```
logo

NTDB id 1043305 ACEN3H RS01790 WP 373358447.1  MTHL...IHKSRSKEKGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVSQLNFQSSDTPLELIVQMNPTTLTNIT   77
NTDB id 1065 ABD1 RS15735 WP 000149372.1       MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNIT   80
consensus                                      !!!!*** *! !!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1043305 ACEN3H RS01790 WP 373358447.1  NVIGAALKEHESNPGAEYNFCYKPVSTATNFAQTRGASLLRAGSANNAVVEDGGVAGFCNLTTDYGSNRQAVVTQVAVSI  157
NTDB id 1065 ABD1 RS15735 WP 000149372.1       NVIGAALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSI  160
consensus                                      !!!!!!!! !!!!!!!!!!!!!!! !!   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!*!!!!!!!!!!!!!!!!!

logo

NTDB id 1043305 ACEN3H RS01790 WP 373358447.1  PTDAASDIPGSNLPRGTNTSEGTQLPKSMLSTQRIRVITTAFLPAYASTSIETLQRDCLSTSSAKISDNFDTALTAKQTL  237
NTDB id 1065 ABD1 RS15735 WP 000149372.1       PTDAMADIPGSNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTL  240
consensus                                      !!!!  !!!!!!!!!  ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*! !! !!!!!!!!!!!!!!!!*!!! !!!!

logo

NTDB id 1043305 ACEN3H RS01790 WP 373358447.1  AECLANHNVPFSTQVQEFNYTNKLTQITAPGS   269
NTDB id 1065 ABD1 RS15735 WP 000149372.1       ADCLANHNVPFSTQVQEFNYTNKLTEITAPGS   272
consensus                                      !*!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!
```

X non conserved
X similar
X ≥ 50% conserved