logo

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NTDB id 51121 SPAF RS02805 WP 041826241.1  MTEKLVEIKDLEISFGEGSKKFVAVKNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSAGEIIYDGRKINGKNSHSEKSEL   85
NTDB id 479 HSISS4 RS06980 WP 021143812.1  MPEKLVEVKDVEISFGEGSKKFVAVHNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGEIIFDGKKINGHLSHSEKNDL   85
NTDB id 328 STU RS16110 WP 002951426.1      MPEKLVEVKDVEISFGEGRKKFVAVHNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGEIIFDGKKINGYLSHSEKNDL   85
NTDB id 296 STER RS06910 WP 011681415.1     MPEKLVEVKNVEISFGEGRKKFVAVHNANFFINKGETFSLVGESGSGKTTIGRAIIGLNDTSNGEIIFDGKKINGYLSHSEKNDL   85
consensus                                  !*!!!!!*!**!!!!!!! !!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!*!!*!!!!**!!!!!**!
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logo

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NTDB id 51121 SPAF RS02805 WP 041826241.1  IRKIQMIFQDPAASLNERATVDYIISEGLINHHLFNSEEERQEKVKNIMHEVGLLAEHLTRYPHEFSGGQRQRIGIARALVMEPE   170
NTDB id 479 HSISS4 RS06980 WP 021143812.1  IRRIQMIFQDPAASLNERATVDYILSEGLHNFHLYKDEEDRKAKIKKIIKEVGLLEEHLTRYPHEFSGGQRQRIGIARSLVMQPD   170
NTDB id 328 STU RS16110 WP 002951426.1      IRRIQMIFQDPAASLNERATVDYILSEGLYNFHLYKDEEERKAKIKEIIKEVGLLEEHLTRYPHEFSGGQRQRIGIARSLVMQPD   170
NTDB id 296 STER RS06910 WP 011681415.1     IRRIQMIFQDPAASLNERATVDYILSEGLYNFHLYKDEEERKAKIKEIIKEVGLLEEHLTRYPHEFSGGQRQRIGIARSLVMQPD   170
consensus                                  !!*!!!!!!!!!!!!!!!!!!!!!*!!!!*!*!!***!!*!**!*!*!**!!!!!*!!!!!!!!!!!!!!!!!!!!!!*!!!*!*
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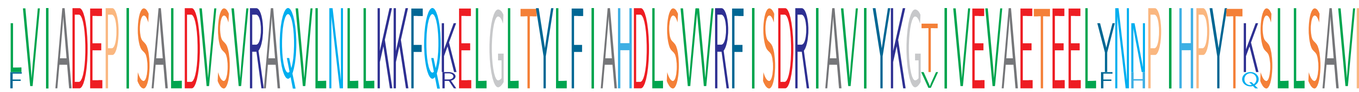
logo



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NTDB id 51121 SPAF RS02805 WP 041826241.1  FVIADEPISALDVSVRAQVLNLLKKFQRELGLTYLFIAHDLSVVRFISDRIAVIYKGVIVEVAETEELFNHPIHPYTQSLLSAVP   255
NTDB id 479 HSISS4 RS06980 WP 021143812.1  LVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGTIVEVAETEELYNNPIHPYTKSLLSAVP   255
NTDB id 328 STU RS16110 WP 002951426.1      LVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGTIVEVAETEELYNNPIHPYTKSLLSAVP   255
NTDB id 296 STER RS06910 WP 011681415.1     LVIADEPISALDVSVRAQVLNLLKKFQKELGLTYLFIAHDLSVVRFISDRIAVIYKGTIVEVAETEELYNNPIHPYTKSLLSAVP   255
consensus                                  *!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!*!!!!!!!!!!*!*!!!!!!*!!!!!!!
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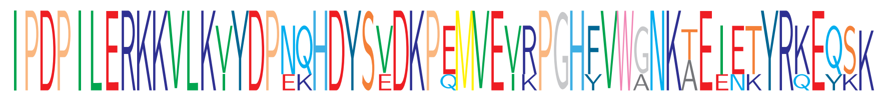
logo



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NTDB id 479 HSISS4 RS06980 WP 021143812.1  IPDPILERKKVLKVYDPNQHDYSVDKPEMVEVRPGHFVWGNKAEIETYRKEQSK   309
NTDB id 328 STU RS16110 WP 002951426.1      IPDPILERKKVLKVYDPNQHDYSVDKPEMVEVRPGHFVWGNKTEIETYRKEQSK   309
NTDB id 296 STER RS06910 WP 011681415.1     IPDPILERKKVLKVYDPNQHDYSVDKPEMVEVRPGHFVWGNKTEIETYRKEQSK   309
consensus                                  !!!!!!!!!!!!!*!!!**!!!!*!!!*!!!**!!!*!!*!! !***!!*!***
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X non conserved

X similar

X ≥ 50% conserved