```
logo
NTDB id 103229 QUE23 RS01655 WP 286348961.1  MLSPFTPSPLDFLPLSFKALKKPPKLLHYVGDLKHLQAPLKVAIVGTRKPTTYTQQITATLAKELAKIGAVVVSGGALGV   80
NTDB id 1210 C694 RS01710 WP 001862983.1     MKSHFQYSTLENIPKAFDILKDPPKKLYCVGDTKLLDTPLKVAIIGTRRPTPYSKQHTITLAREL AKNGAVIVSGGALGV   80
NTDB id 1239 Cj0634 YP 002344063.1           MASEILP...DEFLELFKDLKKPPKKLHYKGNLSLLKQ.DKIAIIGSRRMSVYTKNCVFSLASMLKNAHLCVVSGGALGV   76
consensus                                    ! ! * ** ** **  !* !!*!!!*!***!****!  **!*!!*!*!*** !*** * *!!**!** ****!!!!!!!

logo
NTDB id 103229 QUE23 RS01655 WP 286348961.1  DIIAQKNALPLTIMFAPCSLKRIYPPSNAEVIQQIAKVGLILSEYAEDFQPYRSTFLERNRLVIALSDLVIIPEAQETSG  160
NTDB id 1210 C694 RS01710 WP 001862983.1     DIIAQENALPKTIMLSPCSLDFIYPTNNHKVIQEIAQNGLILSEYEKDFMPIKGSFLARNRLVIALSDVVIIPQADLKSG  160
NTDB id 1239 Cj0634 YP 002344063.1           DITASMAAMPNTIGIFANGLDQIYPRTNEKIIKQIYENALALSEYEDDYLPKNYDFLLRNRLVIALSKAVVVAQADIKSG  156
consensus                                    !!*!* *!*! !!** ***!* !!! *! **!**!* **!*!!!!*!**! * *!! !!!!!!!!!**!****!***!!

logo
NTDB id 103229 QUE23 RS01655 WP 286348961.1  SMASARLALELKKPLFVLPQRLFESPGTNALLAQNKATAIYDIAAFVGDLARKFGLSAPTTPPTAPVPNDFLAFCQKSPS  240
NTDB id 1210 C694 RS01710 WP 001862983.1     SMSSARLAQKYQKPLFVLPQRLNESDGTNELLEKGQAQGIFNIQNFINTLLKDYHLKE.....MPEMEDEFLEYCAKNPS  235
NTDB id 1239 Cj0634 YP 002344063.1           SMQSAKLALELNKPLYVLPQRLGESTATNLLLKENKAKLICDFKEFVSEFAS...........IDMNQDEFLEFCKKGVS  225
consensus                                    !! !!*!!***!!!*!!!!!! !! *!! !!  **! *!***  !* **** * *       *  * **!!**! ! *!

logo
NTDB id 103229 QUE23 RS01655 WP 286348961.1  FEEAYAVFGDLLLEQELLGVVKREHGRVVLL.   271
NTDB id 1210 C694 RS01710 WP 001862983.1     YEEAYLKFGDKLLEYELLGKIKRINHIVVLA.   266
NTDB id 1239 Cj0634 YP 002344063.1           VDEALKIYGQKVYEYELEGKISIEGLFIRVLA   257
consensus                                    **!!* **!****!*!!*!*****   ****
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X non conserved

X similar

X ≥ 50% conserved