

Figure S3

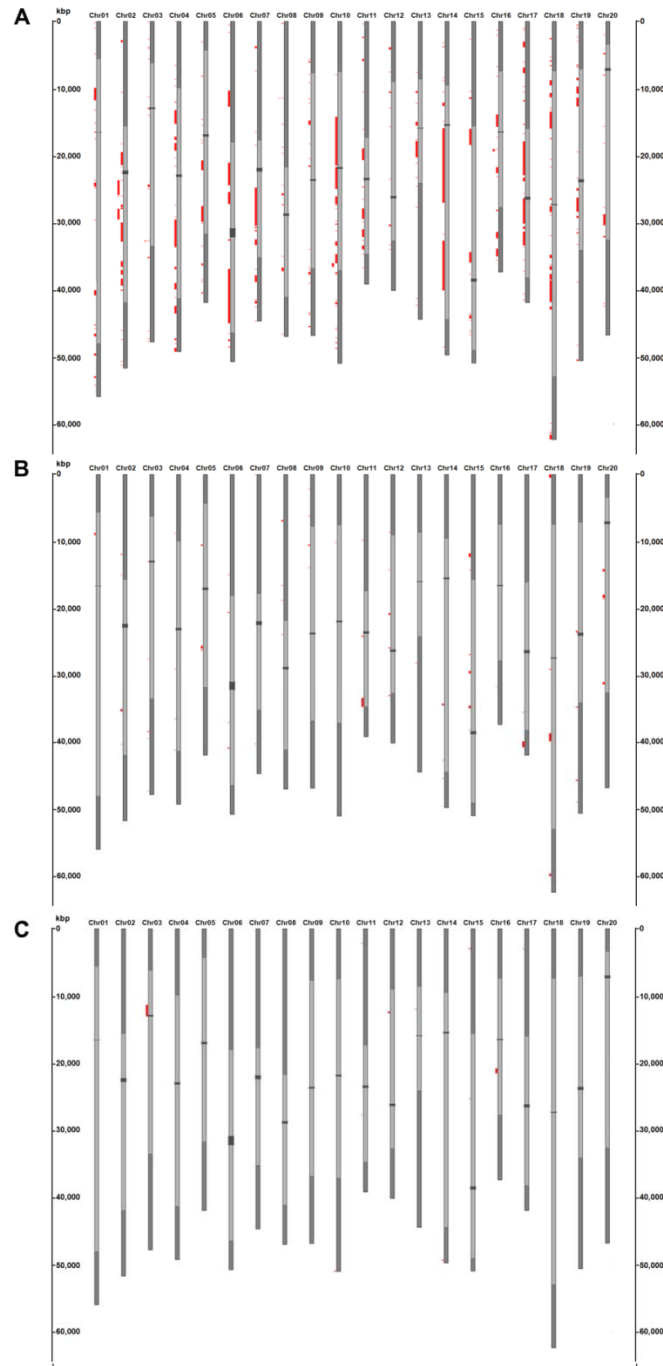


Figure S3 Genome-wide view of homozygous deletion coverage detected in fast neutron-irradiated soybean mutants for each selection category. Light gray lines indicate pericentromeric regions while euchromatic and centromeric regions are shown in dark gray. Homozygous deletion events are shown as red bars, detected in plants from the Forward Screen (A), No-Phenotype (B), and the Reverse Screen (C) classes.