S1 Figure. Phylogenetic relationship reconstructed for each subgenome separately. A and B: Density tree visualizing 1002 NJ trees reconstructed with 100 Kb sliding windows for the Cbp<sub>Cg</sub> and Cbp<sub>Co</sub> subgenomes, respectively. C and D: Whole genome NJ tree showing the absolute divergence between different populations of <i>C. bursa-pastoris</i> for the Cbp<sub>Cg</sub> and Cbp<sub>Co</sub> subgenomes, respectively. The root <i>N. paniculata</i> is not shown. ASI, EUR ME, CO, CG, CR indicate Asian, European and Middle Eastern populations of <i>C. bursa-pastoris</i>, <i>C. orientalis</i>, <i>C. grandiflora</i>, and <i>C. rubella</i>, respectively.