```
logo
NTDB id 47860 NMAA RS04305 WP 002246118.1  MCTRKQQGFTLTELLIVMAIAAIMAMIALPNMSQWIASRRIASHAERVANLLRFSRGEAVRLNLPVYICPVQVKKDGAPNNKCDS   85
NTDB id 1136 NGFG RS02420 WP 003687912.1   MCTRKQQGFTLTELLIVMAIAAIMATIALPNMSGWIASRRIASHAEQVANLLRFSRGEAVRLNLPVYICPVQVKKDGAPNNRCDF   85
consensus                                  !!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!! !!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!

logo
NTDB id 47860 NMAA RS04305 WP 002246118.1  GKKGQGMLAFGDKNGNKIYDNDTADVLLRSVVLNDDINNKRINYAFNHIAFGQTQPTADRVVWTFNQNGTFGYSTNQDLTNTSKF  170
NTDB id 1136 NGFG RS02420 WP 003687912.1   SKKGRGMLAFGDKNGNKAYDNDTVDVFLRSVVLND.TDDSRINYAFNHIAFGSSQPTADRVVWTFNQNGTFGYLPDQNLKDNSKF  169
consensus                                   !!! !!!!!!!!!!!! !!!!! !! !!!!!!!!*    !!!!!!!!!!!! *!!!!!!!!!!!!!!!!!!! * ! !   !!!

logo
NTDB id 47860 NMAA RS04305 WP 002246118.1  VYSDGYIQIVLTDARAVSDADKKFRSAVVLIDSSGRVEVCPRGDRRTRRTMCQYK   225
NTDB id 1136 NGFG RS02420 WP 003687912.1   VYSDGYIQIVLTDARAVSDADKKFRSAVVLINSSGRVEVCRKNDTRAV...CKH.   220
consensus                                  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!** ! !  ***! **
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X non conserved

X similar

X ≥50% conserved