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logo

NTDB id 1043698 ACD268 RS02060 WP 001291293.1  MSEKLVEIKDLEISFGEGSKKFVAVKNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGDIIFDGQKINGKKSRE   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 1043698 ACD268 RS02060 WP 001291293.1  QAAELIRRIQMIFQDPAASLNERATVDYIISEGLYNHRLFKDEEERKEKVQNIIREVGLLAEHLTRYPHEFSGGQRQRIG  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                                      ****!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!*!**!*!!!!*!!*!***!!*!!!!!*!!!!!!!!!!!!!!!!!!!

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NTDB id 1043698 ACD268 RS02060 WP 001291293.1  IARALVMQPDFVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGVIVEVAETEELFN  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 1043698 ACD268 RS02060 WP 001291293.1  NPIHPYTQALLSAVPIPDPILERKKVLKVYDPSQHDYETDKPSMVEIRPGHYVWANQAELARYQKGLN.   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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