

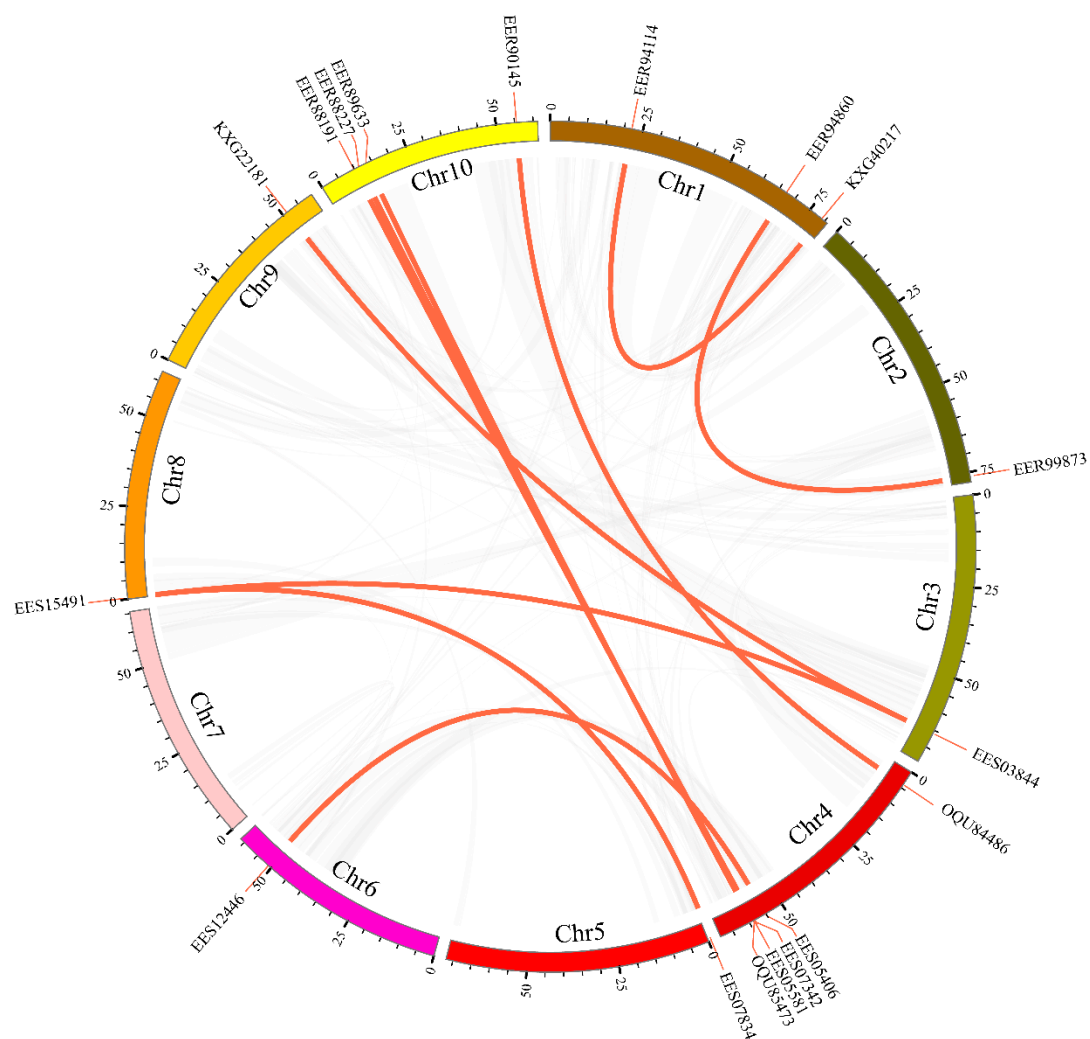
# Genome-wide identification and characterization of the *CCT* gene family in foxtail millet (*Setaria italica*) response to diurnal rhythm and abiotic stress

