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

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

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

X similar

X ≥ 50% conserved