

Conserved in Arabidopsis and cotton

Figure S2. Venn diagram overlap of systemic hypoxia responsive homologues between *Arabidopsis* and cotton. Systemic responsive genes were from lists in Class I and II (Figure 1) for *Arabidopsis* and from closest homologues of leaves responsive genes in cotton (Christianson et al., 2010). These lists were filtered out at fold change > 2 and < 0.5 in expression with p values < 0.05 at any one time point from 1 to 12 h in shoots.