

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWQSENIATLTEIRRHFTLKGSGRMIGAKSSA  
MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWQSENIATLTEIRRHFTLKGSGRMIGAKSSA  
MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWQSENIATLTEIRRHFTLKGSGRMIGAKSSA  
!!

80  
80

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQLQQQSSLEPALEELLQLS  
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQLQQQSSLEPALEELLQLS  
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQLQQQSSLEPALEELLQLS  
!!

160  
160

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

NTLTAETVTGLELGYIEQDSLAEPIIQATHTTEIEINLDELALFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
NTLTAETVTGLELGYIEQDSLAEPIIQATHTTEIEINLDELALFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
HTLTAETVTGLELDYIEQDSLAEPIIQATHTTEIEINLDELALFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
\*!!

240  
240

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQLEARLQFNQSWDAYVEQ  
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQLEARLQFNQSWDAYVEQ  
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQLEARLQFNQSWDAYVEQ  
!!

320  
320

logo

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consensus

HGDRTPPLMPHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLQHTHSQAMGLHEYTSQLKESYEV  
HGDRTPPLMPHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLQHTHSQAMGLHEYTSQLKESYEV  
HGDRTPPLMPHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLQHTHSQAMGLHEYTSQLKESYEV  
!!

400  
400

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

LLDKPALLQSDYIFEIYQKAHQQLQLFDALAAGQRVGVVVKQHQSI EELKLYTQYTPDISNDLPQQDSTSFEPISNIEP  
LLDKPALLQSDYIFEIYQKAHQQLQLFDALAAGQRVGVVVKQHQSI EELKLYTQYTPDISNDLPQQDSTSFEPISNIEP  
LLDKPALLQSDYIFEIYQKAHQQLQLFDALAAGQRVGVVVKQHQSI EELKLYTQYTPDISNDLPQQDSTSFEPISNIEP  
!!

480  
480

logo

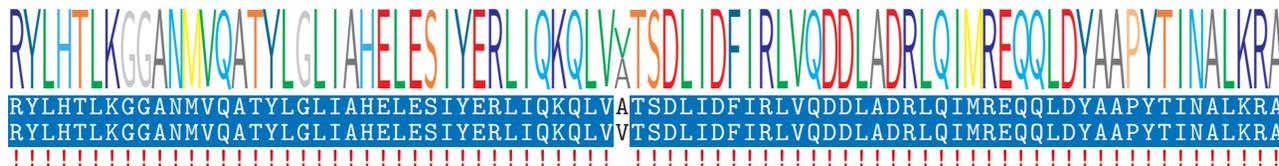
NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

EPEWATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM  
EPEVATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM  
EPEVATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM  
!!

560  
560

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



640  
640

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



720  
720

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



800  
800

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



880  
880

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



960  
960

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1040  
1040

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1120  
1120

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1200  
1200

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1280  
1280

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
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consensus



1360  
1360

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
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consensus



1440  
1440

logo

NTDB id 230029 AB895 RS03975 WP 032059036.1  
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consensus



1506  
1506

non conserved  
 similar  
 ≥ 50% conserved