

**Supplemental Figure 4.** Hierarchical clustering analysis of the relative expression levels of transcription factors during grain development. (a) Hierarchical clustering analysis of the relative expression levels of transcription factors during grain development; (b) Gene ontology (GO) enrichment analysis of the TFs in foxtail millet grain development; (c) KEGG pathway classification top 20 of the TFs in foxtail millet grain development.