Alignments of cDNAs and genomic DNAs were used to define the intergenic and intronic DNA sequences. CDS: coding sequence. Intron: non-coding sequence between the start and stop codons of the CDS. Intergenic: Non-coding sequence outside of the CDS. Divergence scores determined as per[77]. Codon substitution rates were determined as per[71]. SE is the Standard Error. ND: Not determined.