

logo

NTDB id 398126 GII79 RS06040 WP 003245494.1
NTDB id 94 BSU 11300 NP 389012.2
consensus

MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNISQSKYNVDVYKKEGVKTEEDIINAIED
MITRLVMIFSVLLLLSGCGQTPFKGKIEKVGMLFPDTINDLVWGTKGYKGLLNISQSKYNVDVYKKEGVKTEEDIINAIED
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

80
80

logo

NTDB id 398126 GII79 RS06040 WP 003245494.1
NTDB id 94 BSU 11300 NP 389012.2
consensus

FHKRGVNLLYGHGSEYAEVFNLVGEDYPMDFVSNAAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASTWQ
FHKRGVNLLYGHGSEYAEVFNLVGEDYPMDFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASTWQ
FHKRGVNLLYGHGSEYAEVFNLVGEDYPMDFVISNAKAKADNVTSVHFSGEAMGFFGGMTAAHMSKTNQVGVIASTWQ
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160
160

logo

NTDB id 398126 GII79 RS06040 WP 003245494.1
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consensus

PEVDGFIKGAKYENPDI EVNTKYTDHWDDDTTAVKLYQKMKNEGADVVPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL
PEVDGFIKGAKYENPDI EVNTKYTDHWDDDTTAVKLYQKMKNEGADVVPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL
PEVDGFIKGAKYENPDI EVNTKYTDHWDDDTTAVKLYQKMKNEGADVVPAGDGYNVPVIQQIKKDGLYAIGYVTDQSDL
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

240
240

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consensus

GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYDNLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE
GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYDNLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE
GENTVLTSTVQNVDKAYEIIAEQFNKGTLEGGDHYDNLNTGVVEMGTFSPLVDQDFQQRIAKLIKTYNKTGELPKNE
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

317
317

non conserved
 similar
 ≥ 50% conserved