```
logo
NTDB id 1009463 ABJ384 RS01590 WP 349928513.1  MNNKFPATGLHQLGFTLVELIIALALGLLITAAATQLFLGGIVSARIQQANAELQDSGVFGLDYMARDIRLANYGNINNP   80
NTDB id 1066 ABD1 RS15740 WP 000079197.1       MAQSF..ISNKEQGFTLIELIVALALGLILVAAATQLFIGGLLSSRLQKANAEIQDSGIFGLEYMARDIRLLNYGNVVNP   78
consensus                                      ! * !**    *  !!!!*!!!*!!!!!!** !!!!!!!*!!**! !*! !!!!*!!!!*!!!*!!!!!!!! !!!!* !!

logo
NTDB id 1009463 ABJ384 RS01590 WP 349928513.1  ELTNTTQLGGIVLSQGASGSVNVP..................KDIGDSSVT....LQG.....LTSNVSEKSDQLTIQFS  133
NTDB id 1066 ABD1 RS15740 WP 000079197.1       QLTDTTPWGGIVLTGSTATNANNINFIPKVDTNTYIPEALLSRGAGDTVSTVNNHWKGLSNIQNSSNAEVQSDQLTIQFI  158
consensus                                       !! !!* !!!!!*       ! ********************  !!*  !****  !***** *!!    !!!!!!!!

logo
NTDB id 1009463 ABJ384 RS01590 WP 349928513.1  APNDMFNCEGEEVKTGEYVIQRYFLRKDNNGEE.KDLVLACDANKKGAMSVTNIT.DFGGTQLGQVIMPRVDHLRFYLGT  211
NTDB id 1066 ABD1 RS15740 WP 000079197.1       APTNMTNCEGVNVLAGDLIVQRYFLRVDNNGSSQQDYALACDANTPAVSATAQPDIVNGLGDAGQIILPRIDHFHVLLGA  238
consensus                                      !!   ! !!!!  !  !* **!!!!!! !!!!  * !  !!!!!! **     ** *  !    !!*!*!!*!! *  !!

logo
NTDB id 1009463 ABJ384 RS01590 WP 349928513.1  MTKNDSDVISTTYYTINEYITVANAARNAGQ..LVPRITSIKAAVIVRSVDNTNNQLVDPTKPIQFLDKSVTL.TNQTKH  288
NTDB id 1066 ABD1 RS15740 WP 000079197.1       KNAA....GNFAYYTIPQYRVAAQAARDASPAVAAPRILSIQISVLARSTNNAQNKAIDPNQSFLMLDQNVHAADNRTRF  314
consensus                                          ****    !!!!* !   !*!!! ! ***  !!! !!   !* !!  ! *!  *!!  *   !!  !* * ! !**

logo
NTDB id 1009463 ABJ384 RS01590 WP 349928513.1  LRRVYMTTIAIRNGMGDAI   307
NTDB id 1066 ABD1 RS15740 WP 000079197.1       LRRVYSVTIALRNAMGETI   333
consensus                                      !!!!!  !!!*!!*!!* !
```

X non conserved

X similar

X $\geq$ 50% conserved