

Figure S1. SNP density map for rice chromosomes.

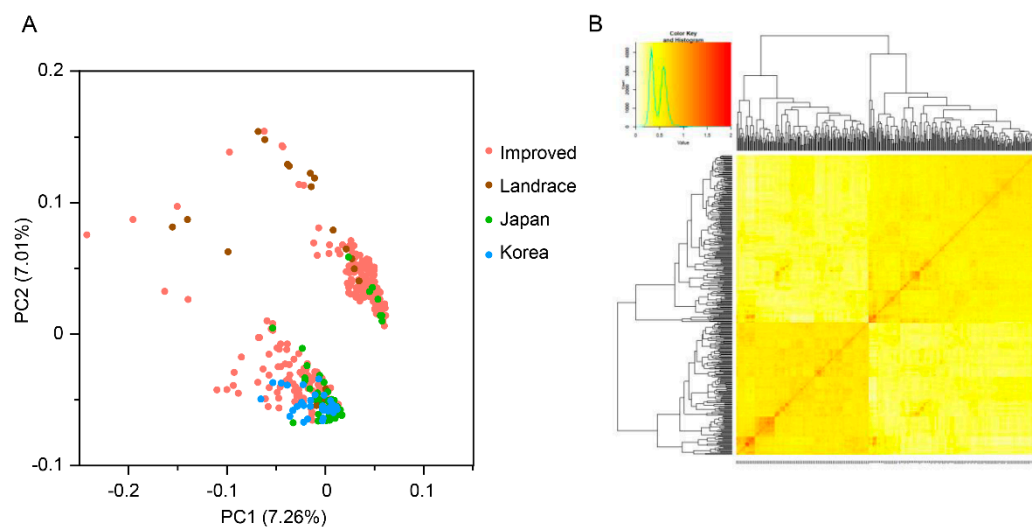


Figure S2. Population structure of 359 rice accessions.

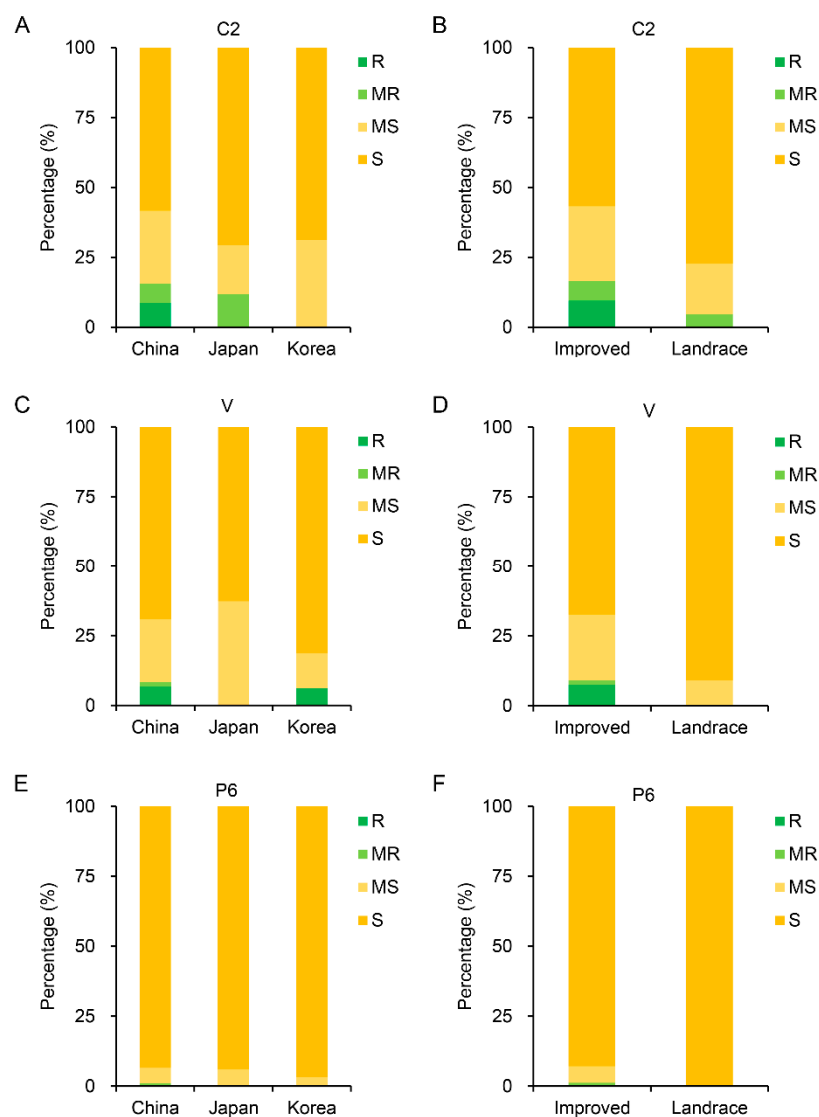


Figure S3. Boxplots for LLs of 359 accessions inoculated with three *Xoo* strains.