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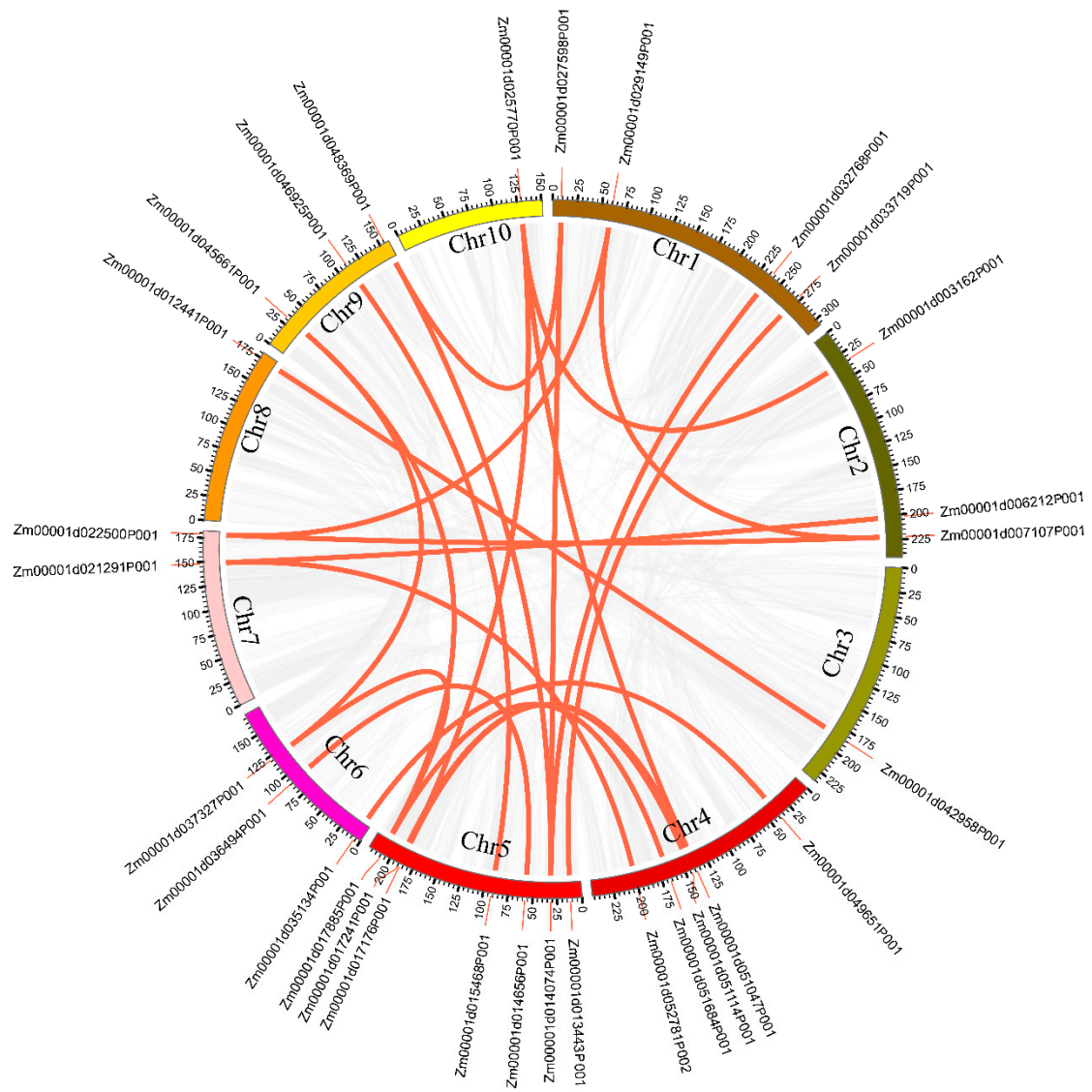
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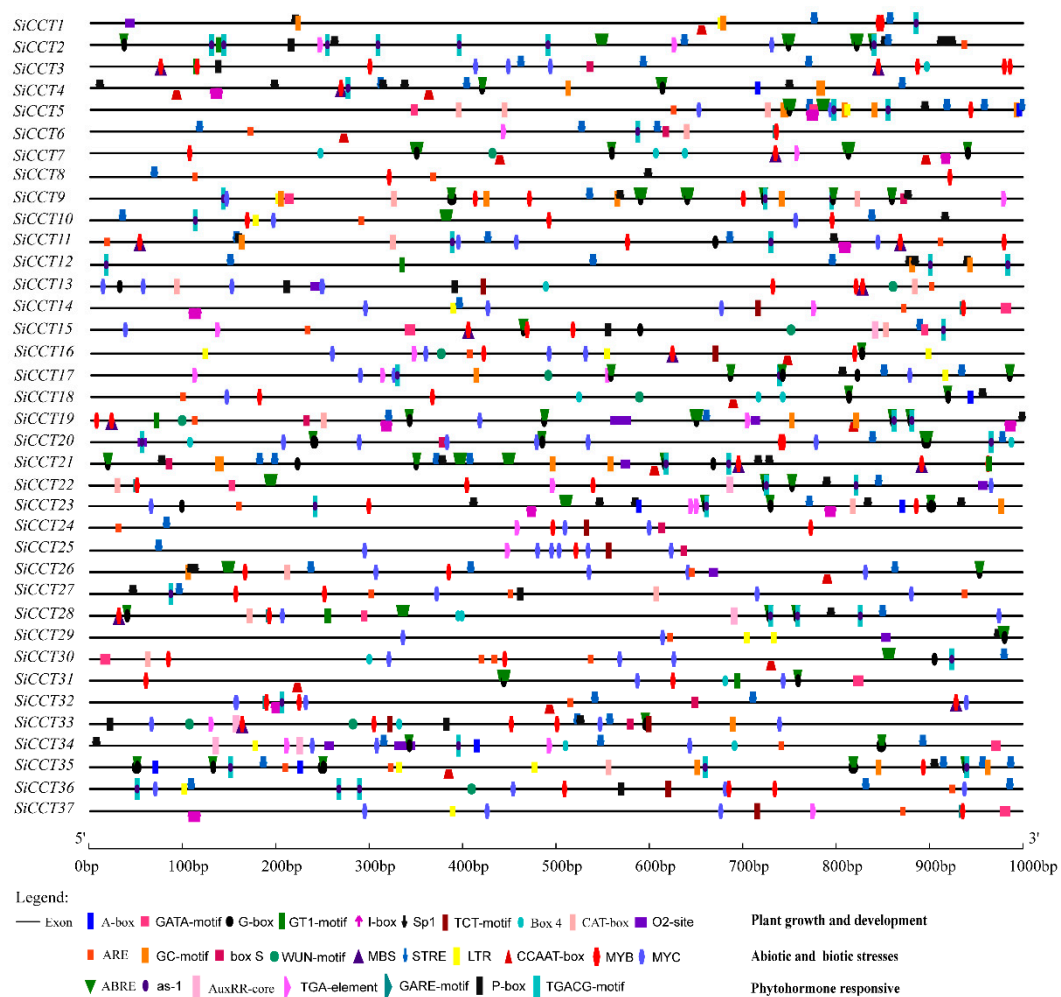
† These authors contributed equally to this work.



