Applying genotypic principal component scores as latent phenotypes in genome-wide and epistatic analyses of soybean agronomic traits.

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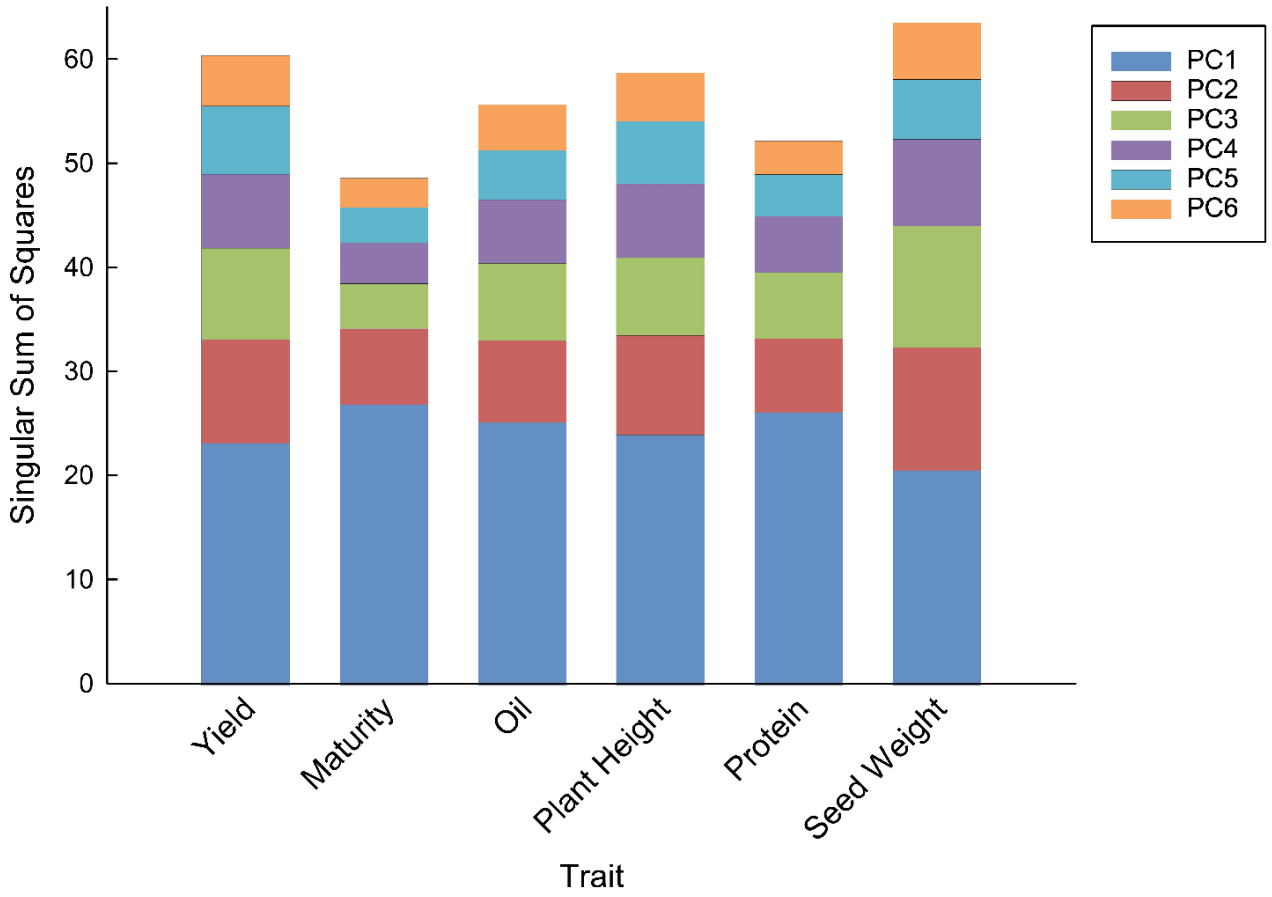

Supplementary Fig. S1 Bar graph representing singular sum of squares of gPCs derived from PCA of agronomic traits of soybean. The sum of squares is an indication of how much variability is accounted for by calculating each PC, the total reaching 100 when all variability is accounted for