

logo

NTDB id 1028 TT RS08215 WP 011173991.1
NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 1041407 ACDZ79 RS11740 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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consensus

ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLD~~TL~~SL~~EE~~VEDLTGLVVEPYQT~~TK~~SAFLYALAKHYPELGLPVPPPPSG 160
ALLPAEKAKELKAIPFALDEEAGVVRVAFNLPLD~~TL~~SL~~EE~~VEDLTGLVVEPYQT~~TK~~SAFLYALAKHYPELGLPVPPPPSG 0
..... 0
..... 0
..... 0

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consensus

EQQKDLKLGELLLQKGWISREALVEEQKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL 240
 EGQKDLKLGELLLQKGWISREALVEEQKTGDLLGRILVRKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAALLL
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 0
 0
 0

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L R S D A L R Y G A V P I G F Q N G E V E 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
 L R S D A L R Y G A V P I G F Q N G E V E 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
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 0
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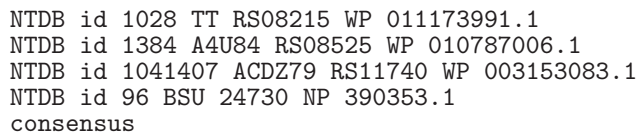
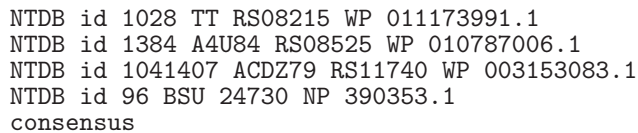
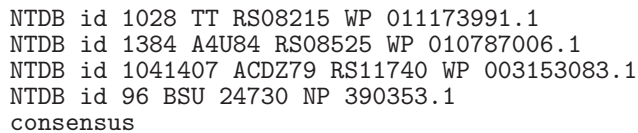
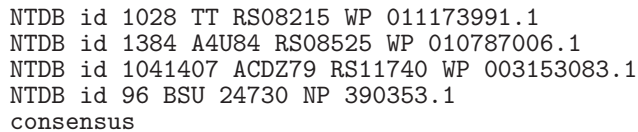
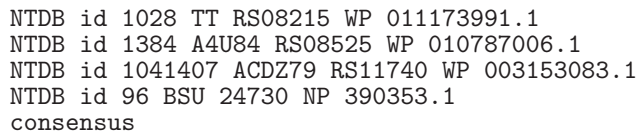
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 KEAL**EVQ**KGL**P**RAK**P**L**G**E**I**L**V**E**L**G**L**AR**P**ED**V**EEAL**Q**K**Q**RRGGGR**L**ED**T**LV**Q**S**G**K**L**R**P**EAL**A**Q**A**V**A**T**M**L**G****P****C**D**P**E**E**D**P**. 399
MS**Q****T****I****C****E****Q****T****T****E****R**I 14
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 0

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[illegible]

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 non conserved
 similar
 $\geq 50\%$ conserved