860 WP 000076101.1   MSQDELQELSLEQLSAKKKKFNLDKFLQQFNTLDMNNYGSWPISVKITCWIFIFFAVLALGYFIVIQPKLQAIDNAQAQESNLLN   85
consensus                                  *  * *     *    **** *** * ** **! **!*! !!**!!! * * *  ****!!!******* *********!**! *

logo
NTDB id 79303 HRS36 RS11270 WP 173237386.1  DFESKQHQASNLQAYRNQLQLMNERFGTMLKQLPAKNEMPGLLEEISKTGIGAGLKFMLFAPQPEVEHDFYVELPIKITVTGSYM  146
NTDB id 1255 GCO85 RS04845 WP 011213355.1   EFEKKQHQAANLQAYRNQLQLMNERFGTMLKQLPAKNEMPGLLEEISKTGIASGLKFELFAPQPEVVHDFYVELPIKITIVGSYM  146
NTDB id 1012 ACIAD RS15185 WP 004923719.1   EFREKQSKLRNLQQYQHQLEEMRASFNQQLTQLPKESEIPGLVEDINMTGVNSGLKFKNIRLENEVKQEFFIEQPISIEATGDYH  170
NTDB id 1056 ABD1 RS15860 WP 000076101.1   EFREKDSKLRNLQQYQVQLQEMQANFNQQLEQLPKETEIPSLVEDINLTGVNSGLKFKNIRLEDEVKQEIFIEQPIAIEATGDYH  170
consensus                                  *! *!**  *!!! ! *!!* !* *!   !*!!!  *!*!*!*!*! *!!***!!!!*   * *!!******! !!*! **! !*

logo
NTDB id 79303 HRS36 RS11270 WP 173237386.1  QLALFLSRVAEMNRIVTLHEFSVEGLPSADKKIVAEDVLVMSITAKIYRYRTQ.............................  199
NTDB id 1255 GCO85 RS04845 WP 011213355.1   QLARFLSRVAEMNRIVTLHEFSVEGVSSGDGKTVSQDELVMAITAKIYRYRAQ.............................  199
NTDB id 1012 ACIAD RS15185 WP 004923719.1   AFGAFVSGIAALPRIVTLHDFVIEAKQS.KEKTD.IPEVDYAVKAKTYRYIGVDEQGNASAVQ....ANTSTTTPAQGGTP  245
NTDB id 1056 ABD1 RS15860 WP 000076101.1   AFGAFVSSIAALPRIVTMHDFTVDVSPAKDNKSD.IPVLNYSIKAKTYRYVGATDTSEQTGKAPAASASTNTTVQPK....  246
consensus                                    **!*!**! **!!!!*!*!********** !*  ** **  * !! !!!*** *    **       * * **
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X non conserved

X similar

X ≥ 50% conserved