

Supplementary Table S2 Positions of predicted *cis*-elements in the promoter sequences of five anthocyanin biosynthetic genes.

Promoter	Start site	End site	Predicted elements
<i>CmCHS promoter</i>	-163	-203	G-box (ABRE)
<i>CmCHI promoter</i>	-280	-320	MYC binding site
	-9	-49	G-box (ABRE)
<i>CmDFR promoter</i>	-399	-439	G-box (ABRE)
	-425	-465	G-box (ABRE)
	-265	-305	MYC binding site
	-369	-409	MYC binding site
	-148	-188	MYC binding site
<i>CmF3H promoter</i>	-195	-235	MYC binding site
	-324	-364	MYC binding site
	-520	-560	MYC binding site
	-715	-755	G-box (ABRE)
<i>CmANS promoter</i>	-209	-249	ABRE
	-831	-871	ABRE