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logo

NTDB id 304981 DV114 RS03445 WP 114926157.1  .MYQTQKGFTLIELLIVITIAAVMAVIALPNMNQWIASRRAASQAEQIANLLRFARNEAVRLNLPVYICPAKIKSDGSHN   79
NTDB id 1136 NGFG RS02420 WP 003687912.1     MCTRKQQGFTLTELLIVMAIAAIMATIALPNMSGWIASRRIASHAEQVANLLRFSRGEAVRLNLPVYICPVQVKKDGAPN   80
consensus                                    **    ! !!!! !!!!!* !!!*!! !!!!!!  !!!!!! !!*!!!*!!!!!! ! !!!!!!!!!!!!!  *! !!   !

logo

NTDB id 304981 DV114 RS03445 WP 114926157.1  GCIHKKNSTSISTSIGMLAYADKNENSSYQNDEDDLSIRYIILNG.DTEKVEYRFDYIPSGTSLSALSSSDPKEVWWKFL   158
NTDB id 1136 NGFG RS02420 WP 003687912.1     NRC.....DFSKKGRGMLAFGDKNGNKAYDNDTVDVFLRSVVLNDTDDSRINYAFNHIAFGSSQ.....PTADRVVWTFN   150
consensus                                     *******  *    !!!!**!!! !  ! !!  !* *! **!! *!  ** ! ! *!* !*! ****** *  ! ! !

logo

NTDB id 304981 DV114 RS03445 WP 114926157.1  PNGTFEHS.....TDGKNYQFSDGHIKISLTDESATDAETKKARATVLLINGNGHVEICAKNDDREICEYTSE   226
NTDB id 1136 NGFG RS02420 WP 003687912.1     QNGTFGYLPDQNLKDNSKFVYSDGYIQIVLTDARAVSDADKKFRSAVVLINSSGRVEVCRKNDTRAVCKH...   220
consensus                                    *!!!! * ***** !   * *!!!*! ! !!!  !     !! !  !*!!!  !*!!*! !!! ! *! ****
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X non conserved

X similar

X ≥ 50% conserved