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logo
NTDB id 1142302 DQN71 RS02500 WP 015605268.1  MTEKLVEIKDLEISFGEGSKKFVAVKNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSAGDILYEGKKINGKQSKA   80
NTDB id 479 HSISS4 RS06980 WP 021143812.1     MPEKLVEVKDVEISFGEGSKKFVAVHNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGEIIFDGKKINGHLSHS   80
NTDB id 328 STU RS16110 WP 002951426.1        MPEKLVEVKDVEISFGEGRKKFVAVHNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGEIIFDGKKINGYLSHS   80
NTDB id 296 STER RS06910 WP 011681415.1       MPEKLVEVKNVEISFGEGRKKFVAVHNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGEIIFDGKKINGYLSHS   80
consensus                                     !*!!!!!*!**!!!!!!! !!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!*!***!!!!!!**!**

logo
NTDB id 1142302 DQN71 RS02500 WP 015605268.1  DEAELIRKIQMIFQDPAASLNERATVDYIISEGLYNYHLFENEEDRQEKVKNIIREVGLLAEHLTRYPHEFSGGQRQRIG  160
NTDB id 479 HSISS4 RS06980 WP 021143812.1     EKNDLIRRIQMIFQDPAASLNERATVDYILSEGLHNFHLYKDEEDRKAKIKKIIKEVGLLEEHLTRYPHEFSGGQRQRIG  160
NTDB id 328 STU RS16110 WP 002951426.1        EKNDLIRRIQMIFQDPAASLNERATVDYILSEGLYNFHLYKDEEERKAKIKEIIKEVGLLEEHLTRYPHEFSGGQRQRIG  160
NTDB id 296 STER RS06910 WP 011681415.1       EKNDLIRRIQMIFQDPAASLNERATVDYILSEGLYNFHLYKDEEERKAKIKEIIKEVGLLEEHLTRYPHEFSGGQRQRIG  160
consensus                                     ****!!!!*!!!!!!!!!!!!!!!!!!!!!!*!!!!!*!*!!***!!*!**!*!*!!*!!!!!*!!!!!!!!!!!!!!!!!!!

logo
NTDB id 1142302 DQN71 RS02500 WP 015605268.1  IARALVMEPDFVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGVIVEVAETEELFN  240
NTDB id 479 HSISS4 RS06980 WP 021143812.1     IARSLVMQPDLVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGTIVEVAETEELYN  240
NTDB id 328 STU RS16110 WP 002951426.1        IARSLVMQPDLVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGTIVEVAETEELYN  240
NTDB id 296 STER RS06910 WP 011681415.1       IARSLVMQPDLVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGTIVEVAETEELYN  240
consensus                                     !!!*!!!*!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!

logo
NTDB id 1142302 DQN71 RS02500 WP 015605268.1  NPIHPYTQSLLSAVPIPDPILERKKVLKVYDPEQHDYSVDQPSMIEIRPGHYVWANKAEADRYKKEYK.  308
NTDB id 479 HSISS4 RS06980 WP 021143812.1     NPIHPYTKSLLSAVPIPDPILERKKVLKVYDPNQHDYSVDKPEMVEVRPGHFVWGNKAEIETYRKEQSK  309
NTDB id 328 STU RS16110 WP 002951426.1        NPIHPYTKSLLSAVPIPDPILERKKVLKVYDPNQHDYSVDKPEMVEVRPGHFVWGNKTEIETYRKEQSK  309
NTDB id 296 STER RS06910 WP 011681415.1       NPIHPYTKSLLSAVPIPDPILERKKVLKVYDPNQHDYSVDKPEMVEVRPGHFVWGNKTEIETYRKEQSK  309
consensus                                     !!!!!!!*!!!!!!!!!!!!!!!!!!!
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