```
logo                                    

NTDB id 373285 FOH38 RS23540 WP 143999073.1  MINRTVLVGRLTKDPELCYTPNGIASCRFMVAVNRTFSNQQGEKEADFISCQAWRKQAENLANFMRKGNLIGVEGRIQTG  80
NTDB id 113 BSU 40900 NP 391970.1            MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQTR  80
NTDB id 624 LCA RS00040 WP 011373726.1       MINRVVLVGRLTRDVDLRYTSSGAAVGTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTR  80
consensus                                    !*!!*!!!!!!!*!**!*!!**!*!***!**!!!!*!*!**!**!!!!!*!* !!**!!!*!!!**!!*!*!!*!!*!!*

logo                                    

NTDB id 373285 FOH38 RS23540 WP 143999073.1  SYEGQNGQRVYTTDVVADSVQFLEPRSNPQQQGGSQSQQSYNNQPSYGSGQPQQQYSCQNAQYNQQHSQFKWPSQQQNQP  160
NTDB id 113 BSU 40900 NP 391970.1            NYENQQGQRVFVTEVQAESVQFLEPKNGGGSGS.....GGYNEGNSGGGQ.....YFCG....GQNDNPFGGNQNN....  142
NTDB id 624 LCA RS00040 WP 011373726.1       NYENQQGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQSNGSQ.................QSGFTSPQQT....  140
consensus                                    *!!*!*!!!!**!*!*******!!***   *  *    * *! **! !**     * *      ** * !  ****

logo                                    

NTDB id 373285 FOH38 RS23540 WP 143999073.1  PMNQPNYTRVDEDPFANSRGPIEVSEDDLPF  191
NTDB id 113 BSU 40900 NP 391970.1            .QRRNQGNSFNDDPFANDGKPIDISDDDLPF  172
NTDB id 624 LCA RS00040 WP 011373726.1       .GNAPAANNTQADPFANNGQAIDISDDDLPF  170
consensus                                      * ****  **!!!!! * *!**!*!!!!!
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X non conserved

X similar

X ≥ 50% conserved