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logo
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NTDB id 1127 NGFG RS10560 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM.FV..................LF..CMLFSCAFSLAAVNINAAS   59
NTDB id 1126 NGFG RS09375 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM.FV..................LF..CMLFSCAFSLAAVNINAAS   59
NTDB id 1125 NGFG RS07215 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM.FV..................LF..CMLFSCAFSLAAVNINAAS   59
NTDB id 1108 NGFG RS06525 WP 003703428.1      MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM.FV..................LF..CMLFSCAFSLAAVNINAAS   59
NTDB id 1014 ACIAD RS13855 WP 004924483.1     ......MSIRHFFTFFWATLLTLFSMQIVFAQGFDQQYDAWKTQQKAHDTRLNSSASHHSLSKPTQQSGTGVKVNINSAT   74
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    ......MQIKTKIVTLFLSLCLPTLPLLANAEETAP.......AAQ...................VEEGIVITVNINTAS   48
NTDB id 1150 A1552VC RS08580 WP 001166105.1   ......MQIKTKIVTLFLSLCLPTLPLLANAEETAP.......AAQ...................VEEGIVITVNINTAS   48
NTDB id 115446 VPUCM RS05150 WP 021449594.1  .............MKWILTLCLVVFAPLSLAA..DT.......KAD...................KYEGIEITVNINTAS   39
NTDB id 1294 VP RS04495 WP 021449594.1       .............MKWILTLCLVVFAPLSLAA..DT.......KAD...................KYEGIEITVNINTAS   39
NTDB id 1395 DSB67 RS04445 WP 010649155.1    .............MKWMLTLCLLILAPMSWAE.TKT.......KAD...................KYEGIEITVNVNSAT   40
NTDB id 1362 HI 1008 AAC22669.1             ........MKTL..FTSVVLCGALVVSSSFAEEKAT.......XQTAQ......SVVTTQAEAQVAPAVVSDKLNINTAT   57
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ........MKKLSRYTFSLLFG.LFSATTFAEEKVT.......TSTEH......TPVVMQQ.VAATQKVNANVVNINTAT   57
consensus                                              *   ** ** ** **** *  *                                          *!*!*!*
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NTDB id 1127 NGFG RS10560 WP 003703428.1      QQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1      QQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1      QQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1      QQELEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1014 ACIAD RS13855 WP 004924483.1     AKELQQ.LHGVGEKKALAIVEYRQQYGKFASVDDLQKVKGIGPKFIEKNRDRLAI..................   128
NTDB id 1159 ABDM36 RS04765 WP 001166105.1    AEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL..................   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1   AEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL..................   103
NTDB id 115446 VPUCM RS05150 WP 021449594.1  AEEIATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL..................    94
NTDB id 1294 VP RS04495 WP 021449594.1       AEEIATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL..................    94
NTDB id 1395 DSB67 RS04445 WP 010649155.1    AQEIATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL..................    95
NTDB id 1362 HI 1008 AAC22669.1             ASEIQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF..................   112
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE.................   113
consensus                                     * !*   ! !*! *!! *!**!!  *! !*****!  !*!!! *** ********
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X non conserved
X similar
X ≥ 50% conserved