```
logo
NTDB id 286 KZH43 RS05975 WP 000782664.1   MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKIDQNLD   85
NTDB id 285 SPD RS06395 WP 000782664.1     MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKIDQNLD   85
NTDB id 284 SPR RS06100 WP 000782664.1     MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKIDQNLD   85
NTDB id 283 SP RS06670 WP 000782668.1      MKMKQISDTTLKITISLEDLMDRGMEIADFLVPQEKTEEFFYAILDELEMPDSFLDTGMLSFRVTPKPDKVDVFVTKSKIDQNLD   85
NTDB id 299 STER RS01050 WP 011680667.1    MEMKQISETTLKIMITMEDLEEHGMELKDFLIPQEKTEEFFYTVMDELDLPDNFKNSGMLSFRVTPRKDRVDVFVTKSDLKEDLD   85
NTDB id 331 STU RS10485 WP 011225365.1     MEMKQINETTLKIMITMEDLEEHGMELKDFLIPQEKTEEFFYTVMDELDLPDNFKNSGMLSFRVTPRKDRVDVFVTKSDLKEALD   85
NTDB id 60807 KE3 RS01295 WP 020916168.1   MEMKQISETTLKITISMEDLEERGMELKDFLIPQEKTEEFFYSVMDELDLPDNFKDSGMLSFRVTPRKDRIDVFVTKSEINKDIN   85
NTDB id 380 SMU RS01250 WP 002262749.1     MEMKQISETTLKITISMEDLEERGMELKDFLIPQEKTEEFFYTVMDELDLPENFKDSGMLSFRVTPRNDRIDVFVTKSEINKNLN   85
consensus                                  ! !!!!**!!!!!*!**!!! **!!!* !!!*!!!!!!!!!! **!!!**!* ! **!!!!!!!!!* !**!!!!!!! *  ***

logo
NTDB id 286 KZH43 RS05975 WP 000782664.1   FEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEAKEEE...........QEQAEQEAESKKEPYIYYILSF   159
NTDB id 285 SPD RS06395 WP 000782664.1     FEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEAKEEE...........QEQAEQEAESKKEPYIYYILSF   159
NTDB id 284 SPR RS06100 WP 000782664.1     FEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEAKEEE...........QEQAEQEAESKKEPYIYYILSF   159
NTDB id 283 SP RS06670 WP 000782668.1      FEDLSDLPDMEELAQMSPDEFIKTLEKSIADKTKDDIEAIQSLEQVEVKEEE...........QEQAEQEAESKKEPYIYYILSF   159
NTDB id 299 STER RS01050 WP 011680667.1    FNDLSDM...EDYSGLSPEEFLKALEDNFMDK..GDIEAHHKLEEHDKTLKEVDET.MT....EPAKEVAEETIREDYTHYVLAF   160
NTDB id 331 STU RS10485 WP 011225365.1     FNDLSDM...EDYSGLSPEEFLKALEDNFMDK..GDIEAHHKLEEHDKTLKEVDET.MT....EPAKEVAEETIREDYTHYVLAF   160
NTDB id 60807 KE3 RS01295 WP 020916168.1   FEDLAAF...DDVSNMSPEEFFKTLEQTMLSK..DDTEAHEKLGEIEEMMEGAVEDVLTEHAQQAEIEPEEDVNPSDYVHYVLDF   165
NTDB id 380 SMU RS01250 WP 002262749.1     LEDLSDF...DDISKMSPEDFFNTLEETMREK..GDAAALDKLAEIEKREEEK...........TQQEKGETKEKRDYVHFVLDF   154
consensus                                  **!!***   *** **!!**!***!! ** *!  *!**!*  !* ** ****           * * ! * ** ** !* **! !

logo
NTDB id 286 KZH43 RS05975 WP 000782664.1   AKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEYGQVLMSHDAVLNLQK   243
NTDB id 285 SPD RS06395 WP 000782664.1     AKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEYGQVLMSHDAVLNLQK   243
NTDB id 284 SPR RS06100 WP 000782664.1     AKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEYGQVLMSHDAVLNLQK   243
NTDB id 283 SP RS06670 WP 000782668.1      AKLADLVAFAKTV.TFEMETSELYKMNERYYLTILVDIENHPSPYPAWLLARMREFADDSDISRSVLQEYGQVLMSHDAVLNLQK   243
NTDB id 299 STER RS01050 WP 011680667.1    SDFDQVVTFAQGLKNVSVEGSEFYKLGDVYYMTILLYLADEPDYYANNMYARFLEYANVADRTRPYLQEHATILMEEDALPVLQA   245
NTDB id 331 STU RS10485 WP 011225365.1     SDFDQVVTFAQGLKNVSVEGSEFYKLGDVYYMTILLYLADEPDYYANNMYARFLEYANVADRTRPYLQEHATILMEEDALPVLQA   245
NTDB id 60807 KE3 RS01295 WP 020916168.1   PSLEAAVAFAKVI.DFPVEASEFYKMDGTYYMTVLIDLQNHPSYYANLMYARMLEYAGAGTKTRAYLQEHAVELLIDDAVEKLKM   249
NTDB id 380 SMU RS01250 WP 002262749.1     PNIQQVINFAKTV.DYDVEASELFKESDAYHMTVLLNLEDKPDYYADLMFARMLEHAGRGTKTRAYLLEHGVQLIKADALQELQM   238
consensus                                    *  ***!!***  ***! !!**!* * !**!*!******!* !   * !!* !*!   * *!  !*! * *!** !!*  !*

logo
NTDB id 286 KZH43 RS05975 WP 000782664.1   IG..   245
NTDB id 285 SPD RS06395 WP 000782664.1     IG..   245
NTDB id 284 SPR RS06100 WP 000782664.1     IG..   245
NTDB id 283 SP RS06670 WP 000782668.1      IG..   245
NTDB id 299 STER RS01050 WP 011680667.1    TKWS   249
NTDB id 331 STU RS10485 WP 011225365.1     TKWS   249
NTDB id 60807 KE3 RS01295 WP 020916168.1   IELV   253
NTDB id 380 SMU RS01250 WP 002262749.1     IG..   240
consensus                                  **
```

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