

Supplementary materials

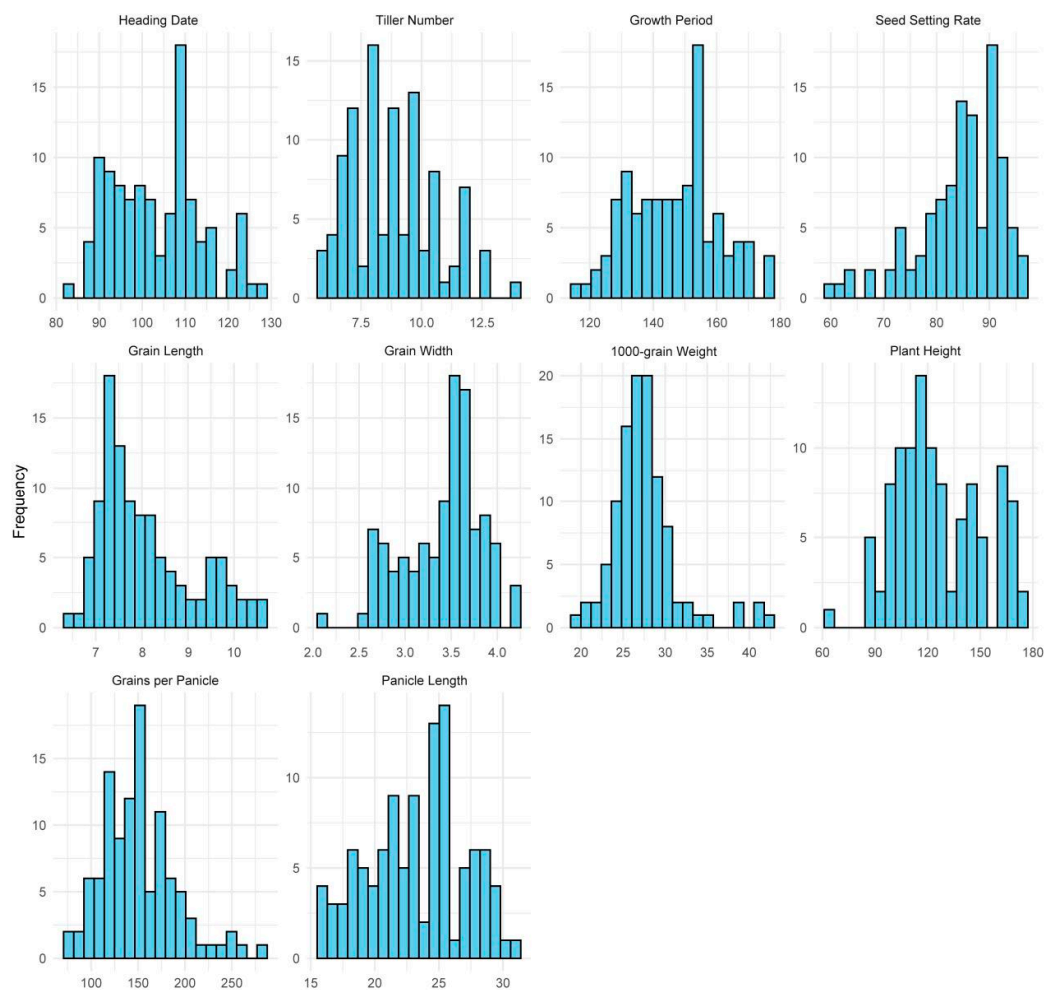


Figure S1. Frequency distribution histograms of ten agronomic traits. Each panel represents one trait.

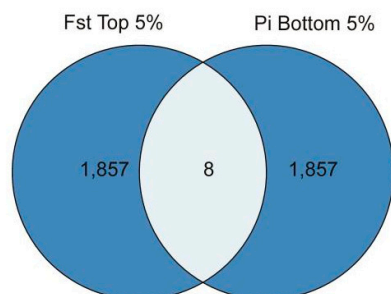


Figure S2. Venn diagram of the intersection between top 5% *Fst* windows and bottom 5% *Pi* windows.

