```
logo
NTDB id 167331 AL533 RS19095 WP 004711686.1  MTHLRSKYIV.QHHQKGATLIVVLIILLIVIAVGVLAIRVAIVSLKVATNSQIGQLNFQSSDTPIQLITQMDPTTLTNIS   79
NTDB id 1065 ABD1 RS15735 WP 000149372.1     MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNIT   80
consensus                                    !!!!  ! ! * **  !!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!* !! !! !!!!!!!*

logo
NTDB id 167331 AL533 RS19095 WP 004711686.1  NVLGAALKENESHPGSEYNFCYKPVSTTVSFAQTRDASLLRAGTANNAVVEDGGVAGFCDLTTDYGSNRQAVVTQIAVSI  159
NTDB id 1065 ABD1 RS15735 WP 000149372.1     NVIGAALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSI  160
consensus                                    !!*!!!!! *!!*!! !!!!!!!! !!*!!!!!!! !!!!!!!*!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!*!!!!

logo
NTDB id 167331 AL533 RS19095 WP 004711686.1  PTDAVNDVPGSNLPRGTNASEGTALPKSMLSTQRIRVISTSFLPAYASTSMQTLQADCLSTNSAKISDNFDSSLSNKQTL  239
NTDB id 1065 ABD1 RS15735 WP 000149372.1     PTDAMADIPGSNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTL  240
consensus                                    !!!!* !*!!!!!!!  ! !!!! !!!!!!!!!!!!!!*! !!!!!!!!!*  !! !!!!! !!!!!!!!!! !* !!!!

logo
NTDB id 167331 AL533 RS19095 WP 004711686.1  ADCLANHNVPFSTQIQEFNYTNKLTETMAPGS  271
NTDB id 1065 ABD1 RS15735 WP 000149372.1     ADCLANHNVPFSTQVQEFNYTNKLTEITAPGS  272
consensus                                    !!!!!!!!!!!!!!*!!!!!!!!!!!  !!!!
```

X non conserved

X similar

X $\geq$ 50% conserved