```
logo

NTDB id 77630 H7R46 RS01675 WP 005067126.1  MTHL...IHKSRSKEKGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVSQLNFQSSDTPLELIVQMNPTTLTNITNVIGA   82
NTDB id 1065 ABD1 RS15735 WP 000149372.1    MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNITNVIGA   85
consensus                                   !!!!*** *! !!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 77630 H7R46 RS01675 WP 005067126.1  ALKEHESNPGAEYNFCYKPVSTATNFAQTRGASLLRAGSANNAVVEDGGVAGFCNLTTDYGSNRQAVVTQVAVSVPTDAASDIPG   167
NTDB id 1065 ABD1 RS15735 WP 000149372.1    ALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSIPTDAMADIPG   170
consensus                                   !!! !!!!!!!!!!!!!!! !!   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!*!!!!!!!!!!!!!!!!!*!!!!   !!!!

logo

NTDB id 77630 H7R46 RS01675 WP 005067126.1  SNLPRGTNTSEGTQLPKSMLSTQRIRVITTAFLPAYASTSIETLQRDCLSTSSAKISDNFDTALTAKQTLAECLANHNVPFSTQV   252
NTDB id 1065 ABD1 RS15735 WP 000149372.1    SNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTLADCLANHNVPFSTQV   255
consensus                                   !!!!!  ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*! !! !!!!!!!!!!!!!!!!*!!! !!!!!*!!!!!!!!!!!!!!

logo

NTDB id 77630 H7R46 RS01675 WP 005067126.1  QEFNYTNKLTQITAPGS   269
NTDB id 1065 ABD1 RS15735 WP 000149372.1    QEFNYTNKLTEITAPGS   272
consensus                                   !!!!!!!!!! !!!!!!
```

X non conserved

X similar

X $\geq$ 50% conserved