Zhang_review

Comments

- The data are easy to locate in the GitHub repository.
- The `.Rmd` file runs successfully in my local environment.
- I really like how you included separate plots for maize and teosinte, as well as a combined plot showing both — that helps with clarity.
- There's one issue in the *SNPs per Chromosome in Maize* and *SNPs per Chromosome in Teosinte* plots: The chromosomes on the x-axis are not sorted in ascending order (e.g., Chromosome 10 appears between Chromosome 1 and 2).
  This can be fixed by converting the chromosome labels to numeric using `as.numeric` before plotting.

Nice work overall — it's clear and well-organized.