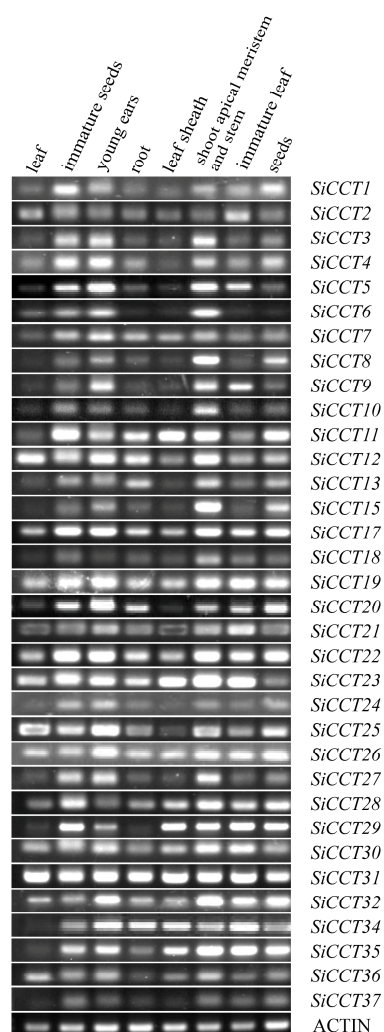
**Figure S1.** Interchromosomal relationships of sorghum CCT genes.



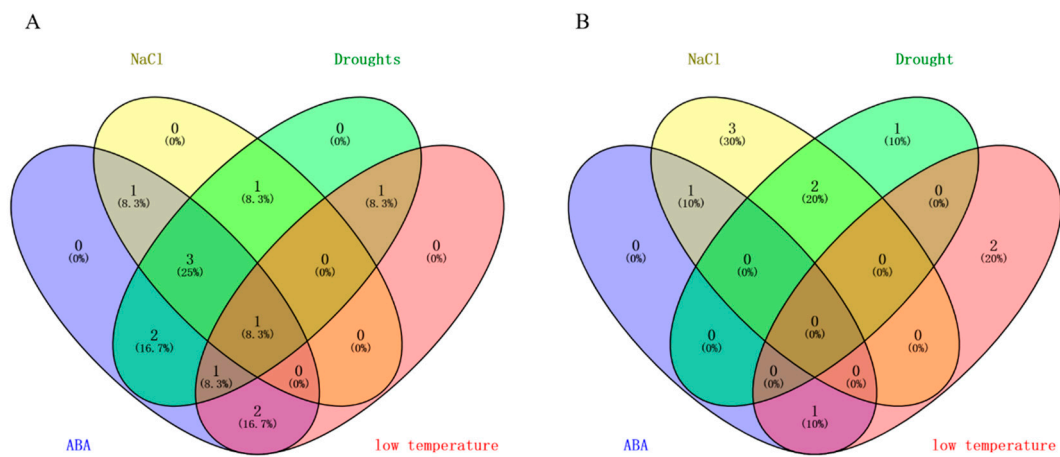
**Figure S2.** Interchromosomal relationships of maize CCT genes.



**Figure S3.** Distribution map of cis-acting elements on each gene.



**Figure S4.** Expression profiles of SiCCT genes in different tissues of millet. The site of expression is shown at the top, and the gene name is shown on the right.



**Figure S5.** The detailed numbers of simultaneously up-regulated (**A**) and down-regulated (**B**) *SiCCT* genes by ABA (abscisic acid), NaCl, drought, and low temperature.