

Description of Additional Supplementary Files

File Name: Supplementary Data 1

Description: Information for all soybean accessions used in this study.

File Name: Supplementary Data 2

Description: SNPs significantly associated with seed protein and oil content and 100-seed weight.

File Name: Supplementary Data 3

Description: TE genotypes of accessions used for biparental QTL mapping experiments.

File Name: Supplementary Data 4

Description: Differentially expressed genes identified by RNA-Seq.