```
logo
NTDB id 1170 A1552VC RS11080 WP 000418747.1  MELFYFYPWLFPVLATLFGLIVGSFLNVVIYRLPKIMEREWRAECAASFPEYGITPPEGKLTLSLPRSTCPHCQTPIRVI   80
NTDB id 360470 FAY30 RS17845 WP 149871142.1 ........MVVILILFLYGLILGSFFNVIGLRIPL.......................KESIVTPSSACPTCGHQLKPY   48
NTDB id 103 BSU 28070 NP 390685.2            ..........MLSILFIFGLILGSFYYTAGCRIPL.......................HLSIIAPRSSCPFCRRTLTPA   46
consensus                                              *********!!!*!!!***** !*!*                       ***** !*!*!! !   ***

logo
NTDB id 1170 A1552VC RS11080 WP 000418747.1  DNIPLLSWLALRGQCSHCKAPISARYPLIELLTALMSLVIATHFPFGVFAVALLFFSYVLIAATFIDFDTLLLPDQLTLP  160
NTDB id 360470 FAY30 RS17845 WP 149871142.1 ELIPVLSYLFQRGKCRGCQSRISPIYPIFELLTGLLFMTAPLVIGWSGELVVALTLISMFMIIVVSDIHYMIIPDKILIW  128
NTDB id 103 BSU 28070 NP 390685.2            ELIPILSFLFQKGKCKSCGHRISFMYPAAELVTACLFAAAGIRFGISLELFPAVVFISLLIIVAVTDIHFMLIPNRILIF  126
consensus                                    **!!*!!*!***!*!* !   *!! *!!* !!*!***** *** ******** ** ******** **!******!******

logo
NTDB id 1170 A1552VC RS11080 WP 000418747.1  LLWGGIALALLGFSPVSLSDAVIGAMAGYLSLWSIYWLFKLLTGKEGMGYGDFKLLAALGAWLGWQQLPVIVLLSSVVGV  240
NTDB id 360470 FAY30 RS17845 WP 149871142.1 F.AGIFLLERIVWPLSPWWDSLLGAITGFVVLLL.....IALVSKGGMGGGDVKLYAVLGLVLGVKLVLLSFFLSTLLGA  202
NTDB id 103 BSU 28070 NP 390685.2            F.LPFLAAARLISPLDSWYAGLLGAAAGFLFLAV.....IAAITHGGVGGGDIKLFAVIGFVLGVKMLAAAFFFSVLIGA  200
consensus                                    *  * ********** *******!!**!** ! *     *******!*!*!!*!!*!**! *!!**** * ***!***!*

logo
NTDB id 1170 A1552VC RS11080 WP 000418747.1  IFGLIQLRQQKKGIDMAFPFGPYLAIAGWFALLWGDKVIDWYFTTWVGQPL  291
NTDB id 360470 FAY30 RS17845 WP 149871142.1 VFGGLALLLKIVKRRQPVPFGPFIAAGTLIAYYWGSEIIDFYIQFLQQGF.  252
NTDB id 103 BSU 28070 NP 390685.2            LYGAAAVLTGRLAKRQPLPFAPAIAAGSILAYLYGDSIISFYIKMALG...  248
consensus                                    **!*****  ********!!*!**!*****!***!* *!**!*   **
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X non conserved

X similar

X ≥ 50% conserved