Figure S1 - Independent contrast plots of recombination rate vs genome compactness and chromosome number. Lines on each plot are the best-fit linear regression. The Pearson's correlation coefficient (*r*) and p-value (*p*) are given on each plot.
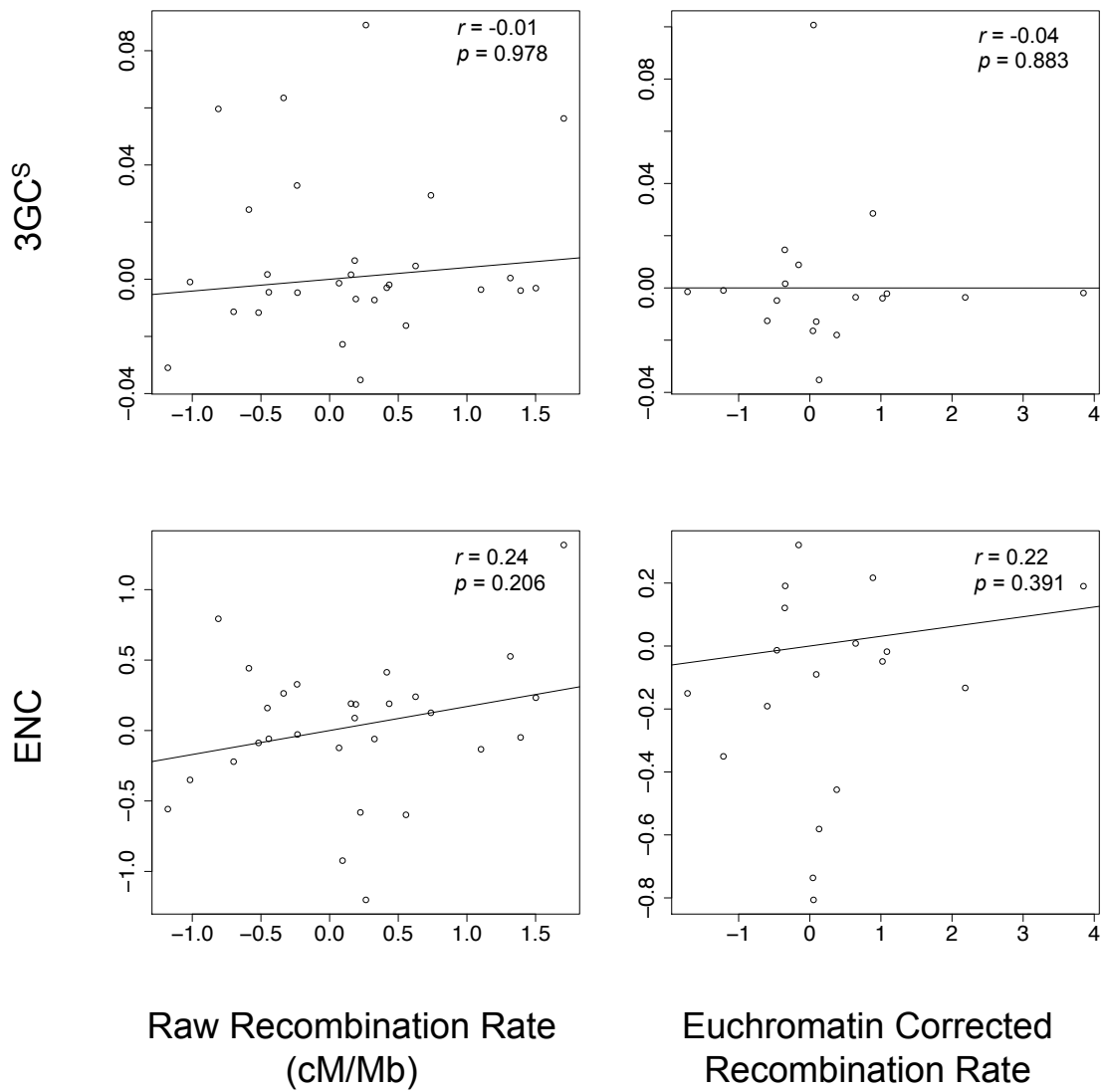
Figure S2 - Independent contrast plots of recombination rate vs GC content at 3rd position synonymous sites ($3GC^S$) and codon bias (ENC). Lines on each plot are the best-fit linear regression. The Pearson's correlation coefficient (r) and p-value (p) are given on each plot.