```
logo
NTDB id 327135 EG344 RS10670 WP 123909413.1  .MISEEYLYAIALRESSQIGDINFHKLVKTFGSAENAWKKAKREYKKLEGIGQKTVADIGNETHLKFAEKELLFCEKNSI   79
NTDB id 1342 RA0C RS05130 WP 004918085.1     MVNAEEILYSIALRRSPLVGDVIFASLVSEVGSAKEAWGLSKLELRKISGVGKQIASNIGDKNLLKFAEKEIEFCIKNEI   80
consensus                                    **  !! !! !!!! !* *!!* !* !!   !!!  !!   ! ! *!* !*!      !!   *!!!!!!!* !! !! !

logo
NTDB id 327135 EG344 RS10670 WP 123909413.1  QVRLRHLDEAPSLLNECIDAPAILYQKGNIDESLQKISLVGTRNMTSYGKQFIIDFFEATQSSKYVSVSGLALGVDKEVH  159
NTDB id 1342 RA0C RS05130 WP 004918085.1     KIHLRHLNQLPRLLGECYDAPAILFQKGDYEDTANNISIVGTRNATAYGKSFLEDLIVQFKNKNIQVISGLALGTDGIAH  160
consensus                                     **!!!!   ! !! !! !!!!!!*!!!  *** * !!*!!!!! ! !!! !* !            *!!!!!! !   !

logo
NTDB id 327135 EG344 RS10670 WP 123909413.1  EQSICSQKPTIAVLAHGFEFLYPAKNRKLSEKILQEGGVLLTEFCSARKPDRENFIQRNRIVAGLSPATIVVETGFGGGS  239
NTDB id 1342 RA0C RS05130 WP 004918085.1     QEALKNNIKTSAVLAHGLHIIYPSKHKILAEEILNNGGALFSEFCSDEKPDREHFLQRNRIVAGLSANTIVVESAYAGGS  240
consensus                                       ** * *! !!!!!! * *!! !** ! ! !!* !! ! *!!!!  !!!!!*!*!!!!!!!!!!* !!!!!****!!!

logo
NTDB id 327135 EG344 RS10670 WP 123909413.1  VSTATFANDYNREVFALPGKITDVHSQGCNQLILHNKATAISTLKDLISMLGLHNPKEKIAELFPYSETTIQLTDNQKNV  319
NTDB id 1342 RA0C RS05130 WP 004918085.1     ISTASFANQYNREVYALAGRLTDKYSQGCNHLIFQNKAKIISSIKGLLKDLDLDNTS.KVLELFPSVE..VSLTPVQEPV  317
consensus                                    *!!!*!!! !!!!!*!!*!**!! *!!!!!*!! *!!!  !!**! !*  ! !*!* *!* !!!!  !*** !!* ! *!

logo
NTDB id 327135 EG344 RS10670 WP 123909413.1  YQFIKQNPQISLDDLAQEIDLSSHKILPIILELELLGKVKSFSGRQFIAN   369
NTDB id 1342 RA0C RS05130 WP 004918085.1     YDIIKEVNDISLDDLSEKLNQPISKVFPVLLELELLGLIRANSGRRYSII   367
consensus                                    !  !!  * !!!!!!   *  * *!* !**!!!!!!! **  !!! *
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X non conserved

X similar

X $\geq$ 50% conserved