```
logo                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 1147638 CKV88 RS07060 WP 002464335.1  ....MPINKVKAFTMIEMLFVLLMVSIILLTTLTSKSLRNLETMNSEMQLSSFILQLDYLKSKAIITQGPILIQFVKHSN   76
NTDB id 33 MW RS07990 WP 001788899.1          MEKQLQIRKQSAFTMIEMLVVMMLISIFLLLTMTSKGLSNLRVIDDEANIISFITELNYIKSQAIANQGYINVRFYENSD   80
NTDB id 17 SA RS07760 WP 001788899.1          MEKQLQIRKQSAFTMIEMLVVMMLISIFLLLTMTSKGLSNLRVIDDEANIISFITELNYIKSQAIANQGYINVRFYENSD   80
consensus                                     ******!*!**!!!!!!!!!*!****!!*!!*!*!!!*!*!!*****!****!!!**!*!*!!*!!**!!*!***!***!*

logo                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                
NTDB id 1147638 CKV88 RS07060 WP 002464335.1  QVKVTENNKTSYFLSINNAEIISINKVNYLYFDKRGNIKPFGSITIKVNKAYYKVIFHIEKGRIRYEKL   145
NTDB id 33 MW RS07990 WP 001788899.1          TIKVIENNKIR.FLKLKVGKIINVAKVDIIAFDKKGNINKFGSITIDNNNSIYRIIFHIEKGRIRYEKL   148
NTDB id 17 SA RS07760 WP 001788899.1          TIKVIENNKIR.FLKLKVGKIINVAKVDIIAFDKKGNINKFGSITIDNNNSIYRIIFHIEKGRIRYEKL   148
consensus                                     **!!*!!!!** !!******!!***!!****!!!*!!!**!!!!!!**!***!**!!!!!!!!!!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved