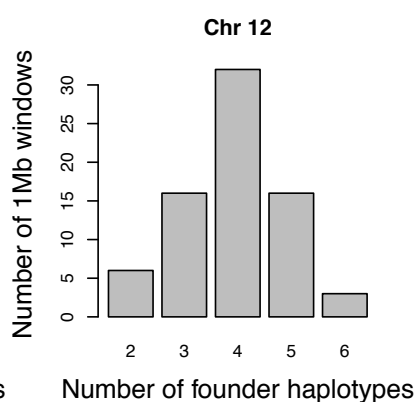
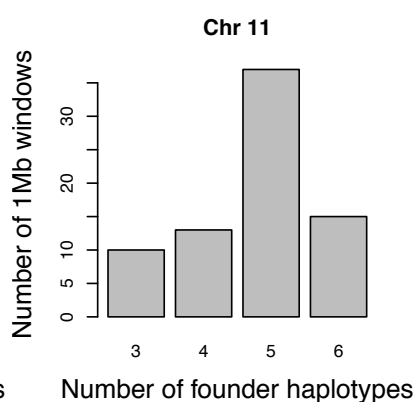
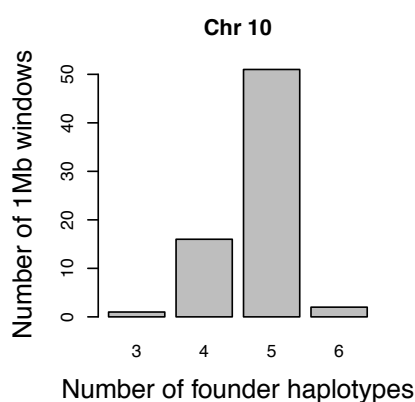
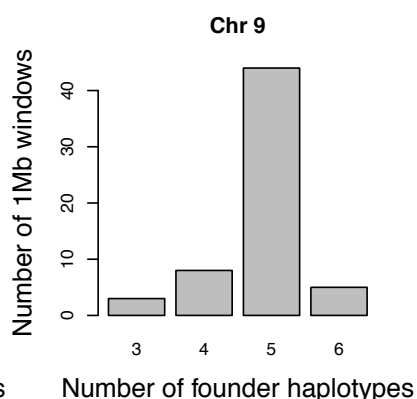
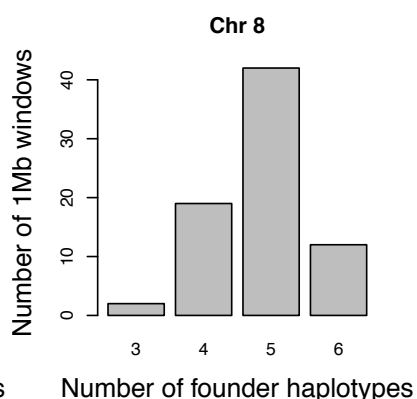
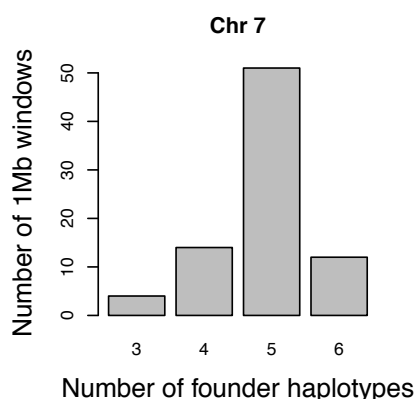
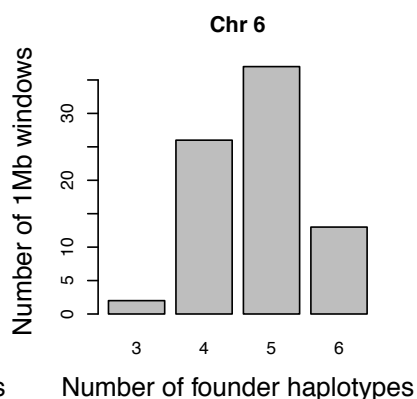
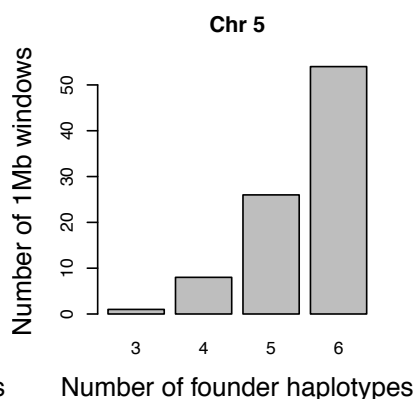
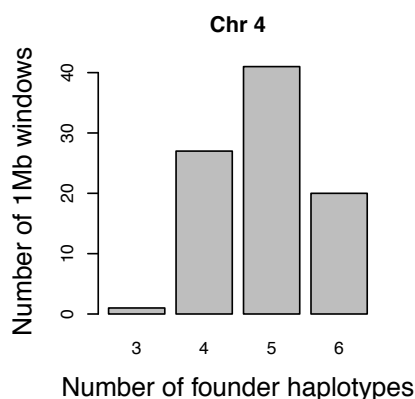
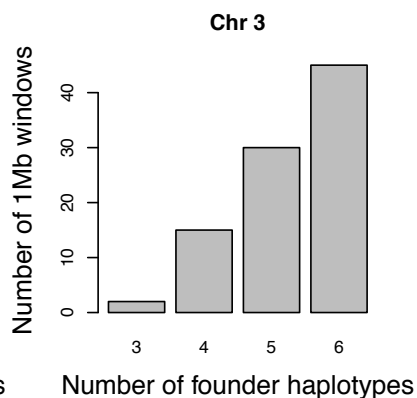
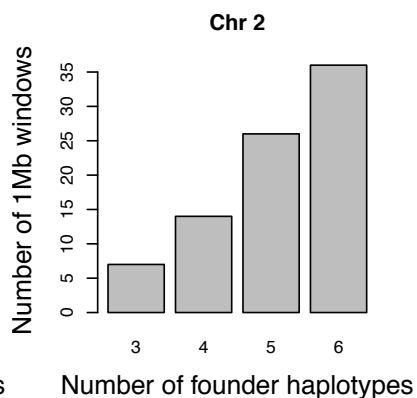
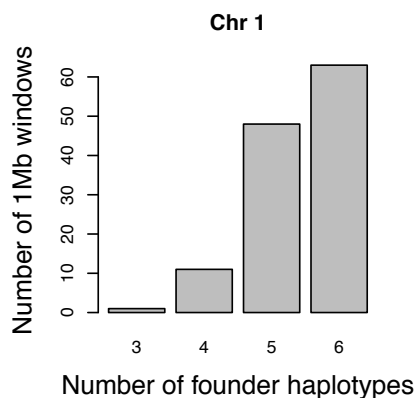
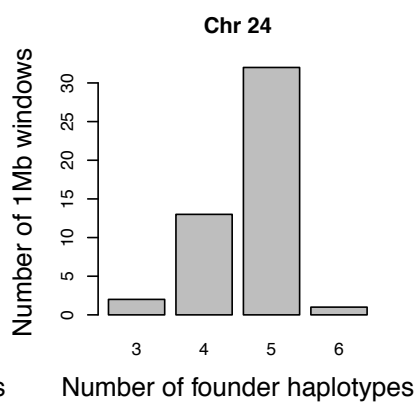
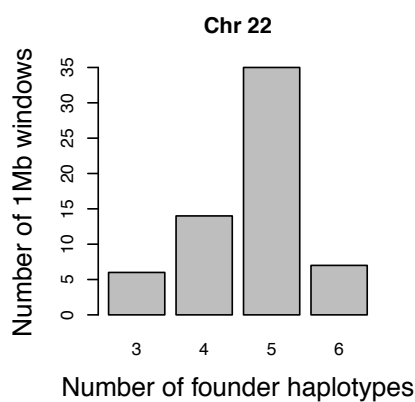
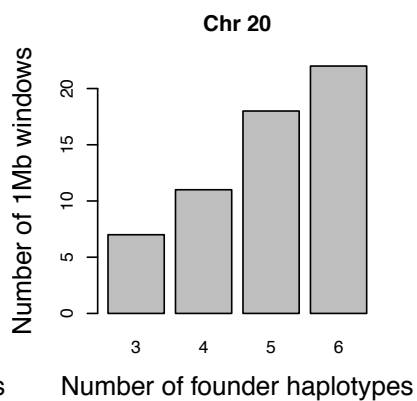
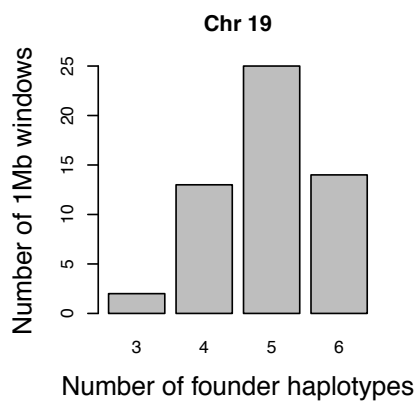
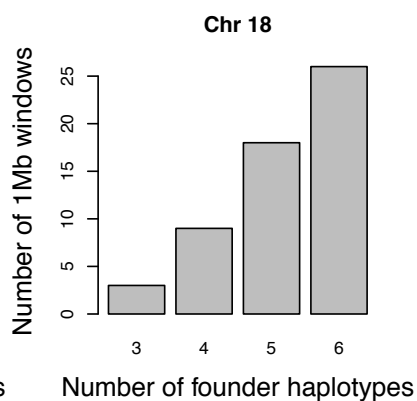
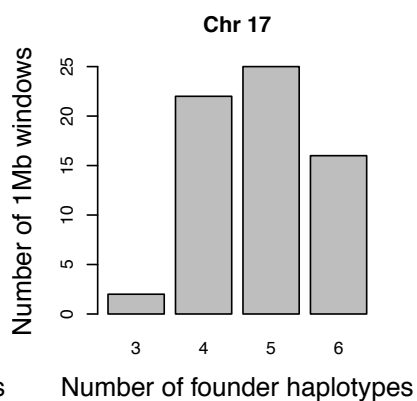
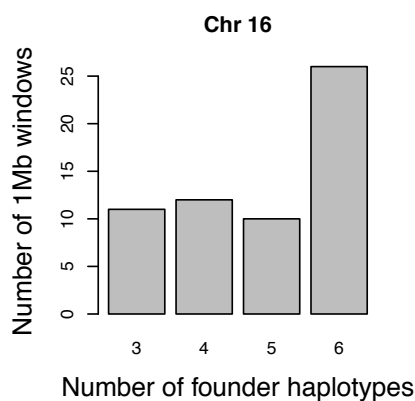
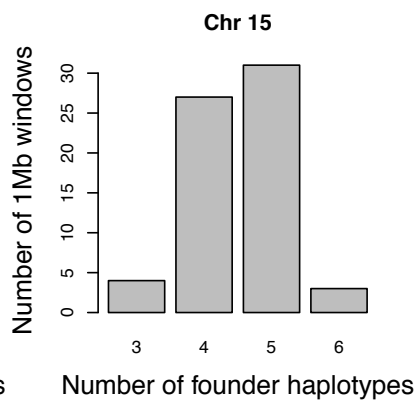
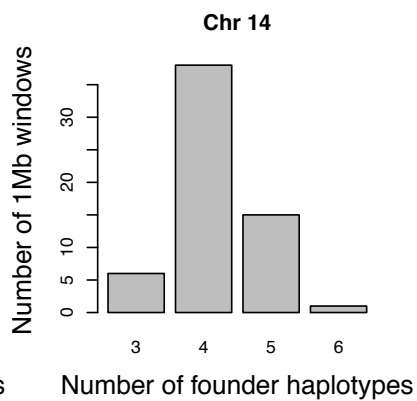
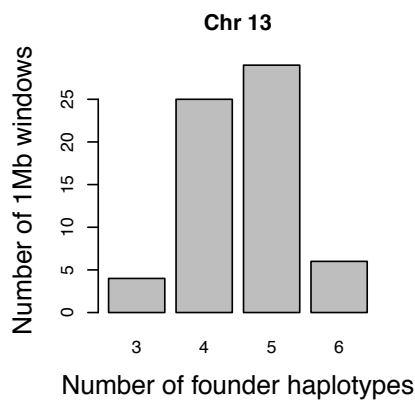
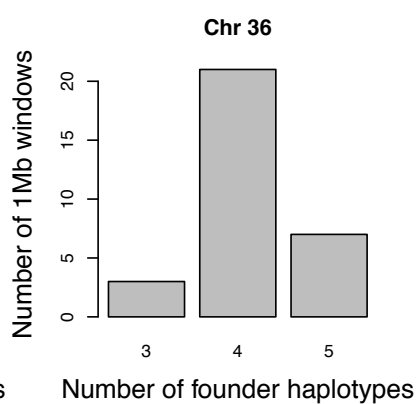
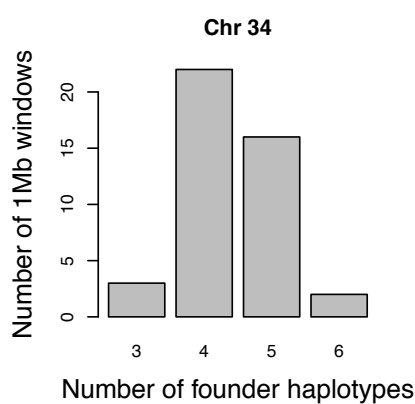
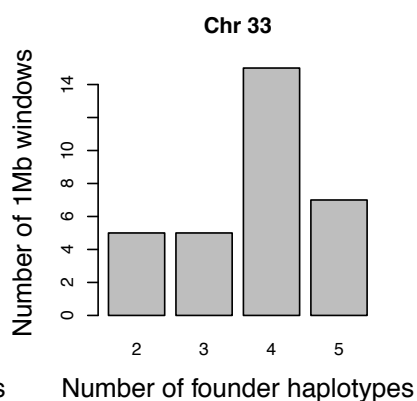
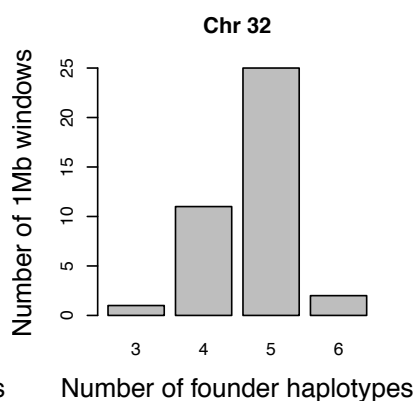
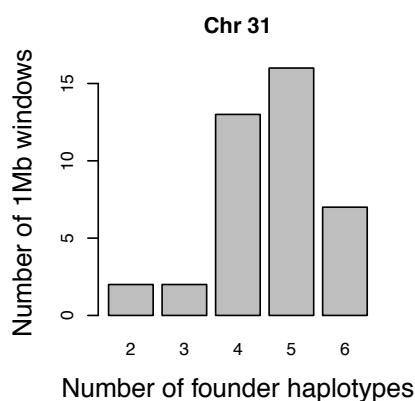
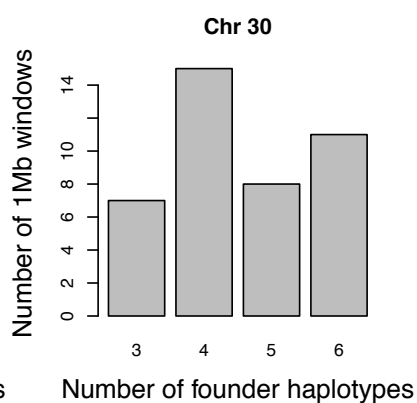
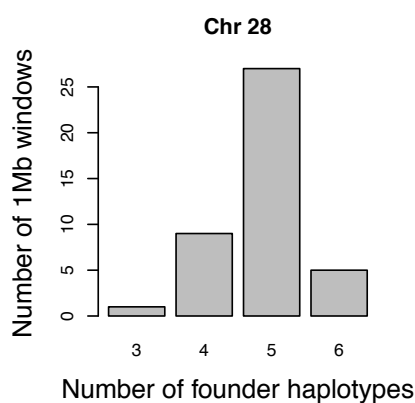
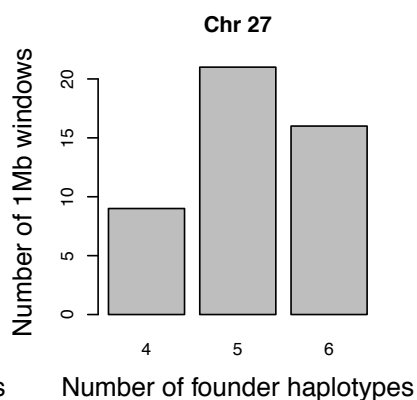
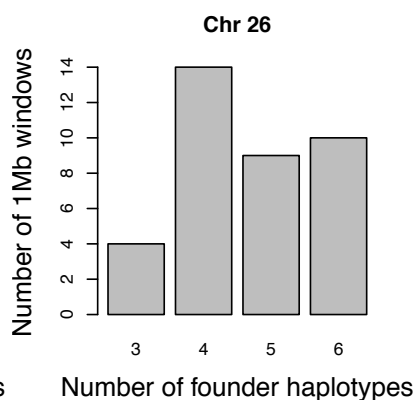
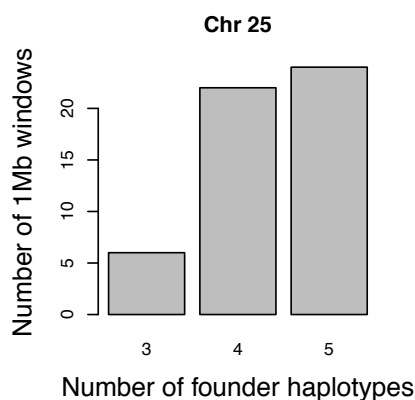
