

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

MKEILKTLTEAMLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA  
MKEILKTLTEAMLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA  
MKEILKTLTEAMLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA  
!!

80  
80

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

ELAMTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDNFKQKRAHTLDFRPLILLGQQLKQQQSLEPALEELLQLS  
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDNFKQKRAHTLDFRPLILLGQQLKQQQSLEPALEELLQLS  
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDNFKQKRAHTLDFRPLILLGQQLKQQQSLEPALEELLQLS  
!!

160  
160

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
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consensus

HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
\*!!

240  
240

logo

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consensus

GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ  
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ  
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ  
!!

320  
320

logo

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consensus

HGDRTPPLMPHGLVSQLQLDVSSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMGLHEYTSQLKESYEV  
HGDRTPPLMPHGLVSQLQLDVSSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMGLHEYTSQLKESYEV  
HGDRTPPLMPHGLVSQLQLDVSSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMGLHEYTSQLKESYEV  
!!

400  
400

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
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consensus

LLDKPALLQSDYIFEIYQKAHQQLQLFDALAAGQRVGVVVKQHQSILEELKLYTQYTPDLSNDLPQQDSTSFEPISNIEP  
LLDKPALLQSDYIFEIYQKAHQQLQLFDALAAGQRVGVVVKQHQSILEELKLYTQYTPDLSNDLPQQDSTSFEPISNIEP  
LLDKPALLQSDYIFEIYQKAHQQLQLFDALAAGQRVGVVVKQHQSILEELKLYTQYTPDLSNDLPQQDSTSFEPISNIEP  
!!

480  
480

logo

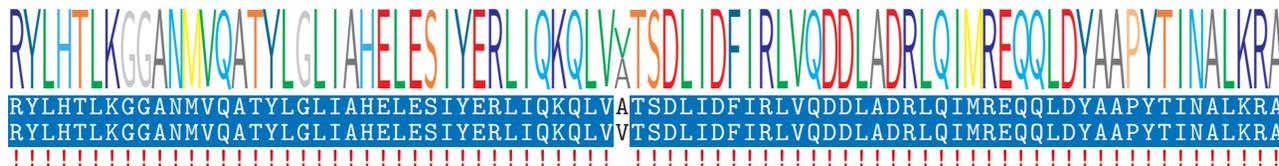
NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

EPEWATVENFSDSADWALGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDITDLNIWVGEQENFAALNNLM  
EPEVATVENFSDSADWALGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDITDLNIWVGEQENFAALNNLM  
EPEVATVENFSDSADWALGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDITDLNIWVGEQENFAALNNLM  
!!

560  
560

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



640  
640

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



720  
720

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



800  
800

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



880  
880

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



960  
960

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1040  
1040

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1120  
1120

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1200  
1200

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1280  
1280

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
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consensus



1360  
1360

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
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consensus



1440  
1440

logo

NTDB id 208119 BS103 RS14465 WP 000658894.1  
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consensus



1506  
1506

non conserved  
 similar  
 ≥ 50% conserved