```
logo
NTDB id 985629 MHB66 RS07695 WP 046310923.1  MYNVSERMIFHRLKGLISPSLLTKWWKVDPELYINEETHHFKQDRELQTIDFTRLKKAEENEFPIFQHIVQAYLKQNIHM   80
NTDB id 114 BSU 16110 NP 389493.1            MDQAAVCLTICRINQLLSPSLLLKWWKADPSMSLTSPVLQTV...TRDQIKAAALKNEIEQFYPKLPRVLAAYREQGINT   77
consensus                                    ! *   **   !*  !*!!!!! !!!! !! * *  * **  ***    !    !!   !* *!  **** !!  ! !*

logo
NTDB id 985629 MHB66 RS07695 WP 046310923.1  IPITSPLYPSTLKNIYDPPPVLFLKGNVSYLNEEKSLGVVGTRVPSSYGEACVKKIVGELVKEDWMIVSGLAKGIDGLAH  160
NTDB id 114 BSU 16110 NP 389493.1            IPISSKQYPFWLKSIYDPPAVLFAKGDMTLLSKGRKIGIVGTRNPTAYGKQVVNHLTKEICRKGWVIVSGLASGIDGMSH  157
consensus                                    !!!*!* !!  !! !!!!!*!!! !! ** !   * *!*!!!! !* !!  *! **  !***  !*!!!!!! !!!!* !

logo
NTDB id 985629 MHB66 RS07695 WP 046310923.1  KECIRKKGKTIGIIAGGFQHLYPKEHEQMAQYMGEHHLLLSEHPPYVKPEKWHFPMRNRLISALTKGTIVIQCKEKSGSL  240
NTDB id 114 BSU 16110 NP 389493.1            AASIKAKGRTIGVIAGGFQHIYPRENLQLADHMAKHHILLSEHPPETKPQKWHFPMRNRIISGLSEGVIVVQGKEKSGSL  237
consensus                                      *!* !!*!!!*!!!!!!!*!!*!* !*! *!* !!*!!!!!!!  !! !!!!!!!!!*!!*!* ! !!*!*!!!!!!!

logo
NTDB id 985629 MHB66 RS07695 WP 046310923.1  ITAYQALEQGKEVFAVAGSIFDPNSTGPARLIQQGAKLVHSTKDILEEFSFRSVQYTEPT  300
NTDB id 114 BSU 16110 NP 389493.1            ITAYQALEQGREVFAVPGSLFDPYAGGPIKLIQQGAKAIWSAEDIFEELPERNVQYTEPF  297
consensus                                    !!!!!!!!!!*!!!!!*!!*!!!   !! *!!!!!!! **!  !! !! * ! !!!!!!
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X non conserved

X similar

X ≥ 50% conserved