



Figure S1. Principal component (PC) analysis of all log transformed gene expression data from *Coffea canephora* cv. Conilon Clone CL153 (CL153) and *C. arabica* cv. Icatu (Icatu) plants submitted to three different water availability conditions: well-watered (WW), moderate water deficit (MWD), and severe water deficit (SWD). A, B, C, D, and E correspond to biological replications. Each treatment is represented by the colors depicted in Figure 1 and contains three biological replicates, depicted in Table S1. The percentage of variance is indicated in each axis.