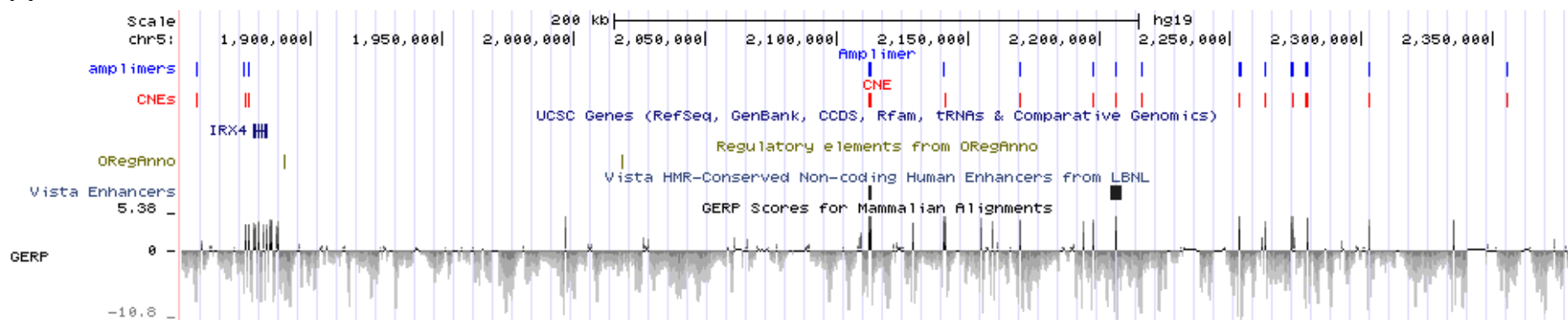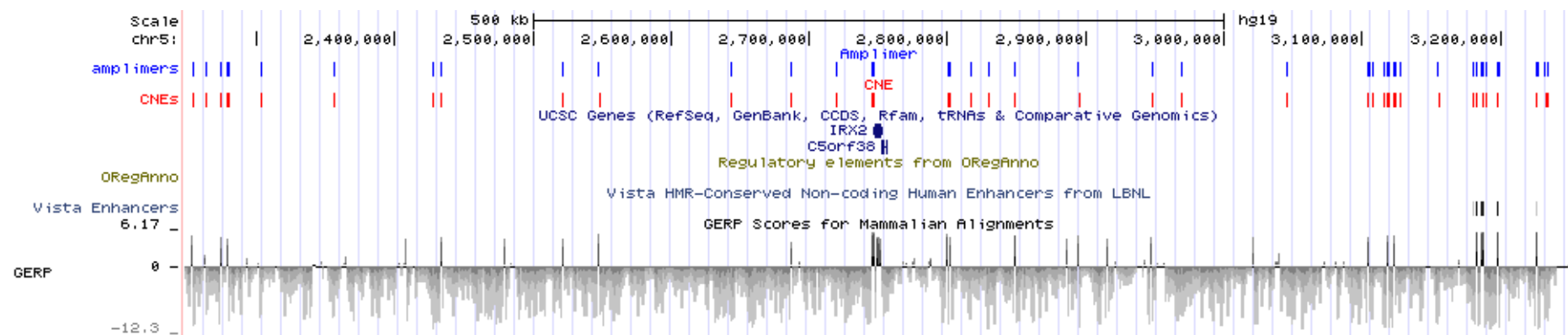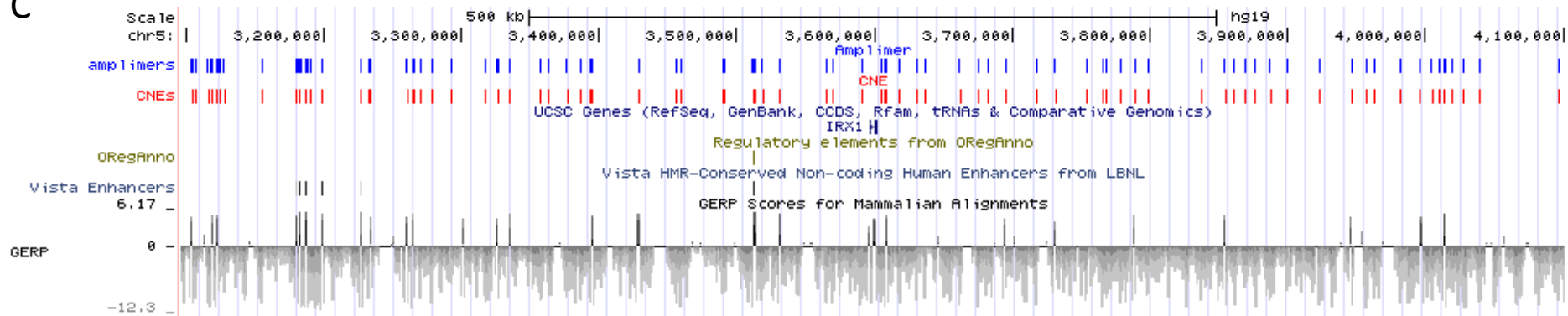Figure S1.

Figure S1 Regional plots of sequenced CNEs and the amplimers used to capture them surrounding *IRX4* (A), *IRX2* (B) and *IRX1* (C). Map positions are from GRCh37/hg19. Regulatory regions annotated with ORegAnno or Vista are depicted (genome.ucsc.edu). Evolutionary constraint is indicated using Genomic Evolutionary Rate Profiling (GERP) scores.