```
logo
NTDB id 1291 VP RS12200 WP 005479697.1        .MDSIAKRPRTRLSPLKRKQQLMEIALEVFARRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDEVLNHVVRQFSNFL   79
NTDB id 321223 EAE30 RS16160 WP 123016836.1   MDASIERRPRTRLSPQKRKHQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLTFVVRQYSNFL   80
NTDB id 1161 ABDM36 RS11380 WP 000340098.1    MDASIEKRPRTRLSPQKRKLQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL   80
NTDB id 1147 A1552VC RS01790 WP 000340098.1   MDASIEKRPRTRLSPQKRKLQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL   80
consensus                                     ***!!**!!!!!!!!!*!!!*!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!**!!!!*!!!!

logo
NTDB id 1291 VP RS12200 WP 005479697.1        SDNIDLDIHARENIANITNAMIELVSQDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLVQNMFIKAIERGEVCDQHDS   159
NTDB id 321223 EAE30 RS16160 WP 123016836.1   TDNIDLDLHARQNLRNIANGIVDLANDDCHWLKVWFEWSASTREEVWPLFVSSNHTNQLLLKNMFTKGIERGELDNRYDA   160
NTDB id 1161 ABDM36 RS11380 WP 000340098.1    TDHIDLDLDVKTNLQTLCKEMVKLAMTDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLIRNMFMKAMERGELCEKHDV   160
NTDB id 1147 A1552VC RS01790 WP 000340098.1   TDHIDLDLDVKTNLQTLCKEMVKLAMTDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLIRNMFMKAMERGELCEKHDV   160
consensus                                     *!*!!!!** **!** ** ****!***!!!!!!!!!!!!!!!!!*!!!!!!!!*!*!!!!!**!!!*!**!!!!*****!*

logo
NTDB id 1291 VP RS12200 WP 005479697.1        EHLANLFHGICYSLFVQANRFKGEAELKELVSAYLDMLCIYNREH   204
NTDB id 321223 EAE30 RS16160 WP 123016836.1   EDMATLFHGLCYSLFVQTNRVPNENSVHQLVDNYLDMLCIYAKAS   205
NTDB id 1161 ABDM36 RS11380 WP 000340098.1    DNMASLFHGIFYSIFLQVNRLGEQEAVYKLADSYLNMLCIYKN..   203
NTDB id 1147 A1552VC RS01790 WP 000340098.1   DNMASLFHGIFYSIFLQVNRLGEQEAVYKLADSYLNMLCIYKN..   203
consensus                                     ***!*!!!!**!!*!*!*!!*** *****! **!! !!!!!**
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X non conserved

X similar

X $\geq$ 50% conserved