

logo

NTDB id 256714 CV702 RS08005 WP 017864798.1  
NTDB id 607 V4T04 RS01905 WP 010906048.1  
consensus

MKKLFTFNKKKKEVEESSAKRSIMLRWAFANTVFCFITFTLFATLTYQLTISSFIKEEQQLLTRSMDSVEEVLEKADAPL  
MKKLFTFNKKKKEVEESSAKRSIMLRWAFANTVFCFITFTLFATLTYQLTISSFIKEEQQLLTRSMDSVEEVLEKADAPL  
!!

80  
80

logo

NTDB id 256714 CV702 RS08005 WP 017864798.1  
NTDB id 607 V4T04 RS01905 WP 010906048.1  
consensus

NSSNLNTYIEATSKIQNGESEGMSLGSIIGTRKAFYIYDLNHKLLYSTNRHTFGFQNNANEMKEIRGENPGYLVQRKII  
NSSNLNTYIEATSKIQNGESEGMSLGSIIGTRKAFYIYDLNHKLLYSTNRHTFGFQNNANEMKEIRGENPGYLVQRKII  
NSSNLNTYIEATSKIQNGESEGMSLGSIIGTRKAFYIYDLNHKLLYSTNRHTFGFQNNANEMKEIRGENPGYLVQRKII  
!!

160  
160

logo

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consensus

SKSTGQVVGYLQAFYDTTTTYHRISNLLLTVLLILEIVALIVAQLIGYFMANYFMKPLEKLYQGMQEMANDPTNDFEPIEI  
SKSTGQVVGYLQAFYDTTTTYHRISNLLLTVLLILEIVALIVAQLIGYFMANYFMKPLEKLYQGMQEMANDPTNDFEPIEI  
SKSTGQVVGYLQAFYDTTTTYHRISNLLLIIVLLILEIVALIVAQLIGYFMANYFMKPLEKLYQGMQEMANDPTNDFEPIEI  
!!

240  
240

logo

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consensus

QSGDEIEELAHVYNDMMLKMKAYLEQQNRFVSDVSHELRTPLAVLDGHINLLNRWGKNDPEVLDES LQASLDEVDRM KKM  
QSGDEIEELAHVYNDMMLKMKAYLEQQNRFVSDVSHELRTPLAVLDGHINLLNRWGKNDPEVLDES LQASLDEVDRM KKM  
QSGDEIEELAHVYNDMMLKMKAYLEQQNRFVSDVSHELRTPLAVLDGHINLLNRWGKNDPEVLDES LQASLDEVDRM KKM  
!!

320  
320

logo

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consensus

LEEMLALARLENVDLSSEELDCDVGKVCNRALKNFQLLHDDFEIVLDNRLIYPTHARISENHFEQGLRILLDNAAKYSPD  
LEEMLALARLENVDLSSEELDCDVGKVCNRALKNFQLLHDDFEIVLDNRLIYPTHARISENHFEQGLRILLDNAAKYSPD  
LEEMLALARLENVDLSSEELDCDVGKVCNRALKNFQLLHDDFEIVLDNRLIYPTHARISENHFEQGLRILLDNAAKYSPD  
!!

400  
400

logo

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consensus

DRKEIVITVSEDEQFVITSVSDKIGIGISEEDINHLFERFFRADKARNREIGGTGLGLSILARLAENYQGEIEVNSELGLG  
DRKEIVITVSEDEQFVITSVSDKIGIGISEEDINHLFERFFRADKARNREIGGTGLGLSILARLAENYQGEIEVNSELGLG  
DRKEIVITVSEDEQFVITSVSDKIGIGISEEDINHLFERFFRADKARNREIGGTGLGLSILARLAENYQGEIEVNSELGLG  
!!

480  
480

logo

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NTDB id 607 V4T04 RS01905 WP 010906048.1  
consensus

STFTLKFPKIK  
STFTLKFPKIK  
STFTLKFPKIK  
!!!!!!!!!!!!!!!!

491  
491

 non conserved  
 similar  
  $\geq 50\%$  conserved