Combined Fragment Data

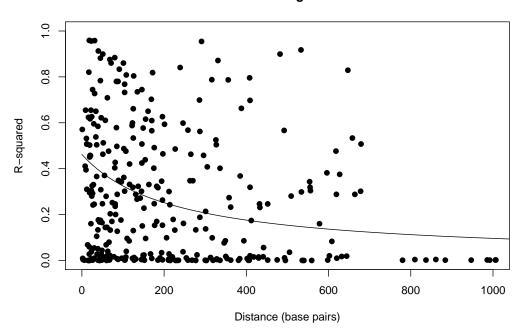


Figure S7: R^2 versus distance for the combined data. Only SNPs with a minor allele frequency $\geq 10\%$ were used in pairwise comparisons. Trend line was fitted as described in the Methods section.