

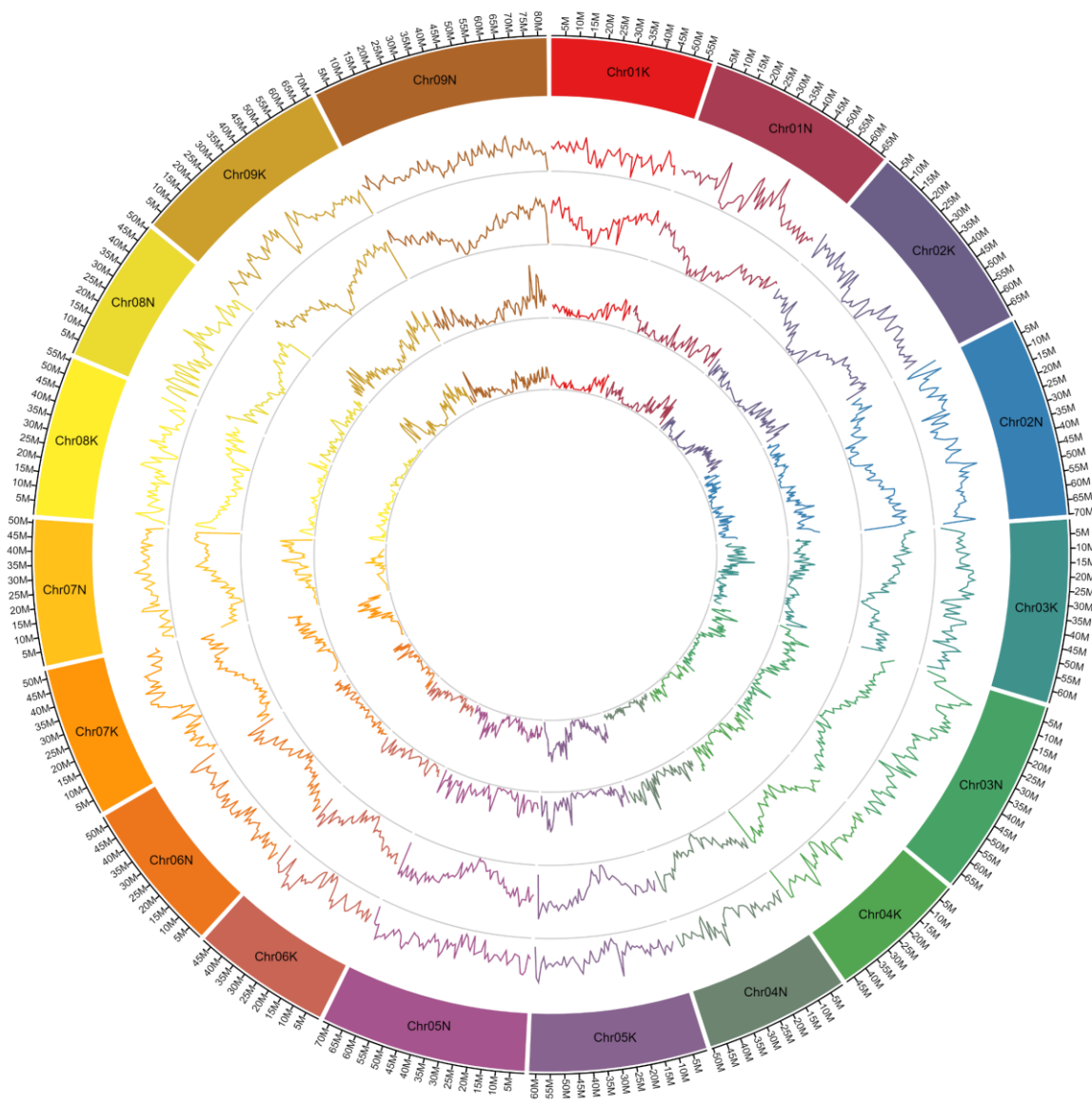


Figure S1. The distribution of, from outer to inner ring, NPs (SNPs and InDels) between AP13 and VS16 identified from GBS data (per Mb, height min=0, height max=3,600), annotated genes in AP13 assembly v5.1 (per Mb, height min=0, height max=160), and NP markers mapped in the Pop1 HH map and the Pop2 HH map (per Mb, height min=0, height max=30). The colored blocks indicate the chromosomes as represented in switchgrass AP13 assembly v5.1.