Supplemental Fig. S2. Unweighted pair group method with arithmetic mean (UPGMA) dendrogram of 52 sugarcane aphid (*Melanaphis sacchari*), one corn leaf aphid (CLA, *Rhopalosiphum maidis*), one pea aphid (Acyrthosiphon pisum Harris), and one yellow sugarcane aphid (YSCA, *Sipha flava*) samples analyzed with 14 simple sequence repeat markers. The purpose of this analysis was to determine if the clonal samples are identical to the predominant sugarcane aphid genotype seen on sorghum. The values for the x-axis are the DICE similarity coefficient values. Only bootstrap values (the numbers above the branches) of greater than 50% are shown. Information for each sample is listed in Supplemental Table S1. Sample names beginning with a ‘S’, ‘Su’, and ‘JG’ were collected on sorghum (Sorghum bicolor), sugarcane (Saccharum spp.), and Johnsongrass (S. halepense), respectively.