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logo

NTDB id 1127 NGFG RS10560 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM......FVLFC..MLFSCAFSLAAVNINAASQQEL.EALPGIGP    71
NTDB id 1126 NGFG RS09375 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM......FVLFC..MLFSCAFSLAAVNINAASQQEL.EALPGIGP    71
NTDB id 1125 NGFG RS07215 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM......FVLFC..MLFSCAFSLAAVNINAASQQEL.EALPGIGP    71
NTDB id 1108 NGFG RS06525 WP 003703428.1     MSRGGATPYRFLLIRHIVARCGLLFATLKGKTMKKM......FVLFC..MLFSCAFSLAAVNINAASQQEL.EALPGIGP    71
NTDB id 1272 LPP RS04360 WP 010946546.1      ........MKARLIAIL..LSFIVIPFSSQAENE.PQK...........TPAKQVLVKGKINLNTADVYSLTGSFKGIGK    58
NTDB id 1249 GCO85 RS04230 WP 010946546.1    ........MKARLIAIL..LSFIVIPFSSQAENE.PQK...........TPAKQVLVKGKINLNTADVYSLTGSFKGIGK    58
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   ......MQIKTKIVTLFLSLCLPTLPLLANAEETAPAA.............QVEEGIVITVNINTASAEELATLLKGIGL    61
NTDB id 1150 A1552VC RS08580 WP 001166105.1  ......MQIKTKIVTLFLSLCLPTLPLLANAEETAPAA.............QVEEGIVITVNINTASAEELATLLKGIGL    61
NTDB id 1395 DSB67 RS04445 WP 010649155.1    ............MKWMLTLCLLILAPMSWAE.TKTKA.............DKYEGIEITVNVNSATAQEIATLLIGIGE    53
NTDB id 421976 G3M63 RS13920 WP 150301204.1 ........MKAI.......YLAVLFVLSSWFALP.........VVAE..TSAAP.ASVISVNINNASAAEIAETLQGVGV    53
NTDB id 1382 A4U84 RS07375 WP 062924234.1    ........MKKLSRYTFSLLFG.LFSATTFAEEKVTTSTEHTPVVMQQ.VAATQKVNANVVNINTATASEIQDKLVGIGA    70
NTDB id 1362 HI 1008 AAC22669.1              ........MKTL..FTSVVLCGALVVSSSFAEEKATXQTAQSVVTTQAEAQVAPAVVSDKLNINTATASEIQKSLTGIGA    70
consensus                                             ** ***  ** *****   *  **           *             *  *!*!*!*  **   * !*!
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logo

NTDB id 1127 NGFG RS10560 WP 003703428.1     AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1126 NGFG RS09375 WP 003703428.1     AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1125 NGFG RS07215 WP 003703428.1     AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1108 NGFG RS06525 WP 003703428.1     AKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLAKLKDQASVGAPAPKGPAKPVLPAVKK   131
NTDB id 1272 LPP RS04360 WP 010946546.1      KRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN.............   105
NTDB id 1249 GCO85 RS04230 WP 010946546.1    KRAEAIIAYRENHQGFKSLEELAEVKGIGQRFVDKNREKLKEIFVIN.............   105
NTDB id 1159 ABDM36 RS04765 WP 001166105.1   KKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...................   103
NTDB id 1150 A1552VC RS08580 WP 001166105.1  KKAQAIVDYREANGPFTHIDDLTNVKGIGEATVRNNAARILL...................   103
NTDB id 1395 DSB67 RS04445 WP 010649155.1    KKAQDIVEYRNEHGPFKTAADLTKVKGIGEATVRKNEDRILL...................    95
NTDB id 421976 G3M63 RS13920 WP 150301204.1 AKADAIVAYREANGSFASVESLSEVSGIGAATVEKNRARIALQ..................    96
NTDB id 1382 A4U84 RS07375 WP 062924234.1    KKAQAIVEYREKNGKFLNVEQLTEVSGIGKATLEKNRDCIVIE..................   113
NTDB id 1362 HI 1008 AAC22669.1              KKAEAIVQYREKHGNFXNAEQLLEVQGIGKATLEKNRDRIIF..................   112
consensus                                    **! *!**!!* ** !*****!  !*!!! * * ****** *
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X non conserved

X similar

X ≥ 50% conserved