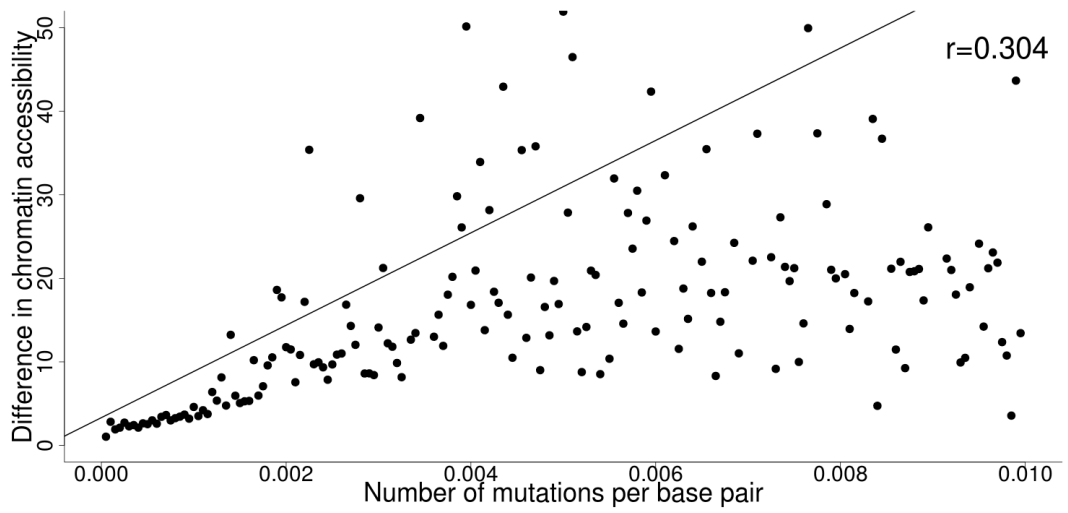


Figure S4



Within-pair differences in chromatin accessibility according to the number of mutations per base pair as an estimate of the density of mutations relative to the size of the open chromatin region.