```
logo

NTDB id 1145821 CS896 RS06740 WP 089087212.1  MKKSLYLSFFLTFSNPLQALVIELLEEIKTSSHKGTFKAKVLDSKEPKQVLGIYNISPHKKLTLTITHISTAIVYQPLDE   80
NTDB id 1207 C694 RS07860 WP 000749005.1      MKKSLCLSFFLTFSNPLQALVIELLEEIKTSPHKGTFKAKVLDSKKPRQVLGVYNISPHKKLTLTITHISTAIVYQPLDE   80
consensus                                     !!!!!*!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!! !*!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo

NTDB id 1145821 CS896 RS06740 WP 089087212.1  KLSLETTLSPNRPTIPRNTQIVFSSKELKEPHSNPIPSLNAPMQKPQNKPSSSQQSPQNFSYPESKLGSKNSKNSLLQPL   160
NTDB id 1207 C694 RS07860 WP 000749005.1      KLSLETTLNPNRPTIPRNTQIVFSSKELKESHPHQMPSLNAPMQKPQNKPHSSQQPSQNFSYPEPKLGSKNSKNSLLQPL   160
consensus                                     !!!!!!!! !!!!!!!!!!!!!!!!!!!!!!*!****!!!!!!!!!!!!!!!!*!!!!**!!!!!!!*!!!!!!!!!!!!!!!!

logo

NTDB id 1145821 CS896 RS06740 WP 089087212.1  VTPSKISPANEVKTPTNDANPPLKHSSEDQENNLFVTPPTEKTLPNNTPNADASENNENNENNENNRDNVEKQAIRDPNI   240
NTDB id 1207 C694 RS07860 WP 000749005.1      AIPSKISPTNETQTPTNDTKPPLKHSSEDQESNLFITPPTEKTLPNNTSNADISENNESNE....NKDNVEKQAIRDANI   236
consensus                                       !!!!!! !!  !!!!!  !!!!!!!!!!! !!!*!!!!!!!!!!!!!*!!! !!!!! !!****!*!!!!!!!!!!*!!

logo

NTDB id 1145821 CS896 RS06740 WP 089087212.1  KEFACGKWVYDDENLQAYRPSVLKRVDKDKEVTTDITPCDYSTAENKSGKIITPYTKISVHKTEPLEDPQTFEAKNNFAI   320
NTDB id 1207 C694 RS07860 WP 000749005.1      KEFACGKWVYDDENLQAYRPSILKRVDEDKQTATDITPCDYSTAENKSGKIITPYTKISVHKTEPLEEPQTFEAKNNFAI   316
consensus                                     !!!!!!!!!!!!!!!!!!!!!!*!!!!! !!   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!

logo

NTDB id 1145821 CS896 RS06740 WP 089087212.1  LQARSSTEKCKRARARKDGTTRQCYLIEEPLKQAWESEYEITTQLVKAIYERPKQDDQIEPTFYETSELAYSSTRKSEIT   400
NTDB id 1207 C694 RS07860 WP 000749005.1      LQARSSTEKCKRARARKDGTTRQCYLIEEPLKQAWESEYEITTQLVKAIYERPKQDDQVEPTFYETSELAYSSTRKSEIT   396
consensus                                     !!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X $\geq$ 50% conserved