

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1034065 AB6A22 RS11885 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTI^GDKRLGAALLDAGLLTDEELQRALERHREVGGS^LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK
MSVLTI^GDKRLGAALLDAGLLTDEELQRALERHREVGGS^LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 80
..... 0
..... 0
..... 0

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NTDB id 1034065 AB6A22 RS11885 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

ALLPAEKAKELKAI^PFALDEEAGV^RVAF^LNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG
ALLPAEKAKELKAI^PFALDEEAGV^RVAF^LNPLDTLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 160
..... 0
..... 0
..... 0

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consensus

EGQKDLKLGELL^LLQK^GWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL
EGQKDLKLGELL^LLQK^GWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL 240
..... 0
..... 0
..... 0

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consensus

LRSDALRYGAVPI^GFQNGEVEV^LSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL
LRSDALRYGAVPI^GFQNGEVEV^LSDPRHKEAVAQLLNRPARFYALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 320
..... 0
..... 0
..... 0

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consensus

KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLG^QTYVEQ^TER
KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLG^QTYVEQ^TER 399
..... MSQ^YTICE^QQ^TERI 14
..... 0
..... 0
..... *

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consensus

FDIS^PSRWQPNLLPK^LLR^VLV^PRLTGN^RLW^LLVK^DPRNLTAL^EIV^RLM^TR^RY^OYEP^VWITSDELTY^LL^NALS
... PDPGAPLLLPEDLCRRYGVFPHRLEGNRLVLLMKDPRNLTALDDVRLALKRKG^LNYEVAPAVATEAAITK^LI... 471
FDIS^PSRWQKNCEEKQLLLRYLAVPVQETEHK^LWLAVDDENNLTACEIFAFM.TH... KQIEP^VVVIASDELKYL^LNALS 89
..... 0
..... 0
..... *

 non conserved
 similar
 $\geq 50\%$ conserved