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 1020538 QRP18 RS04985 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 1020538 QRP18 RS04985 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 $\geq$ 50% conserved