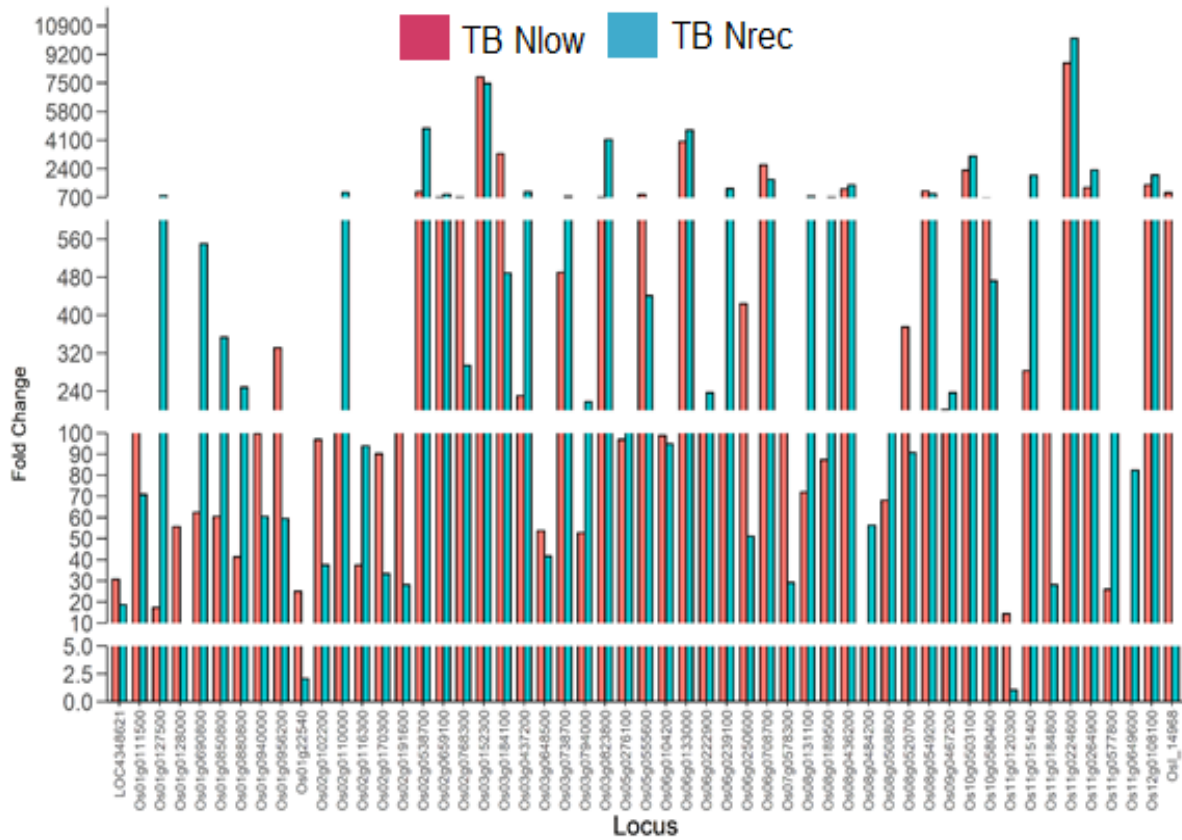
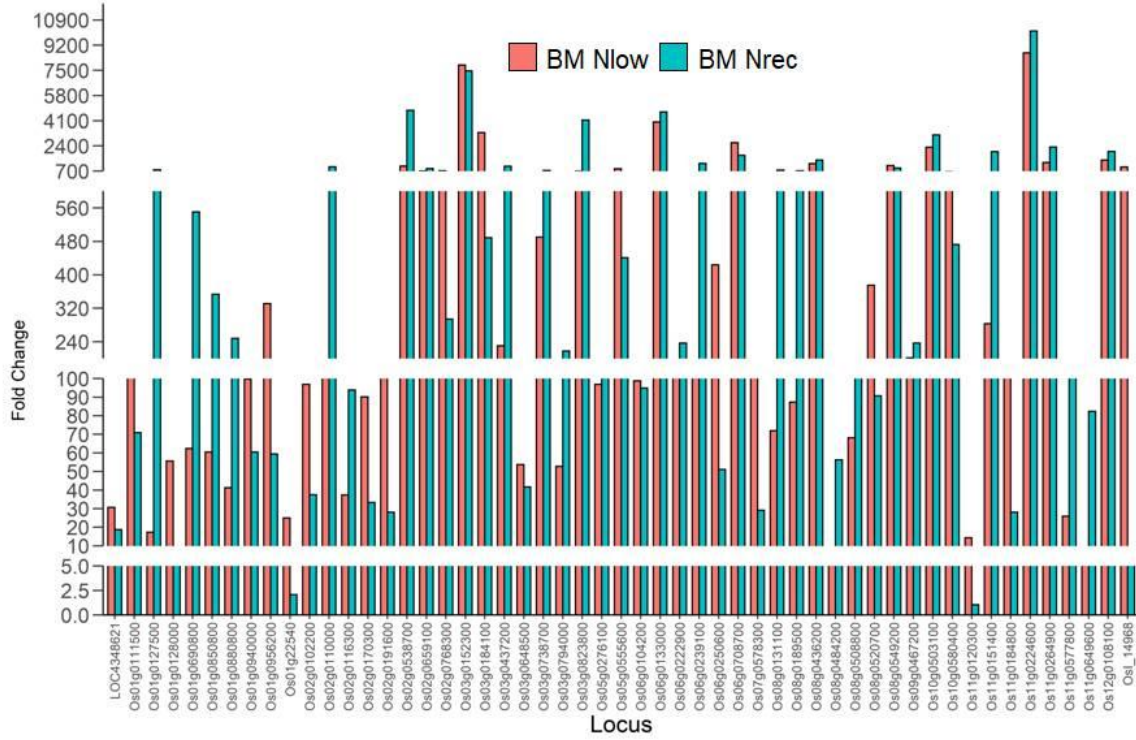


