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NTDB id 1159 ABDM36 RS04765 WP 001166105.1VEEGIVITVNINTASAEELATLLKGGIGLKKAAQAVDYREANGPFTHIDDLTNVKGIGEATV	94
NTDB id 1150 A1552VC RS08580 WP 001166105.1VEEGIVITVNINTASAEELATLLKGGIGLKKAAQAVDYREANGPFTHIDDLTNVKGIGEATV	94
NTDB id 1395 DSB67 RS04445 WP 010649155.1KYEGIEITVMVNSATAQEITATLLIGIGGKKAQDIVEYRNEHGPFKTAADLTQVKGIGEATV	86
NTDB id 1362 HI 1008 AAC22669.1	QAEA.....QVAPAVVSDKLNINTATASEITQKSLTGICAKKAEATVQYREKHCNFXNAEQLLVEVQGICKATL	103
NTDB id 1382 A4U84 RS07375 WP 062924234.1	QQ.V.....AATQKVNANVVNINTATASEITQDKLVGIGAKKAAQAVVEYREKNGKFLNVEQLTEVSGIGKATL	103
NTDB id 595 KW2 RS08725 WP 021037687.1S.....S..VSNQISTTSGPKVNINKADITELQK.LTGIGQKKAQDIDDFRMKNGDFKSLDELKQVSGFGDKTL	208
NTDB id 521 SMSK321 RS04960 WP 000443804.1SSTSCKDKKLNINKASLEELKQ.VKCIGGKRAQDIDHREANGKFKSVDELKQVSGIGAKTI	206
NTDB id 493 SM12261 RS04300 WP 000387351.1SSTSKEKKVNINKASLEELKQ.VKCIGGKRAQDIDHREANGKFKSVDELKQVSGIGAKTI	206
NTDB id 263 KZH43 RS04195 WP 000387330.1SSISKEKKVNINKASLEELKQ.VKCIGGKRAQDIDHREANGKFKSVDELKQVSGIGGKTI	206
NTDB id 222 SPD RS04535 WP 000387330.1SSISKEKKVNINKASLEELKQ.VKCIGGKRAQDIDHREANGKFKSVDELKQVSGIGGKTI	206
NTDB id 188 SPR RS04295 WP 000387330.1SSISKEKKVNINKASLEELKQ.VKCIGGKRAQDIDHREANGKFKSVDELKQVSGIGGKTI	206
NTDB id 153 SP RS04730 WP 000387344.1SSTSKEKKVNINKASLEELKQ.VKCIGGKRAQDIDHREANGKFKSVDELKQVSGIGGKTI	206
NTDB id 1069 ABD1 RS03060 WP 043960841.1	KLNISKSSHSNSSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGGIGKKAQAVVEYRQKNGSFKNIDELKQVKGIGPAIF	127
NTDB id 1041 HON27 RS14740 WP 005124713.1	KLNITKSSHSNGSKISQTKNFNDSTGQIHLNQANVNEFQQ.LKGGIGKKAQAVVEYRQKNGSFKNIDELKQVKGIGPAIF	127
NTDB id 105 BSU 25590 NP 390437.1	QGGG.....GSV....QSDGGKALVNINTATLEELQG.TSCVGPSSKAEATVAYREENGFRFQITIEDITKQVSGIGESF	195
NTDB id 36 MW RS08230 WP 001793467.1	QIGVNSV.....HV....KNGNTNNTKVNINTASVSELSM.VPGVGGAKANAIVEYRNQQGAFQETDDLKQVKGIFGSKTF	219
NTDB id 20 SA RS08005 WP 001802022.1	QIEVNSV.....HV....KNGNTNNTKVNINTASVSELSM.VPGVGGAKANAIVEYRNQQGAFQETDDLKQVKGIFGSKTF	219
NTDB id 1014 ACIAD RS13855 WP 004924483.1	RLNSSASHHSLSKP...TQQSGTGKVNINSATAKELQQ.LHCVGEEKALAAIVEYRQQYQKFAVDDLIQVKGIGPKFI	119
NTDB id 1158257 QOR53 RS00050 WP 002853093.1ALTAFLFGAVNINTATLKEKLS.LNGICEAKAKAILEYRKE.ANFTSDDLKQVKGIGDKLF	69
NTDB id 1242 Cj0011c YP 002343483.1ALTAFLFGAVNINTATLKEKLS.LNGICEAKAKAILEYRKE.ANFTSDDLKQVKGIGDKLF	69
NTDB id 1127 NGFG RS10560 WP 003703428.1FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKATAEYRAQNGAFKSVDDLKQVKGIGPAVL	104
NTDB id 1126 NGFG RS09375 WP 003703428.1FSCAFSLAAVNINAASQQELEA.LPGIGPAKAKATAEYRAQNGAFKSVDDLKQVKGIGPAVL	104
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consensus	***** ***	

 non conserved
 similar
 $\geq 50\%$ conserved