

Table S1. Primer sequences used in this study

Experiment	Genes	Forward (5'-3')	Reverse (5'-3')	References
qRT-PCR	AcActin (Achn107181)	TGAGAGATTCCGTTGCCAGAAGT	TTCCTTACTCATGCGGTCTGCGAT	Petriccione, M. et al. Reference gene selection for normalization of RT-qPCR gene expression data from <i>Actinidia deliciosa</i> leaves infected with <i>Pseudomonas syringae</i> pv. <i>actinidiae</i> . <i>Scientific Reports</i> , 2015, 5, 16961.
	AcHY5	ATGAGATCAGGAGAGTGCCG	AATCTCTGAGCCGCTTGTG	Kim, S. et al. High ambient temperature represses anthocyanin biosynthesis through degradation of HY5. <i>Frontiers in Plant Science</i> , 2017, 8, 1787
	AcCHS	ACAGCTTGACCACCTAAATGGGCTTA	CAAAGTCCAATAAACATGCCAAT	
	AcCHI	GGAGGAGTTGACGGAATCCGTTG	AACTTTCGACGGCTTGGCCT	
	AcF3H	TGTACCCGCTCGCGATCCAG	ATTGCGTCTCTTCGCAAGTTCTTG	
	AcF3'H	GGTGGCGTATGCTCCGGAAGA	GTGGCCCGCACTCACCAAGTG	
	AcDFR	GTCGGAGAACGCTGATTGGATGGG	TGGCATTCCATTGGGAGGGGTT	Peng Y. et al. Differential regulation of the anthocyanin profile in purple kiwifruit (<i>Actinidia</i> species). <i>Horticulture Research</i> , 2019, 6(1): 3.
	AcLDOX	AACAGTGGTCGGAACTAGGG	TCCTCCGAAACCAAGTCCTC	
	AcF3GT	TAGCCAAGCAGAGATCCGCTTCC	CAAGAACCTTCTGGTAAGTACTGTTCGA	
	AcbHLH42	AAGGGGAAGGCAGGTGGATTCCG	GCACTGCAGCTAACCAACG	
	AccAO	ATCCGTTGGCACCTTGAA	GCGGCCCTCGTTATTG	
	AcRBCS	GTTACATGGGCCAAGAAAAT	TCAAACCTAACGGCAAGGGAT	
	AccBR	ATCGGATTCAAGTTATCAGGTG	GTTCGTCCCAGCATTGTTAT	
	AcGLUTR	CCGTTCCCACGATCAAGA	TGTCACTCCCGTCACATCTAAG	
	AcPPH	TGGAGGTGCTGTGGATGA	CAAAGATGTTGCTGGATTGG	Liu Y. et al. Expression differences of pigment structural genes and transcription factors explain flesh coloration in three contrasting kiwifruit cultivars. <i>Frontiers in Plant Science</i> , 2017, 8:1507.
	AcPAO	TCCCGACCCCATTCCAA	TCAAAGGCGTCAGGCAAC	
	AcSGR	ACAAGCGACTGTGAGATGCC	GGTTTAGGGTTGGTTTGG	
CDS cloning	<i>AcBHLH42</i>	ATGGCGGCTCCCCCTA	TAGGGTATGATTGGTGTATT	Wang L. et al. A MYB/bHLH complex regulates tissue-specific anthocyanin biosynthesis in the inner pericarp of red-centered kiwifruit <i>Actinidia chinensis</i> cv. <i>Hongyang</i> . <i>the Plant Journal</i> , 2019, 99: 359-378.
	<i>AcMYB10</i>	TAATAATCACTGGTAGTGTAGG	GCAGACTCAACTCCATCTAATATT	Peng Y. et al. Differential regulation of the anthocyanin profile in purple kiwifruit (<i>Actinidia</i> species). <i>Horticulture Research</i> , 2019, 6(1): 3.
Promoter cloning	<i>ProAcF3GT</i>	CTTGGACGAACCAAGTGAG	AATGTACCGACAAGGAGC	Yu, M. et al. Light-and temperature-induced expression of an R2R3-MYB gene regulates anthocyanin biosynthesis in red-fleshed kiwifruit. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5228.
	<i>ProAcLDOX</i>	GGGCATGATGGTTGGTTA	TACAGACGGAAGGCGACA	

