

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1019020 AB0R79 RS13055 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus

MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV 80
MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPVKV 0
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..... 0
..... 0

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NTDB id 1019020 AB0R79 RS13055 WP 012117985.1
NTDB id 96 BSU 24730 NP 390353.1
consensus
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ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLDTLSEEEVDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPPPSG 160
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consensus
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EGQKDLKLGELLLQKGWISREALVEEQKTGDLLGRILVRKGLPEEALYRALAEQKGLFLESTEGIVPDPSAALLL 240
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..... 0
..... 0
..... 0

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consensus

L R S D A L R Y G A V P I G F Q N G E V E V V L S D P R H K E A V A Q L L N R P A R F Y L A L P Q A W E E L F R R A Y P Q K N R L G E V L V Q E G K L S R E A L 320
 L R S D A L R Y G A V P I G F Q N G E V E V V L S D P R H K E A V A Q L L N R P A R F Y L A L P Q A W E E L F R R A Y P Q K N R L G E V L V Q E G K L S R E A L 0
 0
 0
 0

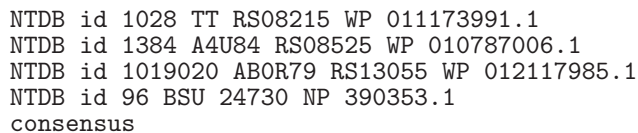
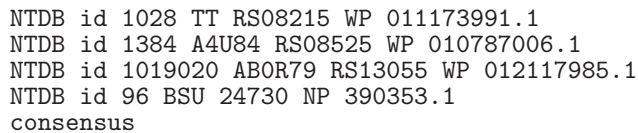
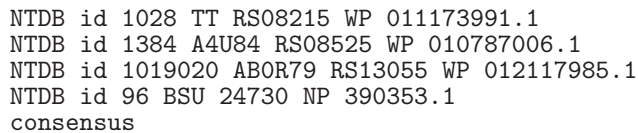
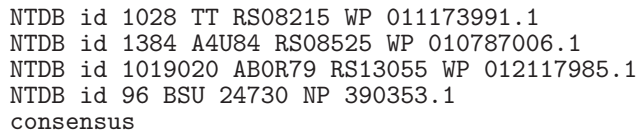
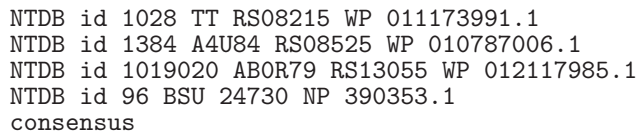
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consensus
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 KEALEVQKGLPRAKPLGEILVELGLARPEDVEEALQKQRRGGGRLEDTLVQSGKLRPEALAQAVATQLGYPYVDPEEDP. 399
MSQYTICQQTTERI 14
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consensus
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FDISPSRWQPNLLKQLRLRYLVYPVRLTECHLWLVKDERNLTLALELYRMLTRKGLNKYEIPVWITSDEITYLLINALS
PDPGAPLLLPEDLCRRYGVFPHRLGNRLVLLKDPKNITLADVRLALKRKGLNVEVAPVATEAAITKLLI... 471
 FDISPSRWQKNCEEKQLLLRYLAVPVQETEHKLWLAVDDENNTACEIFAFM.TH...KQIEPVVIASDEIKYLLINALS 89
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 0

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consensus



 non conserved
 similar
 $\geq 50\%$ conserved