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logo
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NTDB id 108 BSU 35470 NP 391427.1              MNVPVEKNSSFSKELQQTLRSRHLLRTELSFSDEMIEW..HIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHSSG   78
NTDB id 615 LCA RS02545 WP 011374200.1         ............MGQQVIACGRQFTAAQLADTQNN.....NYSLPQIKRRPAFLRVKHRLVCQRCQQV.VPPQT...CLP   59
NTDB id 593 KW2 RS05130 WP 021037147.1         ...........MNLILENLLGHLLLEKDISAFNNFTDQVNNENKL.IKIGAMTSVNANKVRCNRCGTIHIKTNV...KLP   65
NTDB id 1159373 QOR54 RS02050 WP 000432961.1   ...............MENYLGRLWTKAQLSEQ.........LRKIAISLPS.FIKKGSDYICTRCSSS.VAKNC...QLP   51
NTDB id 277 KZH43 RS10090 WP 000867601.1       ............MKVNLDYLGRLFTENELTEE.........ERQLAEKLPA.MRKEKGKLFCQRCNST.ILEEW...YLP   54
NTDB id 236 SPD RS10765 WP 000867601.1         ............MKVNLDYLGRLFTENELTEE.........ERQLAEKLPA.MRKEKGKLFCQRCNST.ILEEW...YLP   54
NTDB id 202 SPR RS10250 WP 000867601.1         ............MKVNLDYLGRLFTENELTEE.........ERQLAEKLPA.MRKEKGKLFCQRCNST.ILEEW...YLP   54
NTDB id 167 SP RS11275 WP 000867616.1          ............MKVNLDYLGRLFTENELTEE.........ERQLAEKLPA.MRKEKGKLFCQRCNST.ILEEW...YLP   54
NTDB id 507 SM12261 RS09240 WP 000867722.1     ............MKVNPNYLGRLFTENELTEE.........ERQLAEKLPA.MRKEKGKLFCQRCNST.ILEEW...YLP   54
NTDB id 535 SMSK321 RS10515 WP 000867726.1     ............MKVNPNYLGRLFTENELTKE.........ERQLAEKLPA.MRKEKGKLFCQRCDSA.ILDEW...YLP   54
consensus                                                  *****  **************          **********  *********!*!!***  *****    ***
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NTDB id 108 BSU 35470 NP 391427.1              KNKLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSSGQQKAANVLIEAISKKEELLIWAVCGAGKTEMLFP   158
NTDB id 615 LCA RS02545 WP 011374200.1         DGRHYCAQCLLFGRLVEGDWLYTVPECHLFENNIPKLTWEGQLTPHQEKAAQAVVEVVQTQKRHVLTAVTGAGKTEMLFQ   139
NTDB id 593 KW2 RS05130 WP 021037147.1         IGAFFCPTCLELGRVRSDEYFYHLPQQDFPEKT..YLRWTGKLTENQEKISDALCQQITNNQKLLVQAVTGAGKTEMIYQ   143
NTDB id 1159373 QOR54 RS02050 WP 000432961.1   TGNYYCRECIVFGRVTSNENLYYFPQKTFSKTN..SLKWKGELTPYQNEVSEELLKGISSKENLLVHAVTGAGKTEMIYH   129
NTDB id 277 KZH43 RS10090 WP 000867601.1       IGAYYCRECLLMKRVRSDQTLYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQ   132
NTDB id 236 SPD RS10765 WP 000867601.1         IGAYYCRECLLMKRVRSDQTLYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQ   132
NTDB id 202 SPR RS10250 WP 000867601.1         IGAYYCRECLLMKRVRSDQTLYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQ   132
NTDB id 167 SP RS11275 WP 000867616.1          IGAYYCRECLLMKRVRSDQTLYYFPQEDFPKQD..VLKWRGQLTPFQEKVSEGLLQVVDKQKPTLVHAVTGAGKTEMIYQ   132
NTDB id 507 SM12261 RS09240 WP 000867722.1     IGTYYCRECLLMKRVRSDQSLYYFPQEDFPKQD..VLKWRGQLTPFQDKVSQGLLQAVDKQEPSLVHAVTGAGKTEMIYQ   132
NTDB id 535 SMSK321 RS10515 WP 000867726.1     IGAYYCRECLLMKRVRSDQVLYYFPQEDFAKQD..ILKWRGQLTPFQEKVSEGLIQAVDKQEPILVHAVTGAGKTEMIYQ   132
consensus                                      *****!**!****!******!***********  *!*!*!*!***!*******************!!*!!!!!!!***
