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NTDB id 979963 AAFA20 RS16960 WP 002014066.1
NTDB id 1054 ABD1 RS15870 WP 002014066.1
NTDB id 1084 FDQ49 RS11070 WP 004711654.1
consensus
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M L K L L K P K H R S I L G I D I S S A V K I L E I S G S G E D I C V E G Y G R E E T P A S A L E G N M I K D V V A N C I K K L V E R L K F T S K N V A L 80
M L R L Y R R P K K G L V G V D I S S T S V K V L E L S V K N N Q Y R V E S Y G L V P L L E N S V V E K N I L N P E A V A D A L T R S I N L A N P Q A I N A A I 80
M L R L Y R K P N K G L M G V D I S S T S V K L L E L S V K N G K Y W V E S Y A L M P L P E N S V V E K N I L N P E A V A E A L E R A M N L A N P Q T T H A A I 80
M L R L Y R K P N K G L M G V D I S S T S V K L L E L S V K N G K Y W V E S Y A L M P L P E N S V V E K N I L N P E A V A E A L E R A M N L A N P Q T T H A A I 80
M L R L Y R K P N K G L M G V D I S S T S V K L L E L S V K N G K Y W V E S Y A L M P L P E N S V V E K N I L N P E A V A E A L E R A M N L A N P Q T T H A A I 80

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AVPTS¹VIHK²TI³EMDAD⁴MS⁵DDERE⁶VQIR⁷MDA⁸EQ⁹YIP¹⁰FLDE¹¹VS¹²LDFE¹³VL¹⁴PD¹⁵RLAN¹⁶PNRV¹⁷NV¹⁸LV¹⁹AT²⁰RTEN²¹VET²²RV²³EV²⁴LE²⁵L

DA¹VS²YV³QINE⁴GL⁵TL⁶ME⁷EL⁸WTE⁹DR¹⁰Y¹¹TI¹²IN¹³MS¹⁴EP¹⁵AK¹⁶SS¹⁷SL¹⁸LD¹⁹V²⁰LA²¹SA²²RA²³EN²⁴VN²⁵TR²⁶VE²⁷AA²⁸HR

AVPDS¹AV²ISK³V⁴VQ⁵INE⁶GL⁷TD⁸LE⁹ME¹⁰EEL¹¹V¹²VI¹³EAD¹⁴K¹⁵YIP¹⁶Y¹⁷P¹⁸I¹⁹DE²⁰IN²¹LDFE²²VL²³GHS²⁴AK²⁵SS²⁶SL²⁷LD²⁸V²⁹LA³⁰SA³¹RA³²EN³³VN³⁴TR³⁵VE³⁶AA³⁷HR 160

AVPTS¹MS²V³VH⁴K⁵TI⁶EMDAD⁷MT⁸DDERE⁹VQIR¹⁰MDA¹¹ERY¹²IP¹³FLDE¹⁴VS¹⁵LDFE¹⁶VL¹⁷MS¹⁸EP¹⁹LAN²⁰PNRV²¹SV²²LA²³AT²⁴RTEN²⁵VD²⁶TR²⁷IE²⁸VE²⁹LE³⁰L 160

AVPTS¹TVIHK²TI³EMDAD⁴MSDDERE⁵VQIR⁶MDA⁷EQ⁸YIP⁹FLDE¹⁰VS¹¹LDFE¹²VL¹³PD¹⁴RLAN¹⁵PNRV¹⁶NV¹⁷LV¹⁸AT¹⁹RTEN²⁰VET²¹RV²²EV²³LE²⁴L 160

AVPTST¹VIHK²TI³EMDAD⁴MSDDERE⁵VQIR⁶MDA⁷EQ⁸YIP⁹FLDE¹⁰VS¹¹LDFE¹²VL¹³PD¹⁴RLAN¹⁵PNRV¹⁶NV¹⁷LV¹⁸AT¹⁹RTEN²⁰VET²¹RV²²EV²³LE²⁴L 160

AVPTST¹VIHK²TI³EMDAD⁴MSDDERE⁵VQIR⁶MDA⁷EQ⁸YIP⁹FLDE¹⁰VS¹¹LDFE¹²VL¹³PD¹⁴RL¹⁵LN¹⁶PNRV¹⁷NV¹⁸LV¹⁹AT²⁰RTEN²¹VET²²RV²³EV²⁴LE²⁵L 160

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consensus
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[illegible]

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consensus

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[illegible]

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consensus
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FLQMGFSPQVDYQKIENDASSLMVACGLALRSFD

L S H M S F G K K V N I D V I N S D A P T L M V A C G L A L R H V E . 354
F L Q M G F S P Q V D I K K V E N D A S S L M V A C G L A L R S F D E 353
F L Q M G F S P Q V D Y Q K I E N D A S S L M V A C G L A L R S F D . 352
F L Q M G F S P Q V D Y Q K I E N D A S S L M V A C G L A L R S F D . 352
F L Q M G F S P Q V D Y Q K I E N D A S S L M V A C G L A L R S F D . 352

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 non conserved
 similar
 $\geq 50\%$ conserved