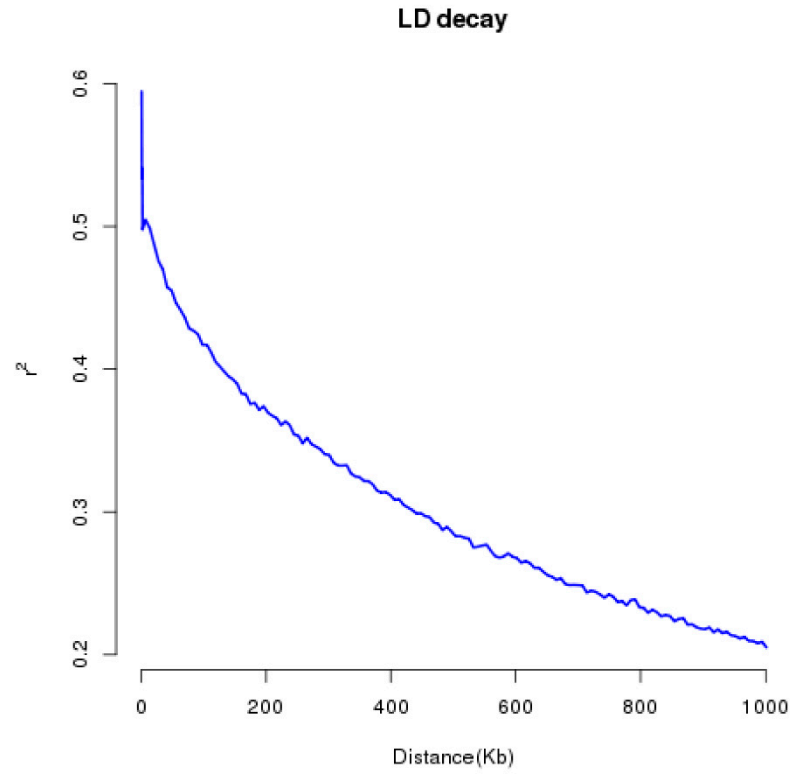


Figure S4. LD ( $r^2$ ) decay analysis of 359 *Japonica* rice accessions.

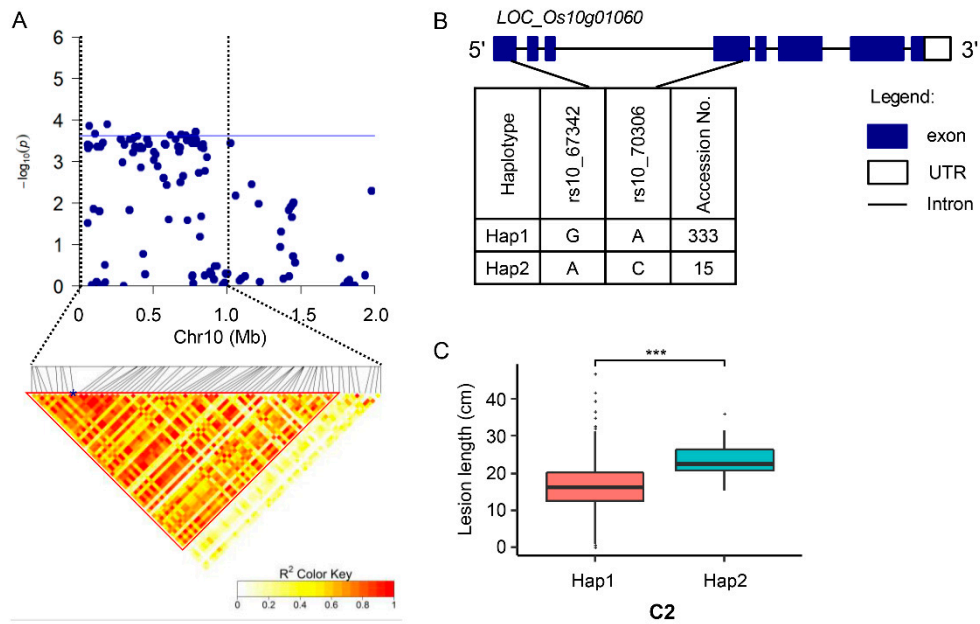


Figure S5. Analysis of associated region of *qBBC2-10.1* and haplotype analysis of *LOC\_Os10g01060*.

