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logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  MRNSRLLLPLAAASATAGITAAAYFSAVFLFFLFLLIILIKTRHAVLILVCFFSFILFFSLYAVADSKNVSVYQRGTYQA   80
NTDB id 107 BSU 25570 NP 390435.1              MRNSRLLLPMAAASATAGITAAAYFPAIFLFILFLLIILIKTRHAFLIIVCFFSFILFFVLYAVTDSQNVSSYRQGTYQF   80
consensus                                      !!!!!!!!!*!!!!!!!!!!!!!!!*!*!!! !!!!!!!!!!!!! !!*!!!!!!!!!! !!!! !! !!! ! !!!!

logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  KAVIDSIPKIDGDRMSMVVKTPDGEKWAASYRIQSAEEKDQLSYLKPGMACEWTGALEEPKHATVPGAFDYNEYLYRQHI  160
NTDB id 107 BSU 25570 NP 390435.1              KAVIDTIPKIDGDRMSMMVETPDKEKWAAAYRIQSAGEKEQLLYIEPGMSCELTGTLEEPNHATVPGAFDYNEYLYRQHI  160
consensus                                      !!!!!*!!!!!!!!!!!*! !!! !!!!! !!!!!! !!*!! !* !!! !! !! !!!! !!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  HWNYSVTSIQKCSEPVNVSYKLLSLRKDIVSFTNSLLPPDSAGIVQALIVGDRFYVEAEVLNAYQKLGVIHLLAISGLHV  240
NTDB id 107 BSU 25570 NP 390435.1              HWNYSVTSIQNCSEPENFKYKVLSLRKHIISFTNSLLPPDSTGIVQALTVGDRFYVEDEVLTAYQKLGVVHLLAISGLHV  240
consensus                                      !!!!!!!!!!! !!!! !  !!*!!!!!*!*!!!!!!!!!!! !!!!!! !!!!!!!! !!! !!!!!!!*!!!!!!!!!!

logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  GILTAGLFYIMIRIGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGIYLAGSLVKWRVHAAGAICTSYIVLLLFNP  320
NTDB id 107 BSU 25570 NP 390435.1              GILTAGLFYIMIRLGITREKASILLLLFLPLYVMLTGAAPSVLRAALMSGVYLAGSLVKWRVRSATAICLSYIVLLLFNP  320
consensus                                      !!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!
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logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  VKQLDALILTHADQDHIGEAETLLKHHKVKRLVIPKGFVSEPKDEKVLQAAREEGVPIEEVKRGDVLQIKDLQFYVLSPE   640
NTDB id 107 BSU 25570 NP 390435.1              IKQLDALILTHADQDHIGEAEILLKHHKVKRLVIPKGFVSEPKDEKVLQAAREEGVAIEEVKRGDVLQIKDLQFHVLSPE   640
consensus                                      *!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!*!!!!!

logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  GPDPASKNNSSLVLWMKTGGMSWILTGDLEKEGEQEVISVFPNLKTDVLKVGHHGSKGSTGEDFIKQIQPETAIISAGEN   720
NTDB id 107 BSU 25570 NP 390435.1              APDPASKNNSSLVLWMETGGMSWILTGDLEKEGEQEVMNVFPNIKADVLKVGHHGSKGSTGEEFIQQLQPKTAIISAGKN   720
consensus                                      *!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!* !!!!*! !!!!!!!!!!!!!!!!!!*!! !*!! !!!!!!! !

logo

NTDB id 1022041 ABZM97 RS13105 WP 367386875.1  NRYHHPHQEVLQILQRHPIRVLRTDQSGTIQYRYKNRVGTFSVYPPYDTSE.....   771
NTDB id 107 BSU 25570 NP 390435.1              NRYHHPHQKVLQLLQRHSIRVLRTDQNGTIQYRYKNRVGTFSVYPPYDTSDITETN   776
consensus                                      !!!!!!!! !!!*!!!!!*!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!******
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X non conserved

X similar

X $\geq$ 50% conserved