```
logo

NTDB id 26061 HAC RS00225 WP 011577035.1  MKNDAYEIILSWFITPLTAILGRFAEFFLYTLHAQFVFNSAVALAFMFFAYKSLKEQNFFSVSTILEALLFVGFFVLFNYALKNP   85
NTDB id 1220 C694 RS00205 WP 000786660.1  MKNDAYEIILSWFITPLTAILGRFAEFFLYTLHAQLVFNSVVALAFMLFAYRSLKEQNFFSASALTEALLFVGFFALFNYALKNP   85
consensus                                 !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!! !!!!!! !!!*!!!!!!!!! ! * !!!!!!!!! !!!!!!!!!

logo

NTDB id 26061 HAC RS00225 WP 011577035.1  LHFYEFFQNAIFILPNMVTHALSQSLNHFSSHTLSLDFIFNHGFYALSFISDLSHSEISVWLFLSVLQAFFLSVLFAIIILVYLE   170
NTDB id 1220 C694 RS00205 WP 000786660.1  MHFYEFFQNAIFIAPNMIAQSLSQSLSNFSDHALSLDFIFNHGFYALSFISDLSHNEMSVWLFLSVLQGLFLSVLFAIIILVYLE   170
consensus                                 *!!!!!!!!!!!! !!!* * !!!!! *!! ! !!!!!!!!!!!!!!!!!!!!!!! !*!!!!!!!!!!!* !!!!!!!!!!!!!!!

logo

NTDB id 26061 HAC RS00225 WP 011577035.1  VHVWCSLGALFLAFGFFKTWRSVVFACLKKCLALGFYKPFLLLVGFLNVSVTKALMDARMQEQQDLSLLLVVALFLCCVFIIGVP   255
NTDB id 1220 C694 RS00205 WP 000786660.1  VHVWCSLGVLFLAFGFFKTWRSVVVICLKKCFALGFYKPFLLLVGFLNVSVTKALIDAHMQEKQDLSLLLVVALFLCCVFIIGVP   255
consensus                                 !!!!!!!! !!!!!!!!!!!!!!!!  !!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!*!!*!!! !!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 26061 HAC RS00225 WP 011577035.1  FFINALFRVQNSLKETYKLATNLGTNMSQNALTSLQYVTTPPVTTALSSVSTSTSESVSKEKETHSPTFKVETTRLDVKIPNFKQ   340
NTDB id 1220 C694 RS00205 WP 000786660.1  FFINALFRVQNSLKETYKLATNLSANLSQNALNSLQYITTPPASS...SVSSSMSESVSKEKETHSPTFKVETTQLDVKIPNFKQ   337
consensus                                 !!!!!!!!!!!!!!!!!!!!!!!!  !*!!!!! !!!!*!!!! *****!!!!*! !!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!

logo

NTDB id 26061 HAC RS00225 WP 011577035.1  KKVKKDTINPKHEI   354
NTDB id 1220 C694 RS00205 WP 000786660.1  KKVKKDTINTKNEI   351
consensus                                 !!!!!!!!!!*!*!!
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X non conserved
X similar
X ≥ 50% conserved