Figure S5: Z Score Expression Graph for Modules 4, 7, 8, and 10
Z score expression values were calculated from the Fragments Per Kilobase of transcript per Million mapped reads for each co-expression module identified from weighted gene co-expression network analysis. Z scores were graphed by sample for all genes in Modules 4, 7, 8, and 10. Organs and samples with peak expression values in each module are indicated on the x-axis with the horizontal bars and text, respectively. 'DAP' refers to days after pollination and 'DAS' refers to days after sowing.