PostGWAS Homework

Instructions:

1.) **Choose ONLY one from the following QTLs from this publication:**
- *qDTY 11. 2*
- *qDTF 11. 2*
- *qPH 11. 2*

2.) **Create a postGWAS analysis report on this QTL using one or a combination of the three tools discussed: RicePilaf, CropGalaxyy, or Rice SNPSeek.**

   You may opt to do a Genome Lift-over and gene list generation, incorporate epigenomic information, do enrichment analysis, etc. You may share your workflow, attach screenshots, and annotate your report step-by-step. The goal of this assignment is for you to familiarize yourself with the use of tools discussed in this module.

3.) **Submit your report on or before 5PM July 3, 2025 through this link:**
   https://forms.gle/EpUpWKuZn4Ue3WnE8