```
logo
                                                 MSIRHFFTFFWATLLTL.......FSMQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPTQ....QSGTGVKVN
NTDB id 1014 ACIAD RS13855 WP 004924483.1        MSIRHFFTFFWATLLTL.......FSMQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPTQ....QSGTGVKVN    69
NTDB id 1069 ABD1 RS03060 WP 043960841.1         ...MQYSMRFWKNKYFVLIMLCLGVISPSVCAQSFDQNFQEWKVKQQMYDQKLNISKSSHSNSSKISQTKNFNDSTGQIH    77
NTDB id 201180 KAB08 RS03015 WP 002017231.1     ...MQYSMRFWKNKYFVLIILCLGVISPSVYAQSFDQNFQEWKAKQQMYDQKLNITKSSHSNGSKISQTKNFNDSTGQIH    77
NTDB id 1041 HON27 RS14740 WP 005124713.1        ...MQYSMRFWKNKYFLLIMLCLGVISPSVCAQSFDQNFQEWKAKQQMYDQKLNITKSSHSNGSKISQTKNFNDSTGQIH    77
consensus                                          ******!!*******************!*!!*!!!****!!**!***!**!!***!*!!* ****************

logo
                                                 INSATAKELQQLHGVGEKKALAIVEYRQQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI
NTDB id 1014 ACIAD RS13855 WP 004924483.1        INSATAKELQQLHGVGEKKALAIVEYRQQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI    128
NTDB id 1069 ABD1 RS03060 WP 043960841.1         LNQANVNEFQQLKGIGEKKAQAIVEYRQKNGSFKNIDEIKNVKGIGPAIFEKNKSRLAL    136
NTDB id 201180 KAB08 RS03015 WP 002017231.1     LNQANVNEFQQLKGVGEKKAQAIVEYRQKNGSFKNIDEIKNVKGIGPAIFEKNKSRLAL    136
NTDB id 1041 HON27 RS14740 WP 005124713.1        LNQANVNEFQQLKGVGEKKAQAIVEYRQKNGSFKNIDEIKNVKGIGPAIFEKNKSRLAL    136
consensus                                        *!*!***!*!!!*!*!!!!!*!!!!!!!**!*!***!****!!!!!***!!!**!!!*
```

X non conserved

X similar

X $\geq$ 50% conserved