

logo

NTDB id 1088322 ACLII2 RS15940 WP 003245494.1  
NTDB id 94 BSU 11300 NP 389012.2  
consensus

MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDITNDLVWGTGKYKGLLNISQSKYNVDVYKEGVKTEEDIINAIED  
MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDITNDLVWGTGKYKGLLNISQSKYNVDVYKEGVKTEEDIINAIED  
!!

80  
80

logo

NTDB id 1088322 ACLII2 RS15940 WP 003245494.1  
NTDB id 94 BSU 11300 NP 389012.2  
consensus

FHKRGVNLLYGHGSEYAEVFNLVGEDYPMDFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQ  
FHKRGVNLLYGHGSEYAEVFNLVGEDYPMDFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQ  
FHKRGVNLLYGHGSEYAEVFNLVGEDYPMDFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASFTWQ  
!!

160  
160

logo

NTDB id 1088322 ACLII2 RS15940 WP 003245494.1  
NTDB id 94 BSU 11300 NP 389012.2  
consensus

PEVDGFIKGAKYENPDIENVTKYTDHWDDDDTTAVKLYQKMKNEGADVVPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL  
PEVDGFIKGAKYENPDIENVTKYTDHWDDDDTTAVKLYQKMKNEGADVVPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL  
PEVDGFIKGAKYENPDIENVTKYTDHWDDDDTTAVKLYQKMKNEGADVVPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL  
!!

240  
240

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GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYDLNTGVVEMGTFSPPLVDQDFQQRIAKLIKTYNKTGELPKNE  
GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYDLNTGVVEMGTFSPPLVDQDFQQRIAKLIKTYNKTGELPKNE  
GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYDLNTGVVEMGTFSPPLVDQDFQQRIAKLIKTYNKTGELPKNE  
!!

317  
317

non conserved  
 similar  
 ≥ 50% conserved