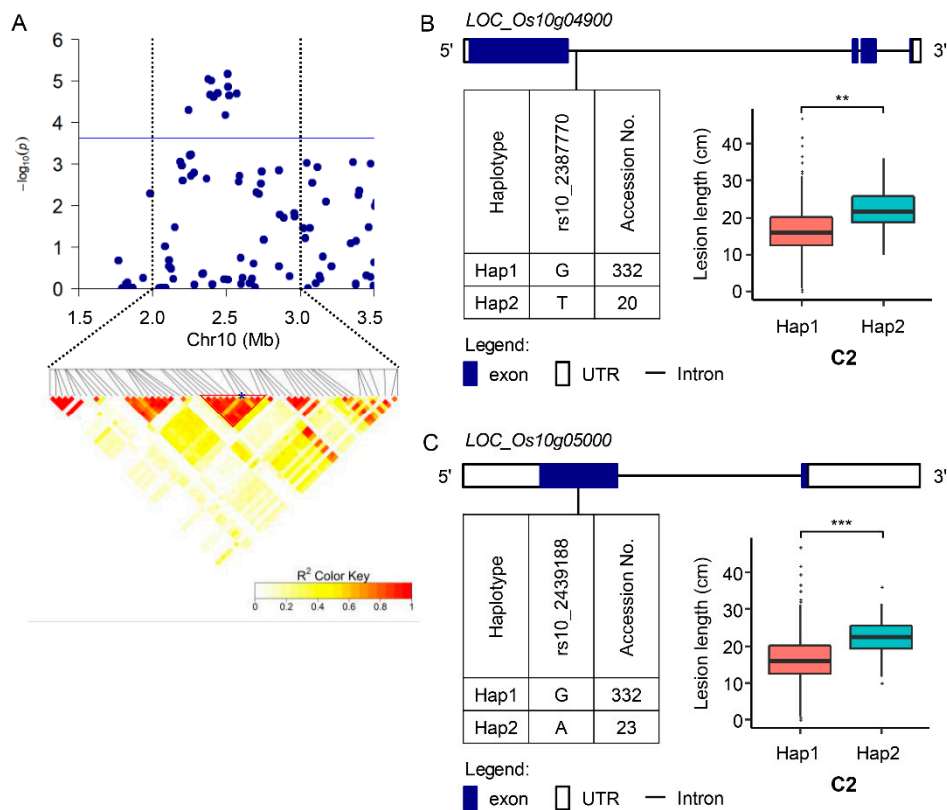


Figure S6. Analysis of associated region of *qBBC2-10.2* and haplotype analysis of *LOC\_Os10g04900* and *LOC\_Os10g05000*.

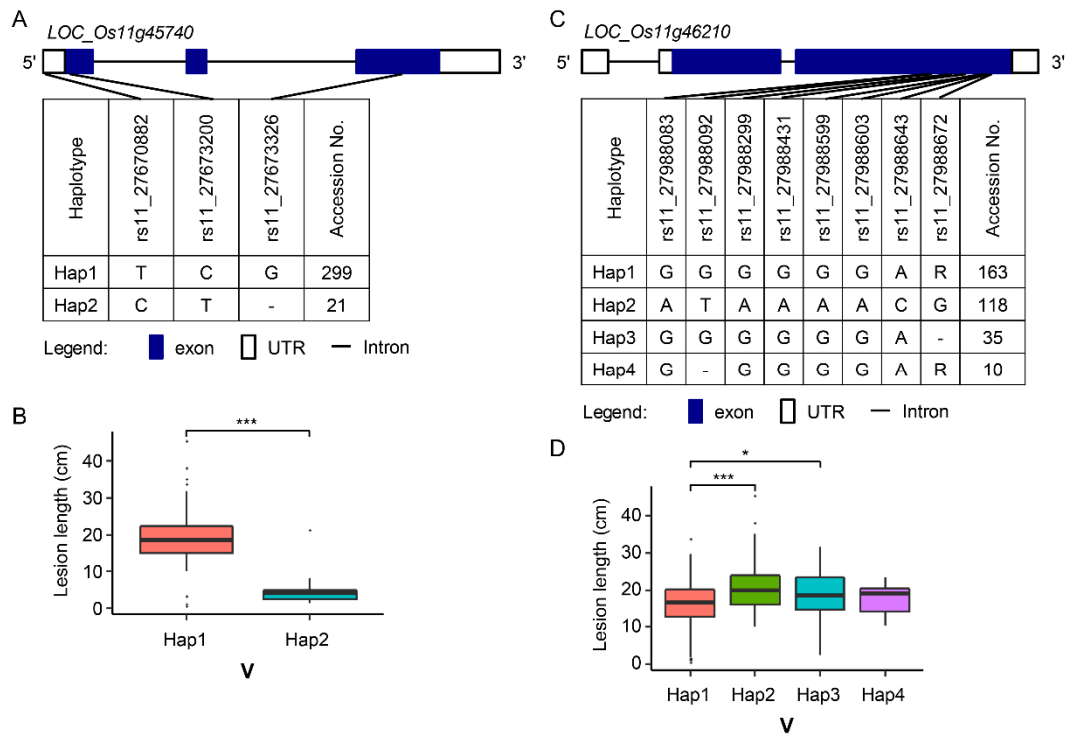


Figure S7. Gene structure and haplotype analysis of *LOC\_Os11g45740* and *LOC\_Os11g46210*.