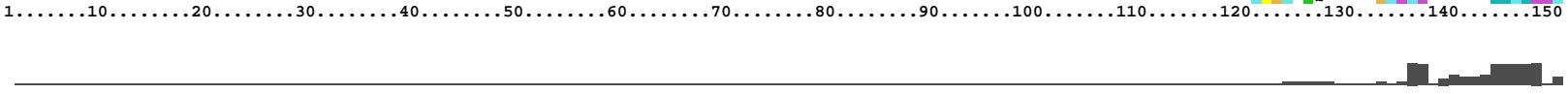
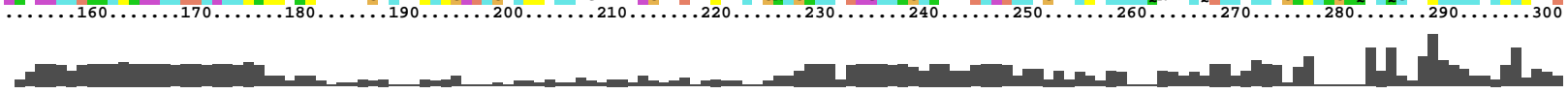


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 gi | 114556049 | ref | XP_524676.2 | M C V C A G C R A A P S R R G A G P L Q V A G G W S E G A D M D - Y D S Y Q H I F Y D - Y 150
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 gi | 194665470 | ref | XP_593398.4 | M C K C A G C F A A P S G R G A G P L Q V A G G S E G A D M D - F D S Y Q H I F Y D - Y 150
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 gi | 50759960 | ref | XP_425790.1 | 150
 gi | 113681627 | ref | NP_001038607.1 | M P G I S A S A S R I K K W E M E Y D Q Y Q H I F Y D D E 150
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gi | 6678704 | ref | NP_032532.1 | D C G E D F Y R S T A P S E D I W K F E L V P S P P T S P P W G S G P G A V D P A S G - - - I N P G E P W P G G G A G D E A E S R G H S K A W G R N Y A S I I R R D C M W S G F S A R E R L E R V S D R L A P G A P - - - R G N P P K A P A T P D G T P - - - S L E A S N P A P A T Q C Q L G E P K 300
 gi | 109475447 | ref | XP_233490.4 | D C G E D F Y R S T A P S E D I W K F E L V P S P P T S P P W G S G P G A V D P A S G - - - I S P G E P W P G G G A G D E A E S R G H S K A W G R N Y A S I I R R D C M W S G F S A R E R L E R V S D R L A P G V P - - - R G N P P K A P A T P D G T P - - - S L E A S N P A P A T Q C Q L G E P K 300
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gi | 6678704 | ref | NP_032532.1 | T Q A - C S G S E S P S D S - E G E E - - - I D V V T V E K R - - - R E L D I R K P V T I T V R A D P L D P C M K H F H I S I H Q O Q H N Y A A R F P P E S C S Q E G D P E P G P Q E E A P E I E A P K E K E E E E E E E E E E - - - I V S P P P V G S E A P Q S C H P K P - V S S D T E D 450
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gi | 6678704 | ref | NP_032532.1 | V T K R K N H N F L E R K R R N D L R S R F L A L R D Q V P T L A S C S K A P K V V I L S K A L E Y L Q A L V G A E K K M A T E K R Q L R C R Q O O L Q K R I A Y L S G Y 535
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