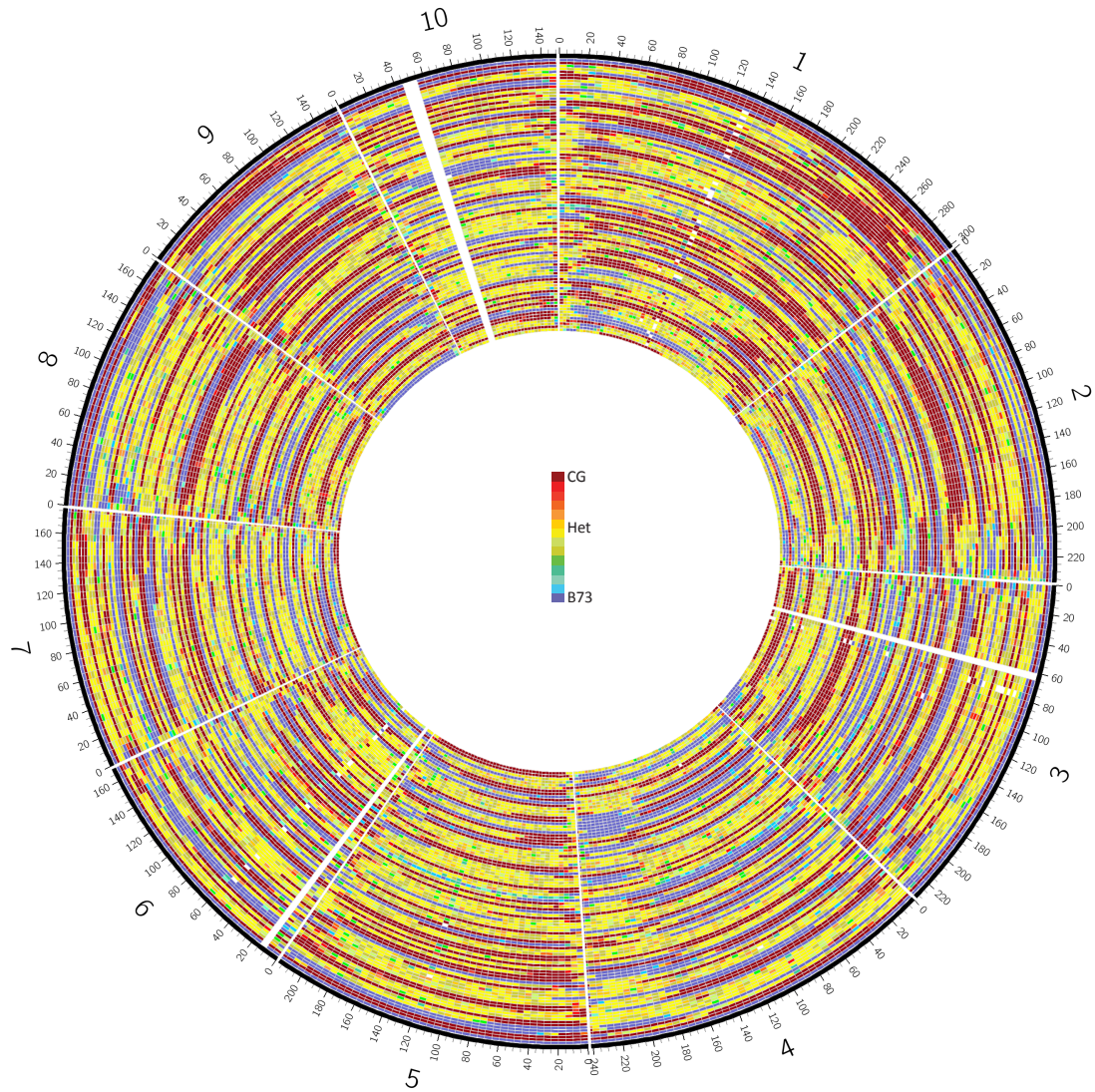




Additional File 6 Supplementary Figure 5: Raw GBS HincII dataset from an F₂ admixture population.

Post-filter, parental-phased variants from the B73 x CG HincII F₂ dataset were paced in 5 Mbp bins spanning the maize genome. Bin heatmaps indicate “mean genotype” value of variants within in the bin. Sample order is given, outermost to innermost, in Additional File 5, Supplementary Table 1.