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logo
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NTDB id 108 BSU 35470 NP 391427.1              GIESALNQGLRVCIATPRTDVVLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAF   238
NTDB id 615 LCA RS02545 WP 011374200.1         GILVALQKGQRVCLAAPRVAVCLELYPRLQAAFATTSIMLMHGEQTEPYRYTQLVLCTTHQLLKFYHAFDTVIVDEVDAF   219
NTDB id 593 KW2 RS05130 WP 021037147.1         LIEQILSHGGSVGLASPRIDVCIELHQRLSRDFTC.QIPLLYHE.GDSYFRSPLVVMTSHQLLRFKEAFDLLIIDEVDAF   221
NTDB id 1159373 QOR54 RS02050 WP 000432961.1   SVAKVIDTGGSVCIASPRIDVCLELYKRLSNDFRC.AITLMHGE.SPSYQRSPLTIATTHQLLKFYHAFDLLIVDEVDAF   207
NTDB id 277 KZH43 RS10090 WP 000867601.1       VVAKVINAGGAVCLASPRIDVCLELYKRLQQDFSC.GIALLHGE.SEPYFRTPLVVATTHQLLKFYQAFDLLIVDEVDAF   210
NTDB id 236 SPD RS10765 WP 000867601.1         VVAKVINAGGAVCLASPRIDVCLELYKRLQQDFSC.GIALLHGE.SEPYFRTPLVVATTHQLLKFYQAFDLLIVDEVDAF   210
NTDB id 202 SPR RS10250 WP 000867601.1         VVAKVINAGGAVCLASPRIDVCLELYKRLQQDFSC.GIALLHGE.SEPYFRTPLVVATTHQLLKFYQAFDLLIVDEVDAF   210
NTDB id 167 SP RS11275 WP 000867616.1          VVAKVINAGGAVCLASPRIDVCLELYKRLQQDFSC.GIALLHGE.SEPYFRTPLVVATTHQLLKFYQAFDLLIVDEVDAF   210
NTDB id 507 SM12261 RS09240 WP 000867722.1     VVAKVINAGGAVCLASPRIDVCLELYKRLQKDFAC.EIALLHGE.SEPYFRTPLVVATTHQLLKFYQAFDLLIVDEVDAF   210
NTDB id 535 SMSK321 RS10515 WP 000867726.1     VVAKVINAGGAVCLASPRIDVCLELYKRLQDDFAC.EISLLYGE.SEPYFRTPLVVSTTHQLLKFYQAFDLLIVDEVDAF   210
consensus                                      ********!**!**!*!!**!**!!**!!***!** *!****** ********!***!*!!!!****!*!**!*!!!!!!
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logo
NTDB id 108 BSU 35470 NP 391427.1          PYSADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRIPARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAV   318
NTDB id 615 LCA RS02545 WP 011374200.1     PFVDNPVLATAVEQACKPQCALLYLTATPTPAIKRAIAAKQMTVSELPLRFHGHLLPEPQRHVAFNWRAQLKKGRLPKRL   299
NTDB id 593 KW2 RS05130 WP 021037147.1     PFRDNDMLYFALENAKKINGTLLYLTATSTDKLDKQIKKHEIKRLFLPRRFHGHPLVIPMFFWKKTFYK...........   290
NTDB id 1159373 QOR54 RS02050 WP 000432961.1 PYVDNPILYQGVKQALKENGTSIFLTATSTTELERKVARKELKKLHLARRFHANPLVIPEMVWVSGIQKSLQTQKLPPKL   287
NTDB id 277 KZH43 RS10090 WP 000867601.1   PYVDNPMLYHAVKNSVKENGLRIFLTATSTNELDKKVRLGELKRLNLPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKL   290
NTDB id 236 SPD RS10765 WP 000867601.1     PYVDNPMLYHAVKNSVKENGLRIFLTATSTNELDKKVRLGELKRLNLPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKL   290
NTDB id 202 SPR RS10250 WP 000867601.1     PYVDNPMLYHAVKNSVKENGLRIFLTATSTNELDKKVRLGELKRLNLPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKL   290
NTDB id 167 SP RS11275 WP 000867616.1      PYVDNPMLYHAVKNSVKENGLRIFLTATSTNELDKKVRLGELKRLNLPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKL   290
NTDB id 507 SM12261 RS09240 WP 000867722.1 PYVDNPMLYHAVKNSVKENGLRIFLTATSTDELDRKVRIGELKRLSLPRRFHGNPLIIPKPVWLSDFNHCLEKSRLSPKL   290
NTDB id 535 SMSK321 RS10515 WP 000867726.1 PYVDNPMLYHAVKNSVKENGLRIFLTATSTDELDRKVRIGELKRLSLPRRFHGNPLIIPKPVWLSDFNRYLEKKCLSPKL   290
