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logo

NTDB id 1035712 ABVF47 RS08300 WP 371687900.1  MFGFFKRKKNTEPEQ.EQASVNPAQPESVEQPDTAAETHTQPEPAASHQSVEKHTANAVVSESTATETTEDII..VPAAV   77
NTDB id 1118 NGFG RS11455 WP 003696286.1       MFSFFRRKKKQETPALEEAQVQETAA.KVESEVAQIVGNIKEDVESLAESVKGRAESAVETVSGAVEQVKETVAEMPSEA   79
consensus                                      !! !!*!!!  !** *! ! !**  ** !! *     *  ***  * !!  *  !! * ! ! !   * ****!

logo

NTDB id 1035712 ABVF47 RS08300 WP 371687900.1  DEPAAIATSTESEVVQAPASENTIVIPVETTEIPSETDKPARLGWAARLKQGLSKSRNQMAKSLASVFGGGKIDEDLYEE  157
NTDB id 1118 NGFG RS11455 WP 003696286.1       GEAAERVESAK.E.....AVAETVGEAVGQVQEAVATTEEHKLGWAARLKQGLAKSRDKMAKSLAGVFGGGQIGEDLYEE  153
consensus                                       !*!     !  *!*****!   !*  *!     *  !  ***!!!!!!!!!!! !!!  !!!!!! !!!!! ! !!!!!!

logo

NTDB id 1035712 ABVF47 RS08300 WP 371687900.1  LETVLLTSDMGIEATEKLLAQVRQRVTLKGLKNGNELRSALKEALYELLQPLQQPLSIPADKKPFVIMMAGVNGAGKTTS  237
NTDB id 1118 NGFG RS11455 WP 003696286.1       LETVLITGDMGMEATEYLMKDVRGRVSLKGLKDGNELRGALKEALYDLIKPLEKPLVLPETKEPFVIMLAGINGAGKTTS  233
consensus                                      !!!!!*! !!!*!!!! !*  !! !!*!!!!! !!!!! !!!!!!!*!* !!  !! *!  ! !!!!!*!!*!!!!!!!!

logo

NTDB id 1035712 ABVF47 RS08300 WP 371687900.1  IGKLAKYFQSQGKSVLLAAGDTFRAAAREQLQQWGERNNVTVIAQESGDSAAVCFDALEAAKARGIDIVLADTAGRLPTQ  317
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGKLAKYFQAQGKSVLLAAGDTFRAAAREQLQAWGGRNNVTVISQTTGDSAAVCFDAVQAAKARGIDIVLADTAGRLPTQ  313
consensus                                      !!!!!!!!! !!!!!!!!!!!!!!!!!!!!!! !! !!!!!!! ! *!!!!!!!!!!* !!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1035712 ABVF47 RS08300 WP 371687900.1  LHLMEEIKKVKRVLQKSLPEAPHEIMLVLDANVGQNAINQVVAFDDALGLTGLIVSKLDGTAKGGILAALAAKRPIPVRF  397
NTDB id 1118 NGFG RS11455 WP 003696286.1       LHLMEEIKKVKRVLQKAIPGAPHEIIVVLDANIGQNAVNQVKAFDDALGLTGLIVTKLDGTAKGGILAALASDRPVPVRY  393
consensus                                      !!!!!!!!!!!!!!!! *! !!!!!**!!!!!*!!!!*!!! !!!!!!!!!!!!!*!!!!!!!!!!!!!!!  !!*!!!*

logo

NTDB id 1035712 ABVF47 RS08300 WP 371687900.1  IGVGEGIDDLRPFDAKAFVDALIEDEE   424
NTDB id 1118 NGFG RS11455 WP 003696286.1       IGVGEGIDDLRPFDARAFVDALLD...   417
consensus                                      !!!!!!!!!!!!!!!*!!!!!!*****
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X non conserved

X similar

X ≥ 50% conserved