

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

MKEILKTLTEAMLPEDSYSEQDAEFLEIFEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA
MKEILKTLTEAMLPEDSYSEQDAEFLEIFEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA
MKEILKTLTEAMLPEDSYSEQDAEFLEIFEEIEEIFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA
!!

80
80

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFKQKRAHTLDFRPLILLGQQQLQQQQSLEPALEELLQLS
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFKQKRAHTLDFRPLILLGQQQLQQQQSLEPALEELLQLS
ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFKQKRAHTLDFRPLILLGQQQLQQQQSLEPALEELLQLS
!!

160
160

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

NTLTAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR
NTLTAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR
HTLTAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETLTLFMEEAEEHLATHQFLDQELHQYDSYNALIRALHTLR
*!!

240
240

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ
GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQYREFVRDSLELLSRYSSSEQL EARLQFNQSWDAYVEQ
!!

320
320

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

HGDRTDPLMPSHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV
HGDRTDPLMPSHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV
HGDRTDPLMPSHGLVSQLQLDVSELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV
!!

400
400

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

LLDKPALLQSDYIFEIYQKAHQQLIQFLDALAAGQRVGVKQHQSI LEEELKLYTQYTPD-SNDLPQQDSTSFEPIYNIEP
LLDKPALLQSDYIFEIYQKAHQQLIQFLDALAAGQRVGVKQHQSI LEEELKLYTQYTPD-SNDLPQQDSTSFEPIYNIEP
LLDKPALLQSDYIFEIYQKAHQQLIQFLDALAAGQRVGVKQHQSI LEEELKLYTQYTPD-SNDLPQQDSTSFEPIYNIEP
!!

480
480

logo

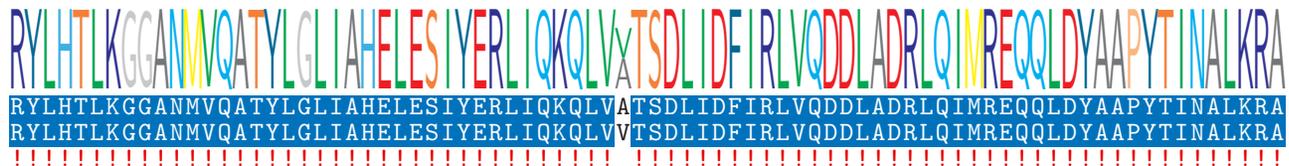
NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus

EPEWATVENFSDSxDIWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM
EPEVWATVENFSDSxDIWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM
EPEVWATVENFSDSxDIWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDIDLNIWVGEQENFAALNNLM
!!

560
560

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



640
640

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



720
720

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



800
800

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



880
880

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



960
960

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



1040
1040

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



1120
1120

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



1200
1200

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



1280
1280

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



1360
1360

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
NTDB id 1046 HON27 RS03130 WP 168726941.1
consensus



1440
1440

logo

NTDB id 110278 ACAK32 RS16135 WP 000658903.1
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consensus



1506
1506

X non conserved
X similar
X ≥ 50% conserved