```
logo

NTDB id 112143 A6L32 RS10635 WP 002248272.1  MCTRKQQGFTLTELLIVMVIAAIMAMIALPNMSQWIASRRIASHAERIANLLRFSRGEAVRLNLPVYICPVQVKKDGTVN   80
NTDB id 1136 NGFG RS02420 WP 003687912.1     MCTRKQQGFTLTELLIVMAIAAIMATIALPNMSGWIASRRIASHAEQVANLLRFSRGEAVRLNLPVYICPVQVKKDGAPN   80
consensus                                    !!!!!!!!!!!!!!!!!! !!!!!! !!!!!!! !!!!!!!!!!!! *!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! *!

logo

NTDB id 112143 A6L32 RS10635 WP 002248272.1  NKCDSGKKGQGMLAFGDKNGNKGYDNDTADVLLRSVVLNDDINDKRINYAFNHIAFGQTQPTTDRVVWTFNQNGTFGYTK  160
NTDB id 1136 NGFG RS02420 WP 003687912.1     NRCDFSKKGRGMLAFGDKNGNKAYDNDTVDVFLRSVVLND.TDDSRINYAFNHIAFGSSQPTADRVVWTFNQNGTFGYLP  159
consensus                                    !*!!  !!! !!!!!!!!!!!!*!!!!! !! !!!!!!!!* ! !!!!!!!!!!!! *!!! !!!!!!!!!!!!!!! *

logo

NTDB id 112143 A6L32 RS10635 WP 002248272.1  DQHLTKQSSFFYSDGYIQIVLTDARAVSDADKKFRSAVVLIDSSGRVEVCPRGDRRTRRTMCQYK  225
NTDB id 1136 NGFG RS02420 WP 003687912.1     DQNLKDNSKFVYSDGYIQIVLTDARAVSDADKKFRSAVVLINSSGRVEVCRKNDTRAV...CKH.  220
consensus                                    !!*!  *! ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!** ! !  ***! **
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X non conserved

X similar

X $\geq$ 50% conserved