```
logo
NTDB id 595 KW2 RS08725 WP 021037687.1          ..MEKILDKLKEYWKIMLLVVCALIAGGIFYIFTNSPKPAETLSVENLSSSSTKSSVSKFNSSSSEKNKNEIMVDLKGAV   78
NTDB id 1165945 AB3225 RS00610 WP 004270751.1   .MWRDYWQRYRRFWWVPCLVII.LLGGGVFWQHSRQQRFLSTN.QQLMAQSTKR.KVKAKPVKQKKSVSKGGFVDIKGAV   76
NTDB id 613 LCA RS05340 WP 231952379.1          MMWQDYWQRYRRFWWVPCLVLLGIIAGGFYWHQSQQKQKVTQN.EQLLATSASKTKPATKKTHQAATKKKGGFVDIKGAV   79
consensus                                        ** *********!****!!******!!***  *** * ****  *****!******  * * *  * *****!!*!!!!

logo
NTDB id 595 KW2 RS08725 WP 021037687.1          AKPNVYQISS.DERLVDLIRQAGGFTDQADQKSINLSAKLKDEEVIYVPKVGESSSSESTDSP...............IG   142
NTDB id 1165945 AB3225 RS00610 WP 004270751.1   AHPGIYPICVAKTRLFEVLKQAGDVTEAADTSQLNLAQTLSDQLIIYVPTKGEAVQAEELVKAADEP.........QAAT   147
NTDB id 613 LCA RS05340 WP 231952379.1          VYPGIYPITVSETRLFEVLKNAGGPTDAADTKRLNLAQKLTDQLIIYIPKKGEVVTDQELIQAPQQIATTSERTQEQAST   159
consensus                                        * !**!*!** **!!******!!* !**!!** *!!***!*!***!!*!**!! ** **** *             ****

logo
NTDB id 595 KW2 RS08725 WP 021037687.1          SSVSNQISTTSGPKVNINKADLTELQKLTGIGQKKAGDIIDFRMKNGDFKSIEDLGKVSGFGDKTLEKLKDEISID   218
NTDB id 1165945 AB3225 RS00610 WP 004270751.1   DNQAGASPISTDAKINLNQADATQLQTLNGIGPKKAEQIIAYRDEQGQFKQIEDLQKVGGIGPKTFEQLRPQICVN   223
NTDB id 613 LCA RS05340 WP 231952379.1          GTSANASSTSDTAKIDLNQADSTQLQTLNGIGPKKAEQIIAYRDEQGKFKQIEDLQKVGGIGPKTFEQLQSQICVN   235
consensus                                        * ******** *!***!*!! !*!!*!*!!!*!!!**!!**!***! !!*!!!!*!!*!*!*!!*!*!* *!***
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X non conserved

X similar

X $\geq$ 50% conserved