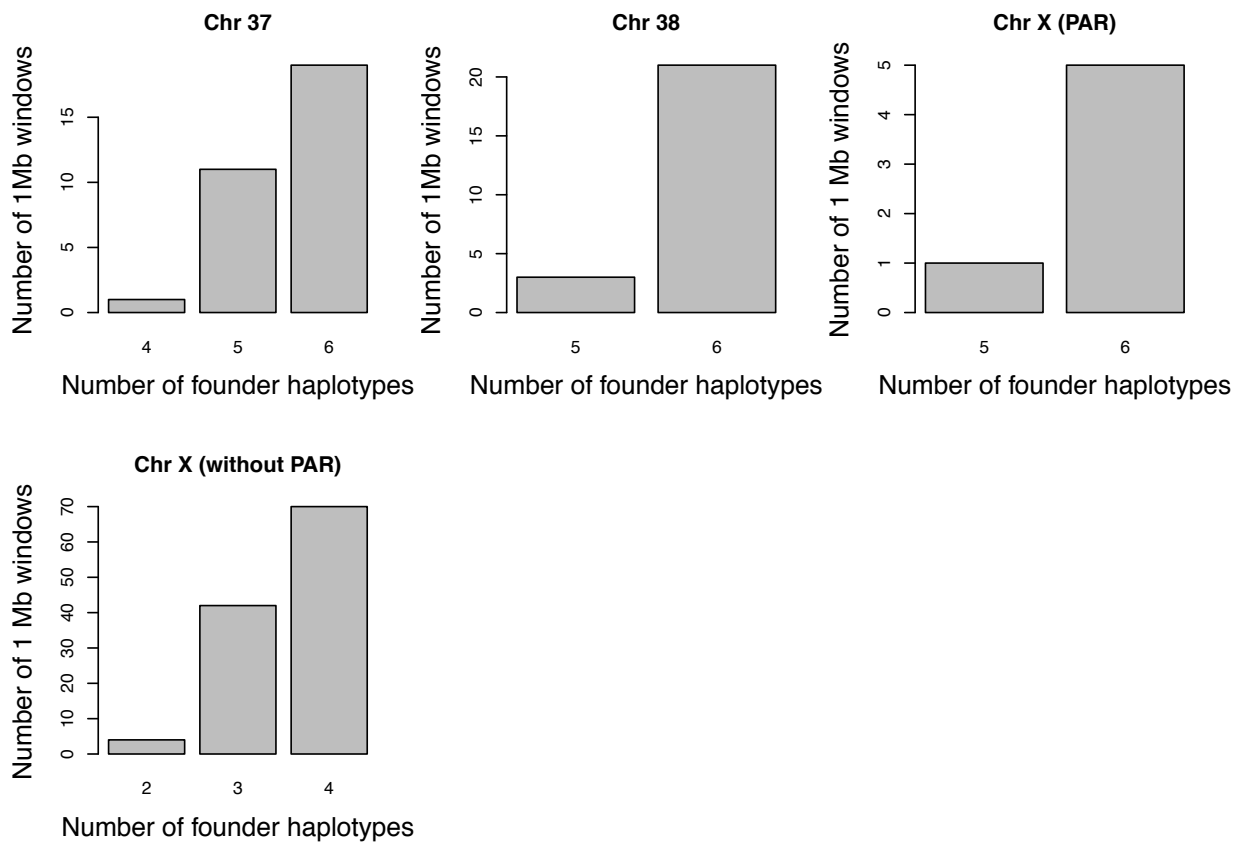










Supplemental Figure S3. Distribution of haplotype diversity (number of different haplotypes of the three founder wolves in 1 Mb windows) across the genome. Only founder haplotype diversity values present at least once on a particular chromosome were displayed.