```
logo

NTDB id 204255 BMU11 RS02425 WP 213033974.1  MTHLIY...KSRSKEKGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVSQLNFQSSDTPLELIVQMNPTTLTNIT   77
NTDB id 1065 ABD1 RS15735 WP 000149372.1     MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNIT   80
consensus                                    !!!!  !***!  !!!!  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!  !!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 204255 BMU11 RS02425 WP 213033974.1  NVIGAALKEHESNPGAEYNFCYKPVSTATNFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTRDYGSNRQAVVTQVAVSV  157
NTDB id 1065 ABD1 RS15735 WP 000149372.1     NVIGAALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSI  160
consensus                                    !!!!!!!!  !!!!!!!!!!!!!!!!  !!   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!  !!!!!!!!!!!!!!!!!*

logo

NTDB id 204255 BMU11 RS02425 WP 213033974.1  PTDAASDIPGSNLPRGTNTSEGTQLPKSMLSTQRIRVITTAFLPAYASTSHETLQRDCLSTSSAKISDNFDTALTAKQTL  237
NTDB id 1065 ABD1 RS15735 WP 000149372.1     PTDAMADIPGSNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTL  240
consensus                                    !!!!   !!!!!!!!!!   ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!  !!  !!!!!!!!!!!!!!!!*!!!  !!!!

logo

NTDB id 204255 BMU11 RS02425 WP 213033974.1  AECLANHNVPFSTQVQEFNYTNKLTQITAPGS  269
NTDB id 1065 ABD1 RS15735 WP 000149372.1     ADCLANHNVPFSTQVQEFNYTNKLTEITAPGS  272
consensus                                    !*!!!!!!!!!!!!!!!!!!!!!!!!  !!!!!!
```

X non conserved

X similar

X ≥ 50% conserved