Supplemental Fig. S2A  A graphical presentation of the output from STRUCTURE software indicating the presence of two separate sub-populations

Supplemental Fig. S2B  The STRUCTURE software plot of the chickpea association mapping population with K=2 clusters
Supplemental Fig. S2C PCA analysis of the chickpea population

Supplemental Fig. S2D The neighbor-joining phylogenetic tree (NJTree) based on Nei’s (1972) standard genetic distance between the 180 chickpea individuals studied