



Fig. S2. Distance clustering dendrogram of gene expression data for different populations. Clustering was performed using Euclidean distances and the average agglomerative method on mean expression values for each population (10,403 genes). *CO* and *CG* correspond to diploid species *C. orientalis* and *C. grandiflora*, respectively. The Asian, European and Middle Eastern populations of *C. bursa-pastoris* are called ASI, EUR and ME, and the two subgenomes are indicated with Co and Cg. F, L, and R stand for flower, leaf and root tissues, respectively. Bootstrap support was generated from 1000 replicates. Color boxes indicate different species and subgenomes.