

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

MKEILKTLTEAMLPEDSYSEQDAEFLEIFEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA  
MKEILKTLTEAMLPEDSYSEQDAEFLEIFEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA  
MKEILKTLTEAMLPEDSYSEQDAEFLEIFEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA  
!!

80  
80

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFKQKRAHTLDFRPLILLGQQQLQQQQSLEPALEEELLQLS  
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFKQKRAHTLDFRPLILLGQQQLQQQQSLEPALEEELLQLS  
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFKQKRAHTLDFRPLILLGQQQLQQQQSLEPALEEELLQLS  
!!

160  
160

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

NTLTAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
NTLTAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
HTLTAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR  
\*!!

240  
240

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ  
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ  
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ  
!!

320  
320

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

HGDRTPPLMPSHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV  
HGDRTPPLMPSHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV  
HGDRTPPLMPSHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV  
!!

400  
400

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

LLDKPALLQSDYIFEIYQKAHQQLIQFLDALAAGQRVGVVVKQHQSILEELKLYTQYTPDISNDLPQQDSTSFEPISNIEP  
LLDKPALLQSDYIFEIYQKAHQQLIQFLDALAAGQRVGVVVKQHQSILEELKLYTQYTPDISNDLPQQDSTSFEPISNIEP  
LLDKPALLQSDYIFEIYQKAHQQLIQFLDALAAGQRVGVVVKQHQSILEELKLYTQYTPDISNDLPQQDSTSFEPISNIEP  
!!

480  
480

logo

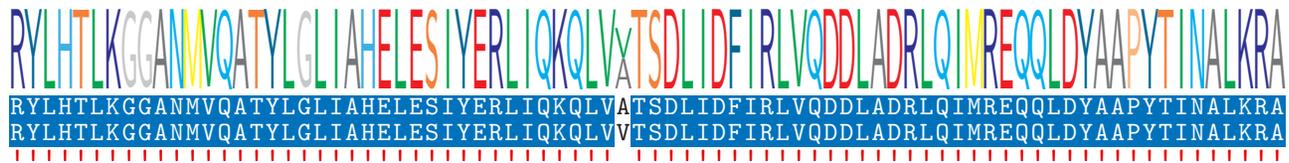
NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus

EPEWATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM  
EPEWATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM  
EPEWATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM  
!!

560  
560

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



640  
640

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



720  
720

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



800  
800

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



880  
880

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



960  
960

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1040  
1040

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1120  
1120

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1200  
1200

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1280  
1280

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1360  
1360

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1440  
1440

logo

NTDB id 110229 ACAK30 RS15700 WP 004745995.1  
NTDB id 1046 HON27 RS03130 WP 168726941.1  
consensus



1506  
1506

non conserved  
 similar  
 ≥ 50% conserved