

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1058720 ACFW98 RS05810 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 80
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK 0
..... 0
..... 0
..... 0

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NTDB id 1058720 ACFW98 RS05810 WP 003153083.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 160
 ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 0
 0
 0
 0

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consensus

EGQKDLKL	GELLLQKGWISREALEEALVEQEK	TGDL	LGRILVRKGLPEEALYRALAEQK	GLEFLESTEGIVPDPSAALLL	240
EGQKDLKL	GELLLQKGWISREALEEALVEQEK	TGDL	LGRILVRKGLPEEALYRALAEQK	GLEFLESTEGIVPDPSAALLL	0
.	.	.	.	.	0
.	.	.	.	.	0
.	.	.	.	.	0

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consensus

LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 320
 LRSDALRYGAVPIGFQNGEVEVWLSDPRHKEAVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGKLSREAL 0
 0
 0
 0
 0

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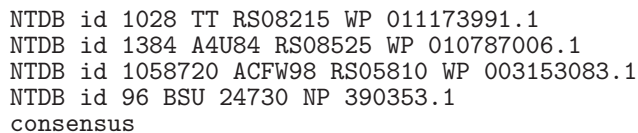
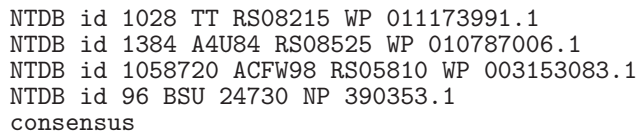
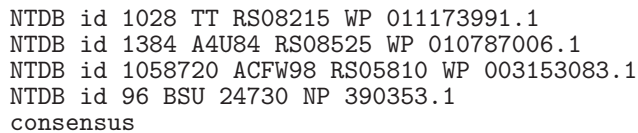
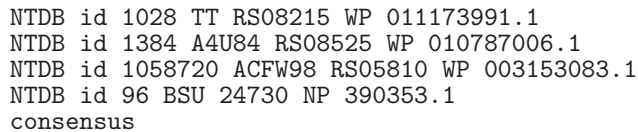
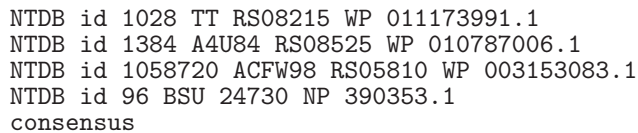
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 KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDP. 399
MSQYTICQQTTERI 14
 0
 0

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consensus
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FDISPSRWQPNLLKQLRLRYLVYPVRLTECHLWLVKDERNLTLALELYRMLTRKGLNKYEIVWITSDEITYLLINALS
 ...PDPGAPLLLPEDLCRRYGVFPHRLEGNRLVLLKDPKNILACDDFAFLKPKGLNKEIAIAAAEAAATKLLINALS 471
 FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLVADDENNLTAEIFAFM.TH...KQIEPVVIASDEIKYLLINALS 89
 0
 0

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consensus



 non conserved
 similar
 $\geq 50\%$ conserved