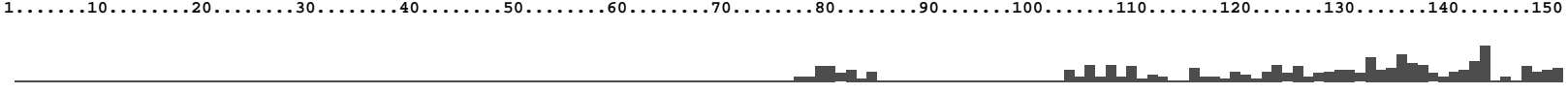
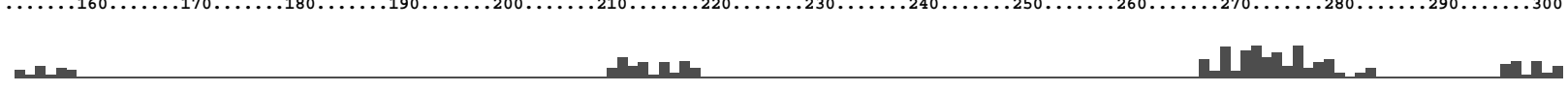


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gi|58866008|ref|NP_001012216.1|-----MLSDDDVN-----EIIIEVENVP--SGVQRNVSSNITLTKRSKAPNFMSLSE---ADT 150
gi|157426860|ref|NP_001098718.1|MSAYYRNNWSEEDPDYDYSGSONRTOGYLKTQGYVDPVPGGLNNDYVPGTRSNPYSVAERTRPDYPGSLAEPNYPRELSDNDYSGTRSNAYSAASRTSPDHTSLPEPDVSEFQSHPHYHRASSRDPDYPGQRNPDYAGSSSSGNYAGSR 150
gi|114661299|ref|XP_001153396.1|MSAYYRNNWSEEDPDYDYSGSONHMGYVLTQDYVDPVPGGLNNDYVPGTRSNPYSVAERTRPDYPGSLAEPNYPRELSDNDYSGTRSNAYSAASRTSPDHTSLPEPDVSEFQSHPHYHRASSRDPDYPGQRNPDYAGSSSSGNYAGSR 150
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gi|119916833|ref|XP_591622.3|MFAYRNWSEEDPDYDYSGSONHMGYVLTQDYVDPVPGGLNNDYVPGTRSNPYSVAERTRPDYPGSLAEPNYPRELSDNDYSGTRSNAYSAASRTSPDHTSLPEPDVSEFQSHPHYHRASSRDPDYPGQRNPDYAGSSSSGNYAGSR 150
gi|189517063|ref|XP_689991.3|-----MSSDDVD-----EIIIEVENVP--SGVQNHVLSIALRKSANPAFCVLSSE---AADR 150



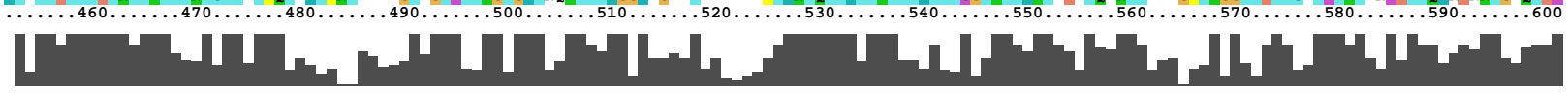
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gi|58866008|ref|NP_001012216.1|IDCQIFS-----HGNDRRHNR-----LRFSSLSNISQIYHG-----PECLGM 300
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gi|114661299|ref|XP_001153396.1|VHPDFGSL-EPDYPGAQNSNDYVPGHRANLNHPGSRKNLEHTSFRINPYVADSLGKPDYVPGADIDPNSPPFFGEPDYPGAEDNOLPSTWREPDYDAENGHDYGSSETPKMTGRVLSRTSSIQPFRRHSDDPVGSLWGENEDYPEGIEM 300
gi|73958977|ref|XP_547109.2|INPDYLDYLGEPDYPGACGNTNHPGSRANSNHPGSRRNPEHAGSGIITPVVDVLGEPDYPGAEINQNSNPFWEVDYVPAE---NSNPFWGKPDYVTEGDRDYGSSETSITREVLVSRSSRPLRHRGDSGLGVIGRENEDYVNTIAM 300
gi|119916833|ref|XP_591622.3|INPDYLDYLGEPDYPGACGNTNHPGSRANSNHPGSRRNPEHAGSGIITPVVDVLGEPDYPGAEINQNSNPFWEVDYVPAE---NSNPFWGKPDYVTEGDRDYGSSETSITREVLVSRSSRPLRHRGDSGLGVIGRENEDYVNTIAM 300
gi|189517063|ref|XP_689991.3|-----LTAHEITQEVEN----- 300



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gi|32441288|ref|NP_083206.1|YFSFLRWLLKFNIFSFVMNFSFIIPQFVGA--KNTLQFTGLEFFTGAGYFGDTVMYYGFYNTSTIRHRMGGASYNMOLAYIFTIGACLIVCFPSLLFSMAKYFRNNFINPH-IYSRGIKLIKICWDFTVTHEKAVKLRKQKNLSTEIRE 600
gi|58866008|ref|NP_001012216.1|YFSFLRWLLKFNIFSFVMNFSFIIPQFVGE--KNTLQFTGLEFFTGAGYFRETVMYYGFYNTSTIRHRMGGASYNMOLAYIFTIGACLIVCFPSLLFSMAKYFRNNFINPH-IYSRGIKLIKICWDFTVTHEKAVKLRKQKNLSTEIRE 600
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gi|73958977|ref|XP_547109.2|YFNFLRWLLKFNIFSFIVTFSFIIPQFVAB--KNTLQFTGLELLGAGYFRDVMYYGFYNTSTIRHGNGRASYNMOLAYIFTIGACLITCFPSLLFSMAKYFRNNFINPH-IYSRGIKLIKICWDFTVTNETAVKLRKQKNLSTEIRE 600
gi|119916833|ref|XP_591622.3|YFNFLRWLLKFNIFSFIVTFSFIIPQFVME--KNHLQFTGLEFLTGAGYFNTVMYYGFYNTSTIQHNSGASYNMOLAYIFTIGACLIVCFPSLLFSMAKYFRNNFINPH-IYSRGIKLIKICWDFTVTHEKAVKLRKQKNLSTEIRE 600
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gi|58866008|ref|NP_001012216.1|NLSELRQENDRLTLNQLTRFSAHVAAWLVSTGTAACCVAVVYLAEYNSFLKTHK---NPGAVLLLFPVVSINLAVPRFYSMFRLVEREYIPRQEVVLLVLRNIFLKISIVGILCYWLNIVALSGEECWEELIGODIYRLLMD 750
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gi|114661299|ref|XP_001153396.1|NLSELRQENSKLTFNQQLTRFSAHVAAWLVSTGVAIAACAAVYVLAEYNEFLKTHS---NPGAVLLLFPVVSINLAVPCISYMFRLVEREYIPRQEVVLLVLRNIFLKISITIGILCYWLNIVALSGEECWEELIGODIYRLLMD 750
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