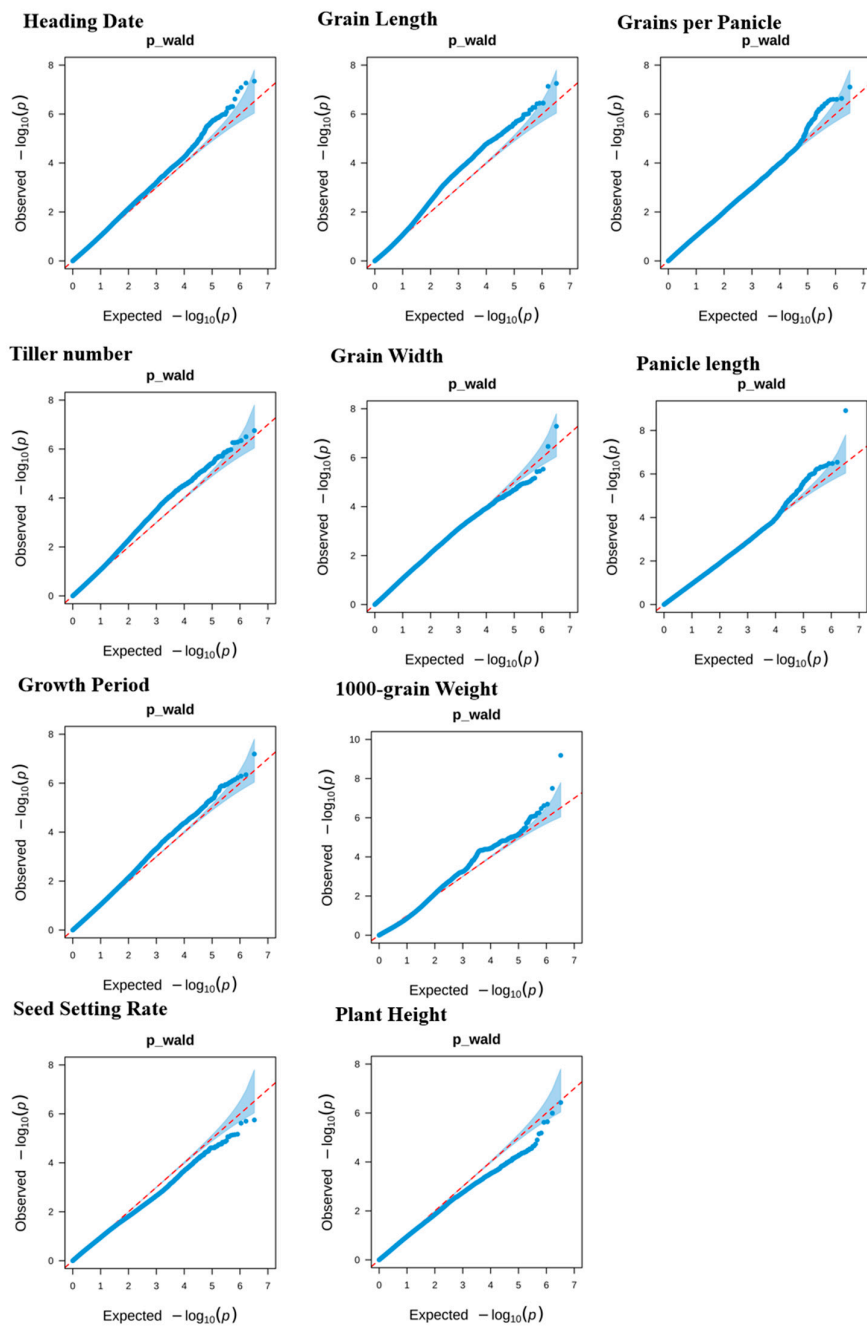


Figure S3. Quantile-quantile (QQ) plots of genome-wide association studies for ten agronomic traits in 109 Zhejiang rice accessions.

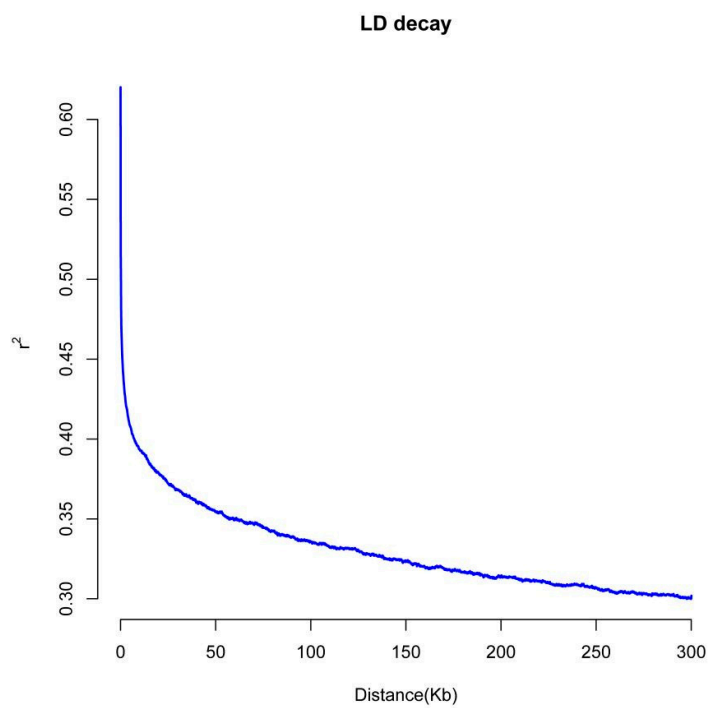


Figure S4. Genome-wide linkage disequilibrium (LD) decay in the 109 rice accessions. LD measured as r^2 plotted against physical distance (kb).