```
logo

NTDB id 1006917 ABLB99 RS01615 WP 004884688.1  MNRSF.LEKQQGFTLIELMVALALGLILVAAATQLFIGGILSSRLQKANAEIQDSGIFGLDYIIRDIRLVNYGNIINPQL   79
NTDB id 1066 ABD1 RS15740 WP 000079197.1       MAQSFISNKEQGFTLIELIVALALGLILVAAATQLFIGGLLSSRLQKANAEIQDSGIFGLEYMARDIRLLNYGNVVNPQL   80
consensus                                      !  !!*  ! !!!!!!!!*!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!*!* !!!!!*!!!!**!!!!

logo

NTDB id 1006917 ABLB99 RS01615 WP 004884688.1  NDTTPWGGIVLTGSTATNNNNVNFVPNTGTNAYIADALLSRGAGDTVSTLNNQWKGLSNIQNSSNATIQSDQLTIQFIAP  159
NTDB id 1066 ABD1 RS15740 WP 000079197.1       TDTTPWGGIVLTGSTATNANNINFIPKVDTNTYIPEALLSRGAGDTVSTVNNHWKGLSNIQNSSNAEVQSDQLTIQFIAP  160
consensus                                       !!!!!!!!!!!!!!!!!! !!*!!*!   !! !!**!!!!!!!!!!!!!!*!!*!!!!!!!!!!!!!! *!!!!!!!!!!!!

logo

NTDB id 1006917 ABLB99 RS01615 WP 004884688.1  SNMTNCEGANVLAGDLVVERYFLRLDNNGSSQQDYALACDANTPSTTPTAQPTSVDGFGDAGQIILPRIDHFHVLLGAKN  239
NTDB id 1066 ABD1 RS15740 WP 000079197.1       TNMTNCEGVNVLAGDLIVQRYFLRVDNNGSSQQDYALACDANTPAVSATAQPDIVNGLGDAGQIILPRIDHFHVLLGAKN  240
consensus                                      *!!!!!!! !!!!!!!*! !!!!!*!!!!!!!!!!!!!!!!!!!!   **!!!!  ! ! !!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1006917 ABLB99 RS01615 WP 004884688.1  AAGNFAYYTIPQYRVAAQAARDATPTQLAPRILSIQISVLARSTNNAQNKAVDPNQTFVMLDQTVHASDTSSRFLRRVYS  319
NTDB id 1066 ABD1 RS15740 WP 000079197.1       AAGNFAYYTIPQYRVAAQAARDASPAVAAPRILSIQISVLARSTNNAQNKAIDPNQSFLMLDQNVHAADNRTRFLRRVYS  320
consensus                                      !!!!!!!!!!!!!!!!!!!!!!!!!*!   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!*!*!!!! !!! !  *!!!!!!!!

logo

NTDB id 1006917 ABLB99 RS01615 WP 004884688.1  TTIALRNAMGEKL   332
NTDB id 1066 ABD1 RS15740 WP 000079197.1       VTIALRNAMGETI   333
consensus                                       !!!!!!!!!! *
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X non conserved
X similar
X ≥ 50% conserved