

logo

NTDB id 374 SMU RS06885 WP 002262930.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIVFDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1143 A1552VC RS01965 WP 001883436.1	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIVFDLQPLINKWMQSENIATLTEIRRHFTLKGSGRMIGAKSSA	80
NTDB id 1417 SGL RS04200 WP 014407071.1	0
NTDB id 24804 NOC RS00860 WP 002812134.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus	0

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NTDB id 374 SMU RS06885 WP 002262930.1	ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQLQQQSLEPALEELLQLS	0
NTDB id 413 AAK55817.1 141..845()	0
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NTDB id 24804 NOC RS00860 WP 002812134.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
consensus	0

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NTDB id 374 SMU RS06885 WP 002262930.1	HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR	0
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consensus	0

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NTDB id 374 SMU RS06885 WP 002262930.1	GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQEYREFVDRSLELLSRYSSSEQLERLLQFNQSWDAYVEQ	0
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 1143 A1552VC RS01965 WP 001883436.1	0
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consensus	0

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NTDB id 374 SMU RS06885 WP 002262930.1	HGDRTDPLMP SHGLV S QLLQLD VSE LLD AELDFEKG I RNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV	0
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NTDB id 1143 A1552VC RS01965 WP 001883436.1	0
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NTDB id 374 SMU RS06885 WP 002262930.1	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAAGQRVGVVKQHQSI L EELKLYTQYTPDISNDLPQQDSTSFEP IYNI EP	0
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NTDB id 1417 SGL RS04200 WP 014407071.1	0
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consensus	0

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NTDB id 374 SMU RS06885 WP 002262930.1	EPEVWATVENFSDSADWAVLGQSVQQDRQYISSVQLCPNDA D LLYKYLXNEYTLG F SCDSDLA W KSE QEN F T E S N T	0
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NTDB id 1046 HON27 RS03130 WP 168726941.1	EPEVWATVENFSDSADWAVLGQSVQQDRQYISSVQLCPNDA D LLYKYLXNEYTLG F SCDSDLA W KSE QEN F T E S N T	558
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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consensus	0

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NTDB id 374 SMU RS06885 WP 002262930.1	LMRYLHTL...KGGANMVQATYLG LIAHELES IYERLIQKQLVVTSD LIDFIRLVQDDLADR...LQIM...REQQL	0
NTDB id 413 AAK55817.1 141..845()	0
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NTDB id 1046 HON27 RS03130 WP 168726941.1	LMRYLHTL...KGGANMVQATYLG LIAHELES IYERLIQKQLVVTSD LIDFIRLVQDDLADR...LQIM...REQQL	626
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NTDB id 374 SMU RS06885 WP 002262930.1
 NTDB id 413 AAK55817.1 141..845()
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 NTDB id 24804 NOC RS00860 WP 002812134.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

DYMMFYETEXMFKRAGQNSNFYRSPVYELAYRSESEYYSLYNXISEIIFBETDVELEPAALAEAELEHHSQYITAVTELATPV

 DYMMAKESEVM.....GFVRS..ELAYASGGRFYGLVNA.....FQRLDVSIEA...AREHAQQRYYITG.....
 DYAAPYTINALKRAGQNSNFQPLPVVEAFDAESEVFSEQKVISEIIIDEIPVELEPALAELETHHDQVFDTAVTELATPV

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706
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 NTDB id 24804 NOC RS00860 WP 002812134.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

EMITTVTSQENEVAANEEDD EAWVEQSFEASELLEMARERLLKQWFERTRALRSLVHSLKGGARMVDFERYE

SGDQIK...TSVLPQSSSYVGSERYLLH...KRYHWAYLELLKRS.....DFDDIL
 EMITTVTSQENEVAANEQDIEAVVEQTFLEEATELLEMA.ESLLKQWFEQRTNRSILLQLQRAVHSLKGGARMVGLEAVQ

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212
785
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0
0

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 NTDB id 24804 NOC RS00860 WP 002812134.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

KYRSLNMYEYKYHFNLYTHLLESAIAWLKDAIFNHNHYSRFDGLEQIYENIQFFETTIIQINERLKRADDFTSYEY

 LVDIDGNVVYSIYKYDNFGTNLLTGK.....YQDTNLGHTFKRL...EQTV.....NEQRKTNEDFTPVVI
 AIAYQLENAFEQFALHFNNSNIYDHLLESIAIWLKDAIFNHNHYSRFDGLEQIYENIQFFETTIIQIPTKLTRADLFSSEP

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0
270
865
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 NTDB id 1417 SGL RS04200 WP 014407071.1
 NTDB id 24804 NOC RS00860 WP 002812134.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus

SDYFYQEDGTEYKEPPPMGAWLESTERTQQGYDHNNALRYSSNLTKMIRESSSESNRSTLYDLDHPSRTLYSAEIAVEKS

 SDFVQEDGKQY....AWLGAPIIQGGYLHSYAMFRLPSNAITKLI AEGSSNPSM.QTILVGGDHRSTRLTSAEIAVEKS
 MTFIQGDGTEPPPMGAWEQTERL....DQNNEMIRVSADLIEKMIDLSGENSINRSRIEMDLSQFSHTLV.....

