

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 204409 BAMY RS12235 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVKALLPA
.....

85
0
0
0

logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 204409 BAMY RS12235 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

EKAKELKAIPFALDEEAGVVRVAFNLPLDTLGLEEVEDLTGLVPEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE
 EKAKELKAIPFALDEEAGVVRVAFNLPLDTLGLEEVEDLTGLVPEPYQTTKSAFLYALAKHYPELGLPVPPPPSGEGQDKLKLGE

170
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF
 LLLQKGWISREALEEALVEQEKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAALLLLRSDALRYGAVPIGF

255
0
0
0

logo

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NTDB id 96 BSU 24730 NP 390353.1
consensus

QNGEVEVVLSDPRHKEAVAQLLNRPARFYLLALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
QNGEVEVVLSDPRHKEAVAQLLNRPARFYLLALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREALKEALEVQKGLPRAKPLGEIL
.....

340
0
0
0

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consensus

VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGSPYTYEQTTERPSRWQPNLLPKQLRLRYLYYP
VELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATMLGSPYTYEQTTERPSRWQPNLLPKQLRLRYLYYP
.....MSQYTICEQQTERRIFDISPSRWQKNCEEKQLLLRYLAVP
.....* * * * *

420
39
0
0

logo

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consensus

HSP70

YRLTEENRLLWLVLYKSDERNITALEIYLRIIMLTRKGGLNKEETAIATAAEAAATKKLTNALSPQEPPFEEAIAEAKFAKKQAEVEVPS
HRLEGNRLIVILMKDPRNILACDDVLRALKRKGLNYEVAPAVATEEAITTKLI.....ERFYGKAELSELTAKEFAKKQAEVEVPVS
VQETEHLWLVAIDENNLTACEIFAFM.TH...KQIEPVIASDELKYLLNALSPPEQQPIYEESLAFVEQE.....QE

* * * * *

498
109
0
0