consensus                                  !******!********!*******!*!!** ************** ***!*!***!**!***********************
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logo
NTDB id 108 BSU 35470 NP 391427.1          KRWIEFHVKEGRPVFLFVPSVSILEKAAACFKGV..HCRTASVHAEDKHRKEKVQQFRDGQLDLLITTTILERGVTVPKV   396
NTDB id 615 LCA RS02545 WP 011374200.1     ERDCQTYL.KTQQILLFVPQVRLLQPVANRLSTLLPAVRIETLHANDPDQIKKITAFRAQQIQLMVTTTILERGVTFKNV   378
NTDB id 593 KW2 RS05130 WP 021037147.1     ..KFIEQRKTGFPLLIFVAEIDFGQDFAKNLQEKFPKEKMAFVASTTKSRKTIVEAFRKKQVSILITTSILERGVTFSSI   368
NTDB id 1159373 QOR54 RS02050 WP 000432961.1 YQLINKQRQTRYPLLLFFPHISEGQVFTEILRQAFPMEKIGFVSSKSTSRLKLVQDFRDNKLSILVSTTILERGVTFPSV   367
NTDB id 277 KZH43 RS10090 WP 000867601.1   KSYIEKQRKTAYPLLIFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCV   370
NTDB id 236 SPD RS10765 WP 000867601.1     KSYIEKQRKTAYPLLIFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCV   370
NTDB id 202 SPR RS10250 WP 000867601.1     KSYIEKQRKTAYPLLIFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCV   370
NTDB id 167 SP RS11275 WP 000867616.1      KSYIEKQRKTAYPLLIFASEIKKGEQLAEILQEQFPNEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCV   370
NTDB id 507 SM12261 RS09240 WP 000867722.1 KSYIEKQRKTGYPLLIFASEIKKGEQLKEIIQEQFPNEKIGFVSSITENRLEQVQAFRDRELTILISTTILERGVTFPCV   370
NTDB id 535 SMSK321 RS10515 WP 000867726.1 KSYIEKQRKTGYPLLIFASEIKKGEQLKEILQKEFPHEKIGFVSSVTEDRLEQVQAFRDGELTILISTTILERGVTFPCV   370
consensus                                  ****************!*************************************!!*********!*!!!!!!!****
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logo
NTDB id 108 BSU 35470 NP 391427.1          QTGVLGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD   463
NTDB id 615 LCA RS02545 WP 011374200.1     AVLVLGAEHMVFNEAVLVQIAGRAGRHKDYAQNPVHFYYQDYTRAIKGACRQIKTQNQRGRRLQSTV   445
NTDB id 593 KW2 RS05130 WP 021037147.1     DVFVINSEHPNFTKSALIQMAGRVGRDSKRPTGLVSFFHSGKSLAMCQAQKEIKKMNRLGGF.....   430
NTDB id 1159373 QOR54 RS02050 WP 000432961.1 DVFVIQANHHLFTKSSLVQISGRVGRALERPEGLLYFLHDGKSKSMHQAIKEIKNMNHIGGF.....   429
NTDB id 277 KZH43 RS10090 WP 000867601.1   DVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHDGLNASIKKAIKEIQMMNKEAGL.....   432
NTDB id 236 SPD RS10765 WP 000867601.1     DVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHDGLNASIKKAIKEIQMMNKEAGL.....   432
NTDB id 202 SPR RS10250 WP 000867601.1     DVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHDGLNASIKKAIKEIQMMNKEAGL.....   432
NTDB id 167 SP RS11275 WP 000867616.1      DVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHDGLNASIKKAIKEIQMMNKEAGL.....   432
NTDB id 507 SM12261 RS09240 WP 000867722.1 DVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHDGLNTSIKKAIKEIQQMNKEAGL.....   432
NTDB id 535 SMSK321 RS10515 WP 000867726.1 DVFVVEANHRLFTKSSLIQIGGRVGRSMDRPTGDLLFFHDGLNTSIKKAIKEIQQMNKEAGL.....   432
consensus                                  ***!*******!****!*!**!!*!!****************** ****!***!* *!*****
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X non conserved

X similar

X ≥ 50% conserved