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logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  MIENWPQKPEGSQWTDDQWKAVVANGRDILVAAAAGSGKTAVLVERIIKKIISEEENPVDVDRLLVVTFTNAAAQEMKNRI   80
NTDB id 119 BSU 10630 NP 388944.2            ...MNIPKPADSTWTDDQWNAIVSTGQDILVAAAAGSGKTAVLVERMIRKITAEENPIDVDRLLVVTFTNASAAEMKHRI   77
consensus                                    ***  **!!  ! !!!!!! !*!  ! !!!!!!!!!!!!!!!!!!!*!*!!  !!!!*!!!!!!!!!!!!! ! !!!*!!

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  GEALEKVLIDGPGSQHIRKQLSLLNKASISTIHSFCLQVIRGYYYMLDVDPRFRIGNQTENELLKEEVLDDILEEEYGIE  160
NTDB id 119 BSU 10630 NP 388944.2            AEALEKELVQRPGSLHIRRQLSLLNRASISTLHSFCLQVLKKYYYLIDLDPGFRIADQTEGELIGDEVLDELFEDEYAKG  157
consensus                                    *!!!!! !*  !!! !!!*!!!!!!*!!!!!*!!!!!!!** !!!**!*!! !!!* !!! !!* *!!!!** !*!!*

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  DNSIFFELVDRYTSDRSDDDLQRMILALHTESRAHPNPEKWLDKLVEAYDVEGK.TIEDLVYASYLLEDVKFQLETAEEH  239
NTDB id 119 BSU 10630 NP 388944.2            .EKAFFELVDRYTTDRHDLDLQFLVKQVYEYSRSHPNPEAWLESFVHLYDVSEKSAIEELPFYQYWKEDIAMVLNGAKEK  236
consensus                                    *   !!!!!!!!!*!!*! !!! **  **  !! !!!!! !!*  !* !!!  !* !!*!**  !* !!*   !  ! !*

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  IRKATELAMLPDGPAPRVETLQADVALLGMLSSAARGSWTNVYEAMQNVSWQTLKRIKKSDYNEDVVKQVDSLRNKAKDE  319
NTDB id 119 BSU 10630 NP 388944.2            LLRALELTKAPGGPAPRADNFLDDLAQIDELIQ.HQDDFSELYKRVPAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKL  315
consensus                                    * *! !!   ! !!!!! *    !*! *  !  **   ** *!  ** !!*   ! *!  ** * **     !!! !!

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  VKKLQEELFSRRPESFLRDFQDMHPVLEKLVKLVKVFTERFQAIKRDKGMVDFTDLEHFCLQILSEQSEDGEMKPSAVAL  399
NTDB id 119 BSU 10630 NP 388944.2            LEKLKTDYFTRSPEQHLKSLAEMKPVIETLVQLVISYGKRFEAAKQEKSIIDFSDLEHYCLAILTAENDKGEREPSEAAR  395
consensus                                    * !!  * !*! !! *!*   *!*!!*! !! !!  *  !! ! ! *! **!!*!!!!*!! !!*   * !!  !!  !

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  QYRNKFAEVLVDEYQDTNFVQESIIKFVTKDSESEGNLFMVGDVKQSIYRFRLAEPGLFLGKYKRFTQEGLDGGMKIDLA  479
NTDB id 119 BSU 10630 NP 388944.2            FYQEQFHEVLVDEYQDTNLVQESILQLVTSGPEETGNLFMVGDVKQSIYRFRLAEPLLFLSKYKRFTESGEGTGRKIDLN  475
consensus                                     !   !*!!!!!!!!!!! !!!!!*  !!  *!  !!!!!!!!!!!!!!!!!!!!! !!! !!!!!!  !   ! !!!!

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  KNFRSRHEVLDSTNFLFKQLMGGKYGEVDYDEQAELKLGASYPENEEVETELLLIQQTEEEVQDSEEGTEVETVQLEARL  559
NTDB id 119 BSU 10630 NP 388944.2            KNFRSRADILDSTNFLFKQLMGGKIGEVDYDEQAELKLGAAYPDNDETETELLLIDNAED.TDASEEAEELETVQFEAKA  554
consensus                                    !!!!!!***!  !!!*!!!*!!  *!!*!!!  !!!!!!! !!* **   !!!*! * !**    !!* !*!  ! !!*
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logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  MAQRIKAMVDSGYEVYDRKTDSMRPVQYRDFVILLRSMPWAPQIMEELKLQGIPVYADLATGYFEATEVNIMMNVFRVID   639
NTDB id 119 BSU 10630 NP 388944.2            IAKEIRKLISSPFKVYDGKKKTHRNIQYRDIVILLRSMPWAPQIMEELRAQGIPVYANLTSGYFEAVEVAVALSVLKVID   634
consensus                                    *!  !* ** !** !!! !  **!**!!!! !!!!!!!!!!!!!!!!* !!!!!!! ! *!!!!! !! * * ! *!!!

