logo

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NTDB id 1127 NGFG RS10560 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSC.....................AFSLAA    52
NTDB id 1126 NGFG RS09375 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSC.....................AFSLAA    52
NTDB id 1125 NGFG RS07215 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSC.....................AFSLAA    52
NTDB id 1108 NGFG RS06525 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKMFVLFCMLFSC.....................AFSLAA    52
NTDB id 1272 LPP RS04360 WP 010946546.1      ........................MKARLI...AILLSFIVIPFSSQAENEPQKTPA.....KQVLVKGK    38
NTDB id 1249 GCO85 RS04230 WP 010946546.1    ........................MKARLI...AILLSFIVIPFSSQAENEPQKTPA.....KQVLVKGK    38
NTDB id 345009 EWA64 RS13420 WP 061895163.1  ........................MKRILVSIALVLSCLF..SFTANANEAPAKVDKH.....EGIEIT    38
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ......................MQIKTKIVTL.FLSLCLPTLPLLANAEETAPAAQVE.....EGIVIT    41
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ......................MQIKTKIVTL.FLSLCLPTLPLLANAEETAPAAQVE.....EGIVIT    41
NTDB id 1294 VP RS04495 WP 021449594.1       ..........................MKW.ILTLCLVVFAPLSLAA..DTKADKY.....EGIEIT    32
NTDB id 1395 DSB67 RS04445 WP 010649155.1    ..........................MKW.MLTLCLLILAPMSWAE.TKTKADKY.....EGIEIT    33
NTDB id 1362 HI 1008 AAC22669.1              ........................MKTL..FTSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDK    50
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ........................MKKLSRYTFSLLFG.LFSATTFAEEKVTTSTEHTPVVMQQ.VAATQKVNANV    50
consensus                                                ** * *    ******      *                 *
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NTDB id 1127 NGFG RS10560 WP 003703428.1     VNINAASQQELE.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1     VNINAASQQELE.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1     VNINAASQQELE.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1     VNINAASQQELE.ALPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1272 LPP RS04360 WP 010946546.1      INLNTADVYSLTGSFKGIGKKRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN.............   105
NTDB id 1249 GCO85 RS04230 WP 010946546.1    INLNTADVYSLTGSFKGIGKKRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN.............   105
NTDB id 345009 EWA64 RS13420 WP 061895163.1  VNINSASAEETAALLKGIGIKKAQQIVADREANGPFQSAEDLARVKGIGPSTVAKNQERIKLK.................   101
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   VNINTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.................   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1  VNINTASAEELATLLKGIGLKKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL.................   103
NTDB id 1294 VP RS04495 WP 021449594.1       VNINTASAEETATMLKGIGEKKAQSIVDYRTEHGPFKTAADLTNVKGIGEATIKKNEDRILL.................    94
NTDB id 1395 DSB67 RS04445 WP 010649155.1    VNVNSATAQEIATLLIGIGEKKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL.................    95
NTDB id 1362 HI 1008 AAC22669.1              LNINTATASEIQKSLTGIGAKKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF.................   112
NTDB id 1382 A4U84 RS07375 WP 062924234.1    VNINTATASEIQDKLVGIGAKKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE.................   113
consensus                                    *!*!*!*  **   * !!! **! *!***!  ** !*****!  !*!!! * * ****** *
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X non conserved

X similar

X ≥ 50% conserved