logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	MSETKQLPLRSEVPEELTWDLSVIFKDDAAFEEAFAALNDRLAEVGNYQGILANGAKAFLAALEYCLDVMRQLETLYVYS	80
NTDB id 480 HSISS4 RS02090 WP 002890133.1	.....MSDNRTHLEEKYTWDLTTIFATDADWETEYESTVQDLKKASAYAGHLLLDSAKNLLGATELYMSLMRRLEKIYVYA	75
consensus	***** * !**** !! !! !! !! !!*! !** *!! !! *!!!

logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	HLRNDQDTTNATDQALHSRANALVAKASEAIAWFEPEVLALEDAKIWGYFDELPELSLYRHYIQQIVDNREHILPAEQEA	160
NTDB id 480 HSISS4 RS02090 WP 002890133.1	SMKNDQDTTVGLYQEYQAKASNLYSQLSEAFAYFEPEFMALDDKKLAEFKEQEPGLGLYDHYFERLLANKDHVLSQEAEE	155
consensus	***!!!!! ! * *! ! !!! !!** * !!!* !! !! *** !* !

logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	LLAGAGEIFGAAADTFSVLNNADLEFPVIEDEKGEKVQLSHGVYGQLLESTDRRVRKDAFEGLYKVYQQFARTFASTLST	240
NTDB id 480 HSISS4 RS02090 WP 002890133.1	LLAAAGDIFNGPTDTFNVLDNADILFPWVSDGQGDVVELTHGNFITLMESKDRDIRKGAYEAMYGTYEQFQHTYAQTLQG	235
consensus	!!!!!*! !! ** !!! !!! !! *! !* *!! !! *!! !!**! ! !!*! !!

logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	NVKRHNYAAKVRHFESARQAALSGNHIPESVYETLVSVVNENLPLLHRYMALRKRLLNLDEMHMYDVYTPLLGESSLKYT	320
NTDB id 480 HSISS4 RS02090 WP 002890133.1	VVKVHNYQAKVRHYNSARHAALAANFIPESVYDSLLESVNKHLPLLHRYLDLRKKVLGLDELKMYDVYTPLSE.TETALT	314
consensus	!! !!!* !!!! ! ** !! *!*!!!** !! !! !!** !!! !!!*!!!!!!!! **

logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	YEEATKEAFKALEPMGEEYLAVVKEAFENRWIDVVENKGKRSGAYSSGCYDTNPYILLNWHDTLDQLFTLVHEMGHSVHS	400
NTDB id 480 HSISS4 RS02090 WP 002890133.1	YEESLKKAEEVLAIFGEEYSKGVHAAFTERWIDVHPNKGKRSGAYSGGAYDTNAFMLLNWQDTLDNLFTLVHETGHSLHS	394
consensus	!!! !! !! * !!! !! * !! !!* !*!!!!!!* !!** !!!!* *!!!!!!!!! !!*!!

logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	YFTRKNQPYVYGDYSIFLAEIASTTNENILTEYLLETQKDPKVRAFVLNHYLDGFKATVFRQTQFAEFEHFIHVEDSKGT	480
NTDB id 480 HSISS4 RS02090 WP 002890133.1	TFTRQTQPYVYGDYPIFLAEIASTTNENILTETLLKEVKDDKTRFAILNHYLDGFKGTVFRQTQFAEFEHAIHEADASGQ	474
consensus	!!! !!!!!!!!* !!!!!!!!!!!!!!! !! !!*! * !!!!!!!*!!!!!!!!! !! !

logo	
NTDB id 1154974 QML79 RS11910 WP 281956400.1	PLTAEFLNDYYGNLNKRYYGDAVTSDPQISYEWSRIPHFYYNYYVYQYATGFSAASALAKKILSKEENALENYLNYLKAG	560
NTDB id 480 HSISS4 RS02090 WP 002890133.1	ILTADFMNKLYADLNEKYYNLKAEDNYEIQFEWERIPHFYMNYYVYQYATGFAAASYLAEKIVHGTEEDKEAYLTYLKAG	554
consensus	*!!!*!*! !! !* !! *!! ! ! *!! !!!!!! !!!!!!!*!! !! !!** ! !! !!!!!

```
                                          SSDYPLEVMKKAGVDMTNTDYLKEAMQVFEQRLvELEALVSKLVK
                                          NN   t I   i     A   KA D  DAAFK  AD L D     EEGEHLS
```

NTDB id 1154974 QML79 RS11910 WP 281956400.1   NSDYPLEVMKKAGVDMTKADYLKEAMQVFAQRLDELEALVSELEK..   605
NTDB id 480 HSISS4 RS02090 WP 002890133.1   SSDYPLEVIKKAGVDMTNTDYLDAAFKVFEDRLVELEALVEKGVHLS   601
consensus   !!!!*!!*!!!!!!!!   !!! ! !!   !! !!!!!!   ***