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logo
T6365 AC98713.1 AF001314:3611-4162          MSSFFYKEILRMTLRFFTVTDEYIAYLRKFESKVHYQYE...NNASTYVGVVLKKNDFNYFIPLSSYKKGNPEKDKAMKK   77
T6366 WP 000182254.1 NC 020240:c63383-62799 .......MTNKDNPKFHTISTEYIDYLREADSKVPFNKDEQ..HSRPYVGVLEKINGHDYFVPLTSRNDKNFN....SQV   67
T10071 WP 000675353.1 NZ LDDO01000030:1726-2241 .............MKFYTISSSYIKYLKDFDDKVPNSEDPTYNNPKAFIGIVLEIEGHKYLAPLTSPKAWHAN....VKE   63
T6317 WP 012609144.1 NC 011767:1958-2473    .............MKFYTISSKYIEYLKEFDDKVPNSEDPTYQNPKAFIGIVLEIQGHKYLAPLTSPKKWHNN....VKE   63
consensus                                             *  **!*!****!!*!!**** !!***************!*** *****! *!!*!***** *    ***

logo
T6365 AC98713.1 AF001314:3611-4162          RSRIVTRLFEIGNINNPLGYLLHHNMIPVPDSELIPLPLDL....KKPKHKMMQKQLIYMKSISEKIENKSEVVYRKAAH   153
T6366 WP 000182254.1 NC 020240:c63383-62799 S....VKLFD..NDEKRIGVLLVNNMIPVPEKECKEIDIAEKTAADPQYGNLMLKQYLFLKENMDRVTNKVEKVYKDVTV   141
T10071 WP 000675353.1 NZ LDDO01000030:1726-2241 SSPAFFKLHENGVPDNQLGLINLKFMIPIIEAEVSLLDLDS..MPDTPYKRMLYKQLQFIRVNEDKISEKSKL.LRNLAL   140
T6317 WP 012609144.1 NC 011767:1958-2473    SSLSCFKLHENGVPENQLGLINLKFMIPIIEAEVSLLDLGN..MPNTPYKRMLYKQLQFIRANSDKIASKSDT.LRNLVL   140
consensus                                   **    **!**** *****!** ** !!!****!*******   ***********!!***** *****!**** *****

logo
T6365 AC98713.1 AF001314:3611-4162          .....EKDGYYLKFSCDFKLLEAKATLYSKKSTFQ...................   183
T6366 WP 000182254.1 NC 020240:c63383-62799 QGKPSHKQKFLKGVCCDFPKLEEKCQEYKERDQAKERDKARRIAYMRQMGRER   194
T10071 WP 000675353.1 NZ LDDO01000030:1726-2241 QGRM........QGTCDFAVLEEKYQHFGKKPEDMEIDD..............   171
T6317 WP 012609144.1 NC 011767:1958-2473    QGKM........QGTCNFSLLEEKYRDFGKEAEDTEEGE..............   171
consensus                                   ****  *  *  ***!*! *!!*!******* ** * **
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X non conserved

X similar

X $\geq$ 50% conserved