

logo

NTDB id 969117 MHH79 RS12735 WP 015483431.1  
NTDB id 96 BSU 24730 NP 390353.1  
NTDB id 1028 TT RS08215 WP 011173991.1  
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK  
.....  
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK

0  
0  
80

logo

NTDB id 969117 MHH79 RS12735 WP 015483431.1  
NTDB id 96 BSU 24730 NP 390353.1  
NTDB id 1028 TT RS08215 WP 011173991.1  
consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAF LNPLD T L S LEEVEDLTGLVVEPYQT T KSAFLYALAKHYPELGLPVPPPPSG  
.....  
ALLPAEKAKELKAIPFALDEEAGVVRVAF LNPLD T L S LEEVEDLTGLVVEPYQT T KSAFLYALAKHYPELGLPVPPPPSG

0  
0  
160

logo

NTDB id 969117 MHH79 RS12735 WP 015483431.1  
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NTDB id 1028 TT RS08215 WP 011173991.1  
consensus

EGQKDLKLGELL LQGWISREALEEALVEQEK TGDLLGRILVRKGLPEEALYRALAEQGLEFLESTEGIVDPDSAALLL  
.....  
EGQKDLKLGELL LQGWISREALEEALVEQEK TGDLLGRILVRKGLPEEALYRALAEQGLEFLESTEGIVDPDSAALLL

0  
0  
240

logo

NTDB id 969117 MHH79 RS12735 WP 015483431.1  
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consensus

LRSDALRYGAVPIGFQNGEVEV LSDPRHKEAVAQ LLNRPARFYALPQAW EELFRRAYPQKNRLGEVLVQEGKLSREAL  
.....  
LRSDALRYGAVPIGFQNGEVEV LSDPRHKEAVAQ LLNRPARFYALPQAW EELFRRAYPQKNRLGEVLVQEGKLSREAL

0  
0  
320

logo

NTDB id 969117 MHH79 RS12735 WP 015483431.1  
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consensus

KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLED TLVQSGKLRPEALAQAVATQLGYPYVDPEEDPP  
.....  
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLED TLVQSGKLRPEALAQAVATQLGYPYVDPEEDPP

0  
0  
400

logo

NTDB id 969117 MHH79 RS12735 WP 015483431.1  
NTDB id 96 BSU 24730 NP 390353.1  
NTDB id 1028 TT RS08215 WP 011173991.1  
consensus

DPGAPLLL PEDLCRRYGVFPHRLEGNRLVLLMKDPRN I LALDDVRLALKRKGLN YEVAPAVATEAAITKLIERFYGKAEL  
.....  
DPGAPLLL PEDLCRRYGVFPHRLEGNRLVLLMKDPRN I LALDDVRLALKRKGLN YEVAPAVATEAAITKLIERFYGKAEL

0  
0  
480