Fold changes in four genotypes using qPCR Based on 52 DETs of RNA-seq. N Recommended was considered as control and N low was taken as treated. Efficient genotypes: ThururBhog (TB), Basmati 370 (BM) and non-efficient - Suraksha (SK) and Kola Joha (KJ)

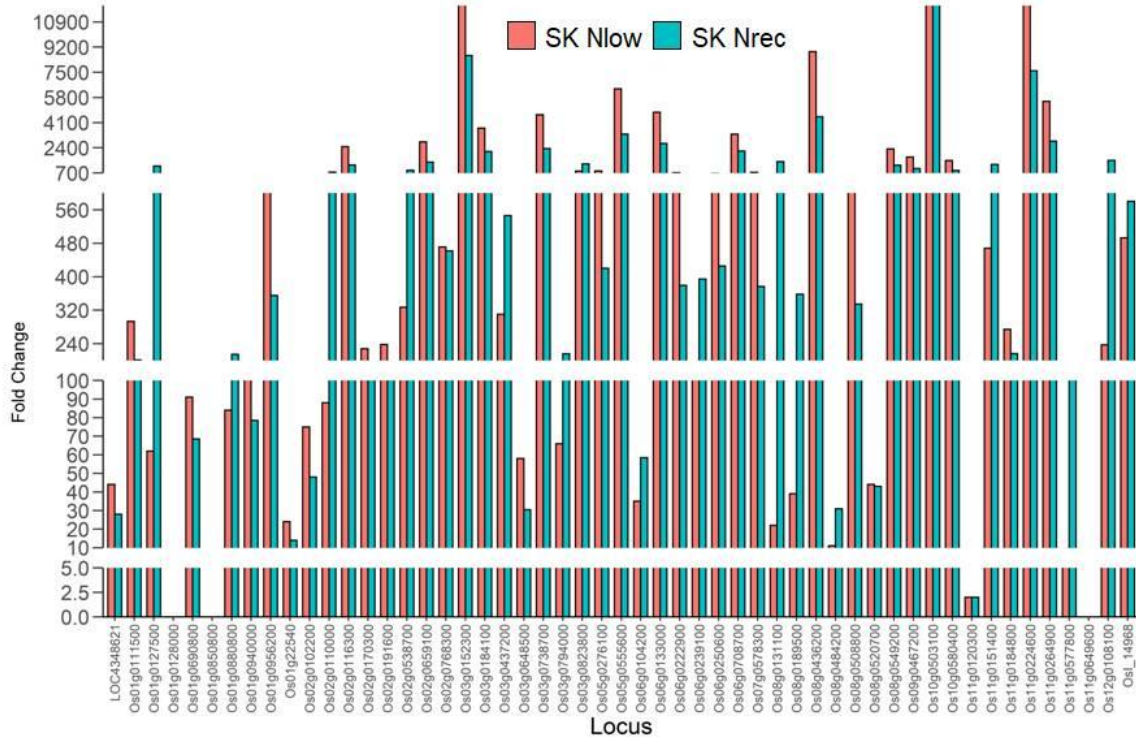
a. TB Nlow and TB Nrec



b. BM Nlow and BM Nrec



c. SK Nlow and SK Nrec



d. KJ Nlow and KJ Nrec

