```
logo                                        MRKQNTLTGIPTSDGQRGSALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGELQVLDLEYTTDSKVT

NTDB id 24190 NMCC RS04460 WP 012221524.1   MRKQNTLTGIPTSDGQRGFALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGELQVLDLEYDTDSKVT   85
NTDB id 1139 NGFG RS02435 WP 003687918.1    MRKQNTLTGIPTSDGQRGSALFIVLMVMIVVAFLVVTAAQSYNTEQRISANESDRKLALSLAEAALREGEFQVLDLEYTADSKVT   85
consensus                                   !!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!  !!!!!

logo                                        FSENCGKGLCTAVNVRTNNNGNEEVFGNIVVQGTPTVEAVKRSCTAKSGKNSTNLCIDNQGVEYKKGTQSVSKMPRYIIEYLGVK

NTDB id 24190 NMCC RS04460 WP 012221524.1   FSENCGKGLCAAVNVRTNN.DNEEAFDNIVVQGKPTVEAVKRSCTA....NSTNLCIDNKGTEYKKGTQSVSKMPRYIIEYLGVK   165
NTDB id 1139 NGFG RS02435 WP 003687918.1    FSENCEKGLCTAVNVRTNNNGNEEVFGNIVVQGTPTVEAVKRSCPAKSGKNSTGLCIDNQGVEYEKGTGNVSKMPRYIIEYLGEK   170
consensus                                   !!!!! !!!! !!!!!!!!* !!! ! !!!!!! !!!!!!!!!!*!****!!! !!!!! ! !! !!!  !!!!!!!!!!!!! !

logo                                        NNQNYYRVTAKAWGKNANTVVVLQSYVSNNDEQ

NTDB id 24190 NMCC RS04460 WP 012221524.1   NGENVYRVTAKAWGKNANTVVVLQSYVSNNDE.   197
NTDB id 1139 NGFG RS02435 WP 003687918.1    NNQNIYRVTAKAWGKNANTVVVLQSYVGNNDEQ   203
consensus                                   !  !*!!!!!!!!!!!!!!!!!!!!!!!! !!!!*
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X non conserved
X similar
X ≥ 50% conserved