```
logo

NTDB id 1143480 AC2117 RS01705 WP 197730966.1  MTHLIHENIKSKSKEKGATLIVVLIILLVVVSVGVLAIRVAIVSLKVATNSQVSQLNFQSSDTPLELIVQMNPTTLTNIT   80
NTDB id 1065 ABD1 RS15735 WP 000149372.1       MTHLNYKNIKGRSKETGATLIVVLIILLIVISVGVLAIRVAIVSLKVATNSQVGQLNFQSSDTPLELIVQMNPTTLTNIT   80
consensus                                      !!!! * !!! *!!! !!!!!!!!!!!!*!*!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1143480 AC2117 RS01705 WP 197730966.1  NVIGAALKEHESNPGAEYNFCYKPVSTSTNFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTNDYGSNRQAVVTQVAVSI  160
NTDB id 1065 ABD1 RS15735 WP 000149372.1       NVIGAALKAHESNPGAEYNFCYKPTSTSVSFAQTRGASLLRAGSANNAVVEDGGVAGFCDLTSDYGSNRQAVVTQVAVSI  160
consensus                                      !!!!!!!! !!!!!!!!!!!!!!! !!!  !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!!!!!!!!!!!!!

logo

NTDB id 1143480 AC2117 RS01705 WP 197730966.1  PTDAASDIPGSNLPRGTNTSEGTQLPKSMLSTQRIRVITTAFLPAYASTSIETLQSDCLSTSSAKISDNFDTALTAKQTL  240
NTDB id 1065 ABD1 RS15735 WP 000149372.1       PTDAMADIPGSNLPRDINLSEGTQLPKSMLSTQRIRVITTAFLPAYASTSLENLQFDCLSTSSAKISDNFDSALTDKQTL  240
consensus                                      !!!!  !!!!!!!!!  ! !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*! !! !!!!!!!!!!!!!!!*!!! !!!!

logo

NTDB id 1143480 AC2117 RS01705 WP 197730966.1  AECLANHNVPFSTQVQEFNYTNKLTQITAPGS  272
NTDB id 1065 ABD1 RS15735 WP 000149372.1       ADCLANHNVPFSTQVQEFNYTNKLTEITAPGS  272
consensus                                      !*!!!!!!!!!!!!!!!!!!!!!!!! !!!!!!
```

X non conserved
X similar
X ≥ 50% conserved