```
logo
NTDB id 99592 AL533 RS19095 WP 004711686.1  MTHLRSKYIV.QHHQKGATLIVVLIILLIVIAVGVLAIRVAIVSLKVATNSQIGQLNFQSSDTPIQLITQMDPTTLTNISNVLGA   84
NTDB id 1065 ABD1 RS15735 WP 000149372.1    MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNITNVIGA   85
consensus                                   !!!!  ! ! * **  !!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!* !! !! !!!!!!!*!!*!!

logo
NTDB id 99592 AL533 RS19095 WP 004711686.1  ALKENESHPGSEYNFCYKPVSTTVSFAQTRDASLLRAGTANNAVVEDGGVAGFCDLTTDYGSNRQAVVTQIAVSIPTDAVNDVPG   169
NTDB id 1065 ABD1 RS15735 WP 000149372.1    ALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSIPTDAMADIPG   170
consensus                                   !!! *!!*!! !!!!!!!! !!*!!!!!!! !!!!!!!*!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!*!!!!!!!!!* !*!!

logo
NTDB id 99592 AL533 RS19095 WP 004711686.1  SNLPRGTNASEGTALPKSMLSTQRIRVISTSFLPAYASTSMQTLQADCLSTNSAKISDNFDSSLSNKQTLADCLANHNVPFSTQI   254
NTDB id 1065 ABD1 RS15735 WP 000149372.1    SNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTLADCLANHNVPFSTQV   255
consensus                                   !!!!!  ! !!!! !!!!!!!!!!!!!!*! !!!!!!!!!*  !! !!!!! !!!!!!!!!! !* !!!!!!!!!!!!!!!!!!*

logo
NTDB id 99592 AL533 RS19095 WP 004711686.1  QEFNYTNKLTETMAPGS   271
NTDB id 1065 ABD1 RS15735 WP 000149372.1    QEFNYTNKLTEITAPGS   272
consensus                                   !!!!!!!!!!!  !!!!
```

X non conserved
X similar
X $\geq$ 50% conserved