```
logo

T6374 YP 007771623.1 NC 021010:296297-296785  ..MIRNGFYIIKDRFFSDMSDPYLKGNKK......QNRPHYYCFEDSNYNGIYWMIPLSSRIDKYKKIVSKRTGKGRNCD   72
T6372 WP 015513411.1 NC 021009:1041468-1041950  ..MEIGHFYYIDDQYFKDFPDSYLMQNKEKVNGQLHDRPCFYAFQDSN.TQLFWMIPFSSQTSKFKGIYNKKMQKYHRCD   77
T6373 WP 015528542.1 NC 021015:c1349281-1348793  MIYQEGYVYHIKDEYFEKVQDSNLMQNKEGG....TYRPTFYCLRDNK.TSLLWMVPLSSRVEKFKAIHDKQVAKYGKCL   75
consensus                                       *  * *! !*! *! ** !**!**!!*        !! *!** !** * *!!*!*!!**!*!*!  !***!* *!*

logo

T6374 YP 007771623.1 NC 021010:296297-296785  IIHIVKLDDSHESAFLIQDMFPISDKYIEREYTIAGNHLRLTSE.HAAKEIEQKARKVLGMLKRGIKFTPTQPDIQKIYE   151
T6372 WP 015513411.1 NC 021009:1041468-1041950  TIVFGEV.LGHEKAFLIQNMCPITEKYMKNEYLDAVANIPVRVDGRLEKEIKDKAGKVLALQRKGAKLIF..PDVLSIEQ   154
T6373 WP 015528542.1 NC 021015:c1349281-1348793  TIVLGEF.DGKEAAFLLQNMFPIREYYLDHIH..TRNNNPVPVKHSIHREVTTRMKKIRQLHSRCKKVVF..PDIDRLEQ   150
consensus                                       *!***** ***! !!!*!*!*!!***!** **  * ***** **  * *!*  ***!**** **! !***  !!* ****

logo

T6374 YP 007771623.1 NC 021010:296297-296785  RLQQDLPATHL.   162
T6372 WP 015513411.1 NC 021009:1041468-1041950  ELLKYM......   160
T6373 WP 015528542.1 NC 021015:c1349281-1348793  IMLAEVKDNATE   162
consensus                                      ** **
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X non conserved

X similar

X ≥ 50% conserved