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logo

NTDB id 263855 C1S74 RS14005 WP 038870500.1  .MDSIAKRPRTRLSPLKRKQQLMEIALEVFARRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDEVLNHVVRQFSNFL   79
NTDB id 1291 VP RS12200 WP 005479697.1       .MDSIAKRPRTRLSPLKRKQQLMEIALEVFARRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDEVLNHVVRQFSNFL   79
NTDB id 1161 ABDM36 RS11380 WP 000340098.1   MDASIEKRPRTRLSPQKRKLQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL   80
NTDB id 1147 A1552VC RS01790 WP 000340098.1  MDASIEKRPRTRLSPQKRKLQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL   80
consensus                                    *  !! !!!!!!!!! !!! !!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!*!!!!*!!!!

logo

NTDB id 263855 C1S74 RS14005 WP 038870500.1  SDNIDLDLHARENIANITNAMIELVSQDCHWLKVWFEWSASTRDEVWPLFVTTNRTNQLLVQNMFIKAIERGEVCDQHEP   159
NTDB id 1291 VP RS12200 WP 005479697.1       SDNIDLDIHARENIANITNAMIELVSQDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLVQNMFIKAIERGEVCDQHDS   159
NTDB id 1161 ABDM36 RS11380 WP 000340098.1   TDHIDLDLDVKTNLQTLCKEMVKLAMTDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLIRNMFMKAMERGELCEKHDV   160
NTDB id 1147 A1552VC RS01790 WP 000340098.1  TDHIDLDLDVKTNLQTLCKEMVKLAMTDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLIRNMFMKAMERGELCEKHDV   160
consensus                                    *!*!!!!** * !*  **  !* !   !!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!* !!!*!!*!!!!*!* !**

logo

NTDB id 263855 C1S74 RS14005 WP 038870500.1  EHLANLFHGICYSIFVQANRSKSEAELTNLVSAYLDMLCIYNREHH   205
NTDB id 1291 VP RS12200 WP 005479697.1       EHLANLFHGICYSLFVQANRFKGEAELKELVSAYLDMLCIYNREH.   204
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 ≥ 50% conserved