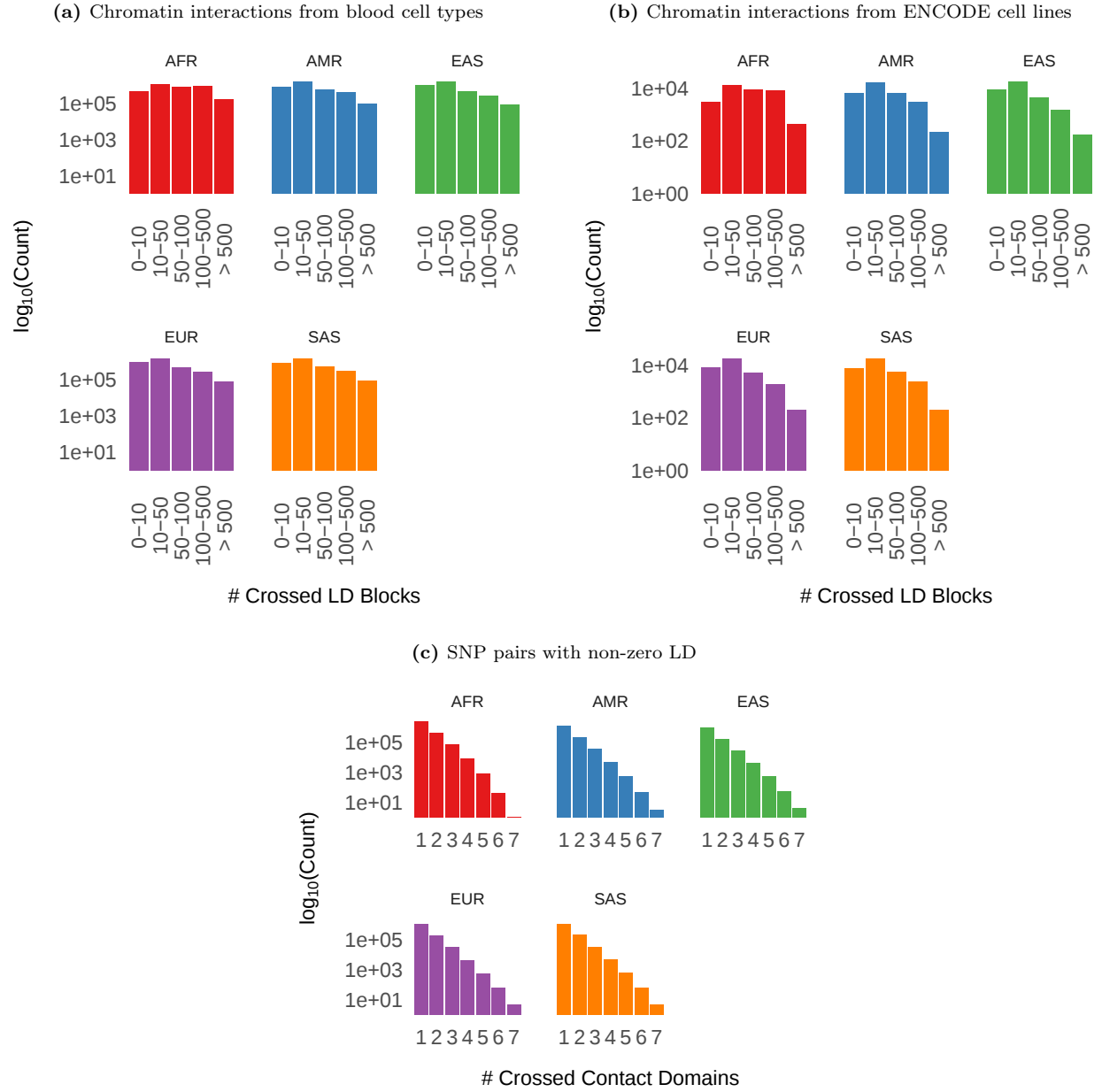




Supplemental Figure 6. Number of LD blocks crossed by statistically significant chromatin interactions across all chromosomes in (A) 17 primary blood cell types (Javierre et al. 2016) and (B) 5 ENCODE cell lines (Rao et al. 2014), grouped by super-population. Many statistically significant interactions cross hundreds or thousands of LD blocks. (C) Number of contact domains crossed by SNP pairs with non-zero LD, summed over 5 ENCODE cell lines, grouped by super-population.