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logo

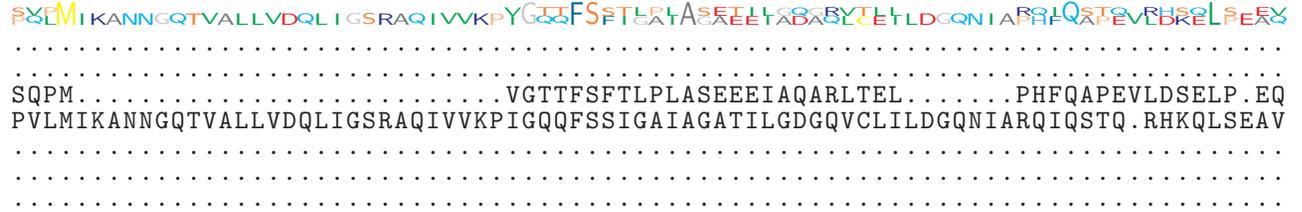
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 NTDB id 1046 HON27 RS03130 WP 168726941.1
 NTDB id 1417 SGL RS04200 WP 014407071.1
 NTDB id 24804 NOC RS00860 WP 002812134.1
 NTDB id 1047 HON27 RS03110 WP 000389061.1
 consensus



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715
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logo

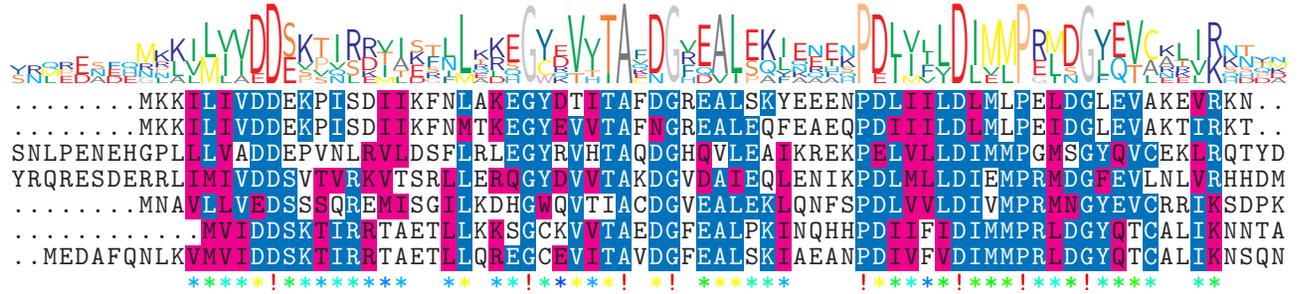
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 consensus



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1370
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logo

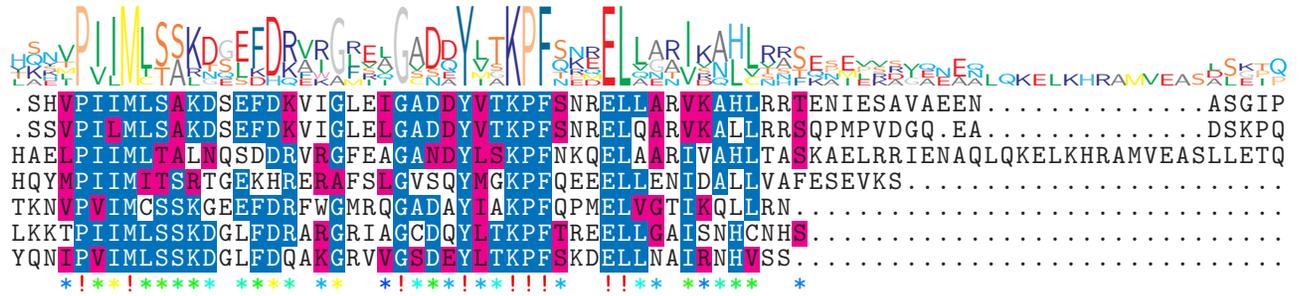
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 consensus



70
70
840
1450
72
68
78

logo

NTDB id 374 SMU RS06885 WP 002262930.1
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 NTDB id 1143 A1552VC RS01965 WP 001883436.1
 NTDB id 1046 HON27 RS03130 WP 168726941.1
 NTDB id 1417 SGL RS04200 WP 014407071.1
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 consensus



135
134
920
1506
121
118
127

