```
logo

NTDB id 294178 DLK06 RS01780 WP 005067126.1  MTHL...IHKSRSKEKGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVSQLNFQSSDTPLELIVQMNPTTLTNIT   77
NTDB id 1065 ABD1 RS15735 WP 000149372.1     MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNIT   80
consensus                                    !!!!*** *! !!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 294178 DLK06 RS01780 WP 005067126.1  NVIGAALKEHESNPGAEYNFCYKPVSTATNFAQTRGASLLRAGSANNAVVEDGGVAGFCNLTTDYGSNRQAVVTQVAVSV  157
NTDB id 1065 ABD1 RS15735 WP 000149372.1     NVIGAALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSI  160
consensus                                    !!!!!!!! !!!!!!!!!!!!!!!! !!   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!*!!!!!!!!!!!!!!!!*

logo

NTDB id 294178 DLK06 RS01780 WP 005067126.1  PTDAASDIPGSNLPRGTNTSEGTQLPKSMLSTQRIRVITTAFLPAYASTSHETLQRDCLSTSSAKISDNFDTALTAKQTL  237
NTDB id 1065 ABD1 RS15735 WP 000149372.1     PTDAMADIPGSNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTL  240
consensus                                    !!!!  !!!!!!!!!!  ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*! !! !!!!!!!!!!!!!!!!*!!! !!!!

logo

NTDB id 294178 DLK06 RS01780 WP 005067126.1  AECLANHNVPFSTQVQEFNYTNKLTQITAPGS   269
NTDB id 1065 ABD1 RS15735 WP 000149372.1     ADCLANHNVPFSTQVQEFNYTNKLTEITAPGS   272
consensus                                    !*!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!
```

X non conserved
X similar
X ≥ 50% conserved