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logo

NTDB id 1161 ABDM36 RS11380 WP 000340098.1      MDASIEKRPRTRLSPQKRKLQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL   80
NTDB id 1147 A1552VC RS01790 WP 000340098.1     MDASIEKRPRTRLSPQKRKLQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLNFVVRQYSNFL   80
NTDB id 1009815 ABFV67 RS02330 WP 110167828.1  MDASIEKRPRTRLSPQKRKQQLMEIALEVFAKRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDDVLTFVVREFSNFL   80
NTDB id 1291 VP RS12200 WP 005479697.1          .MDSIAKRPRTRLSPLKRKQQLMEIALEVFARRGIGRGGHADIAEIAQVSVATVFNYFPTREDLVDEVLNHVVRQFSNFL   79
consensus                                       ***!!*!!!!!!!!!*!!! !!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!**!!!**!!!!

logo

NTDB id 1161 ABDM36 RS11380 WP 000340098.1      TDHIDLDLDVKTNLQTLCKEMVKLAMTDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLIRNMFMKAMERGELCEKHDV  160
NTDB id 1147 A1552VC RS01790 WP 000340098.1     TDHIDLDLDVKTNLQTLCKEMVKLAMTDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLIRNMFMKAMERGELCEKHDV  160
NTDB id 1009815 ABFV67 RS02330 WP 110167828.1  TDHIDLDLHAKQNMHNIASGMVSLAIDDCHWLKVWFEWSASTRDEVWPLFMSSNRTNQLLVKNVFSKAIERSEVCSKHNP  160
NTDB id 1291 VP RS12200 WP 005479697.1          SDNIDLDIHARENIANITNAMIELVSQDCHWLKVWFEWSASTRDEVWPLFVSTNRTNQLLVQNMFIKAIERGEVCDQHDS  159
consensus                                       *!*!!!!** **!** ****!**!***!!!!!!!!!!!!!!!!!!!!!!!!!!*!*!!!!!!!**!*!*!!*!!*!*!**!**

logo

NTDB id 1161 ABDM36 RS11380 WP 000340098.1      DNMASLFHGIFYSIFLQVNRLGEQEAVYKLADSYLNMLCIYKN......  203
NTDB id 1147 A1552VC RS01790 WP 000340098.1     DNMASLFHGIFYSIFLQVNRLGEQEAVYKLADSYLNMLCIYKN......  203
NTDB id 1009815 ABFV67 RS02330 WP 110167828.1  EDLATLFHGICYSLFVQANRARDEADMQKLVNNYLNMLCIYQGDRAVAI  209
NTDB id 1291 VP RS12200 WP 005479697.1          EHLANLFHGICYSLFVQANRFKGEAELKELVSAYLDMLCIYNREH....  204
consensus                                       ***!*!!!!!*!!*!*! !!***  ****! **!!*!!!!!***
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X non conserved

X similar

X ≥ 50% conserved