```
logo

NTDB id 60992 RBAU RS12410 WP 007408321.1  MKRIKRTWPLKDQAAFLKRLGEMTEKGYSLIEGLRLLQLQLHKRQLAELTDGIRRLREGDAFYQVLEALSFHKEAVSICYFAEKH   85
NTDB id 97 BSU 24720 NP 390352.1           .......................MTAGGYTLLDGLRLMELQMNKRQAADLTDSVTCLREGAPFYQVLKSLSFHKEAVGICYFAETH   63
consensus                                  **********************!!  !!*!**!!!!* !!**!!! !*!!! * *!!!! *!!!!!  !!!!!!!! !!!!!! !

logo

NTDB id 60992 RBAU RS12410 WP 007408321.1  GELPGAMKQSGDLLQRKLMQTNQLKKMLRYPVFLISSVCVMFYILQSLIIPQFSGIYQSMNMNTSRTTAFIFAFFRHFHEACALA  170
NTDB id 97 BSU 24720 NP 390352.1           GELPASMIQSGELLERKIAQADQLKRVLRYPLFLIFTVAVMFYMLQSIIIPQFSGIYQSMNMETSRSTDMLFAFFQHIDLVIILL  148
consensus                                  !!!!* ! !!!*!! !!* !  !*!**!!!!*!!! *!*!!!!*!!!*!!!!!!!!!!!!!! !!!*!  *!!!! ! *  * !

logo

NTDB id 60992 RBAU RS12410 WP 007408321.1  LSAAFCLFLYVWFLCKKKSPQDKMLIVVKIPLLGKAAVLFNSYFFSLQLSSLLKSGLSIYDSLTAFKEQSFLPFYREEAEMLITR  255
NTDB id 97 BSU 24720 NP 390352.1           VLFTAGIGIYYWLVFKKKSPARQMLICIRIPLVGKLVKLFNSYFFSLQLSSLLKSGLSIYDSLNAFKHQTFLPFYRCEAEQLIER  233
consensus                                  *    ** *! ! **!!!!!   !!!***!!!*!!   !!!!!!!!!!!!!!!!!!!!!!!!! !!!*!*!!!!!!*!!! !! !

logo

NTDB id 60992 RBAU RS12410 WP 007408321.1  LKAGETIESALSGHPCYEKDLAAAVSHGQANGLLHRELYTYSQFMMERLEQNAAKYTSILQPVIYGVVACMILIVYMSMLMPMYQ  340
NTDB id 97 BSU 24720 NP 390352.1           LKAGESIESAICGSLFYETDLSKVISHGQLSGRLDRELFTYSQFILQRLEHKAQKWTGILQPMIYGFVAAMILLVYLSMLVPMYQ  318
consensus                                  !!!!!*!!!!**!***!! !!   *!!!!  ! !*!!!*!!!!!** !!!* ! !*! !!!!*!!! !!*!!!*!!*!!!*!!!!

logo

NTDB id 60992 RBAU RS12410 WP 007408321.1  MMNQM   345
NTDB id 97 BSU 24720 NP 390352.1           MMNQM   323
consensus                                  !!!!!
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X non conserved

X similar

X ≥ 50% conserved