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  NPMQDIPLAAVLRSPIVGLNDEELATLRAHGKKGSFYEVMSSFLKGAPLEEEQELHDKLEWFYNLLQGWREFARQQSLSD   719
NTDB id 119 BSU 10630 NP 388944.2            NPYQDIPLASVLRSPIVGADENELSLIRLENKKAPYYEAMKDYLAAGD..RSDELYQKLNTFYGHLQKWRAFSKNHSVSE   712
consensus                                    !! !!!!!! !!!!!!!!  * !!  *! * !!***!! !  *! *****   !!* !!  !! *!! !! ! ***!*!*

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  LIWKVYGETGYYDFVGGLPAGKQRQANLRVLYDRARQYEATSFRGLFRFLRFIERILERGDDMGTARALGEQEDVVRIMT   799
NTDB id 119 BSU 10630 NP 388944.2            LIWEVYRDTKYMDYVGGMPGGKQRQANLRVLYDRARQYESTAFRGLFRFLRFIERMQERGDDLGTARALSEQEDVVRLMT   792
consensus                                    !!! !! *! ! !*!!!*!*!!!!!!!!!!!!!!!!!! ! !!!!!!!!!!!!!* !!!!!*!!!!!! !!!!!!!*!!

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  IHKSKGLEFPVVFVAGLGRRFNTQDLMKRFLLHKDFGFGSQFIDPRKRIKYTTLSQLAIKRKMKMELIAEEMRVLYVALT   879
NTDB id 119 BSU 10630 NP 388944.2            IHSSKGLEFPVVFVAGLGRNFNMMDLNKSYLLDKELGFGTKYIHPQLRISYPTLPLIAMKKKMRRELLSEELRVLYVALT   872
consensus                                    !! !!!!!!!!!!!!!!! !!  !! ! *!!*!* !!!* *!*!  !! !*!!* *!*!*!!* !!* !!*!!!!!!!

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  RAKEKLILIGTVKDANKEMEKWLDAREHSEWLLPDHIRAGASCYLDWIAPSLYRHRDSEMLLELGQGNIPGEIYGYSASW   959
NTDB id 119 BSU 10630 NP 388944.2            RAKEKLFLIGSCKDHQKQLAKWQASASQTDWLLPEFDRYQARTYLDFIGPALARHRDLGDLAGVP...AHADISGHPARF   949
consensus                                    !!!!!! !!!**!!**! * !!     ***!!!!** !  ! *!!!*!*! ! !!!!   !  *****  **! !**! *

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  KVEVVDGKTLLAPE..PVQEEKQELLEALREKKAVPLESERKEEVYDRLMWEYGYEDATSHRAKQSVTEIKRNYQSEEG.  1036
NTDB id 119 BSU 10630 NP 388944.2            AVQMIHSYDLLDDDLEERMEEKSERLEAIRRGEPVPGSFAFDEKAREQLSWTYPHQEVTQIRTKQSVSEIKRKREYEDEY  1029
consensus                                     ! ***   !! *****  !!! ! !!!*!   *!!      !   * ! ! !** * ! *! !!!!*!!!!    !* *

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  .SDNAFIKKLRAPIKTRPRFMEKKGLTYAERGTAVHAVMQHVDLKKPITIEVLQEQIARMVNKELLTFEQAEEIAIEKVI  1115
NTDB id 119 BSU 10630 NP 388944.2            SGRAPVKPADGSILYRRPAFMMKKGLTAAEKGTAMHTVMQHIPLSHVPSIEEAEQTVHRLYEKELLTEEQKDAIDIEEIV  1109
consensus                                    *   *  *    **  !! !! !!!!! !!*!!!*! !!!!**! ****!!     **!*  !!!!! !! * ! !! **
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logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  SFFDSDLGRRVLAAKSVEREVPFTMMLSAEEAYQDWQGKSDETILVQGVIDCMIEEEDGITLIDFKTDTIEGKFPGGFDQ  1195
NTDB id 119 BSU 10630 NP 388944.2            QFFHTEIGGQLIGAKWKDREIPFSLALPAKEIYPDAH.EADEPLLVQGIIDCLYETEDGLYLLDYKSDRIEGKFQHGFEG  1188
consensus                                     !!****!  ***!!  *!!*!!** !*! ! !*! **  !!**!!!!*!!!* ! !!!* !*!*!*! !!!!!**!!*

logo
NTDB id 345258 EXW58 RS05570 WP 215580884.1  AKPTLEDRYKVQLSLYAKALEKSLKHPVKEKCLYFFDGNHVVNIDE  1241
NTDB id 119 BSU 10630 NP 388944.2            AAPILKKRYETQIQLYTKAVEQIAKTKVKGCALYFFDGGHILTL..  1232
consensus                                    ! ! !  !!  !* !! !!*!   !**!! **!!!!!! !** ***
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X non conserved

X similar

X $\geq$ 50% conserved