>pAcF3GT-1058

TGGACGAACCAGTGAGTACATATATACATGTATATGGAGGAAGAGAATAGTTAATTAGAAACA
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> pAcLDOX-1500

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 TTCAGAACCTAATTGTCCTAAACTAAAATTAGTTACTTAAGTGAATTTGTCAGATTAGAGGGAC
 TAATGGTGAACCTACCTGAAATTCAAATTAGGATGGATTGAACATGAGGAAAGAAAGAGAAG
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 ACAATAATGATAATTAAAGAACAGAAAAAGTGTAAAGGGTGACATTGGATCTCTATCAGCA
 CGGTTAGACCACCTCGTGTGCCATCCAGTCAGTCCACTCCCCACGTGAGAAACTGATTCAA
 ACCACCTCTCACTTTAAGTTGATGCTAGCCCTAACCCCTACCGAGTCAGCACCTTCTGAAAAA
 ACCAACACATTGCAAGTGAAGCAACAAACACCTCTATAAAACTCCCAACACCTAATCACA
 CACATAATTGTCACAAATTCACACATAAGTACCGAATTACATTATAATTAGGTGTCGAAAAAAATG

Figure S1. The promoter sequences of *AcF3GT* and *AcLDOX*

Note: cis-acting elements are indicated in green.

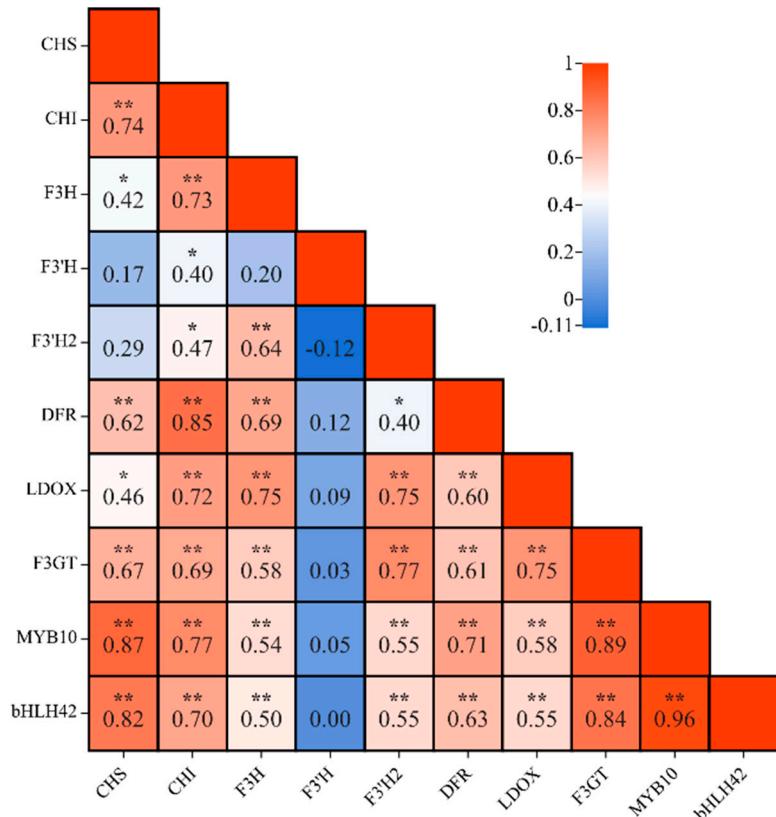


Figure S2. Correlation analysis of gene expression between anthocyanin synthesis genes and regulatory genes.

Note: * and ** represent statistical significance at $p \leq 0.05$ and $p \leq 0.01$ levels, respectively.

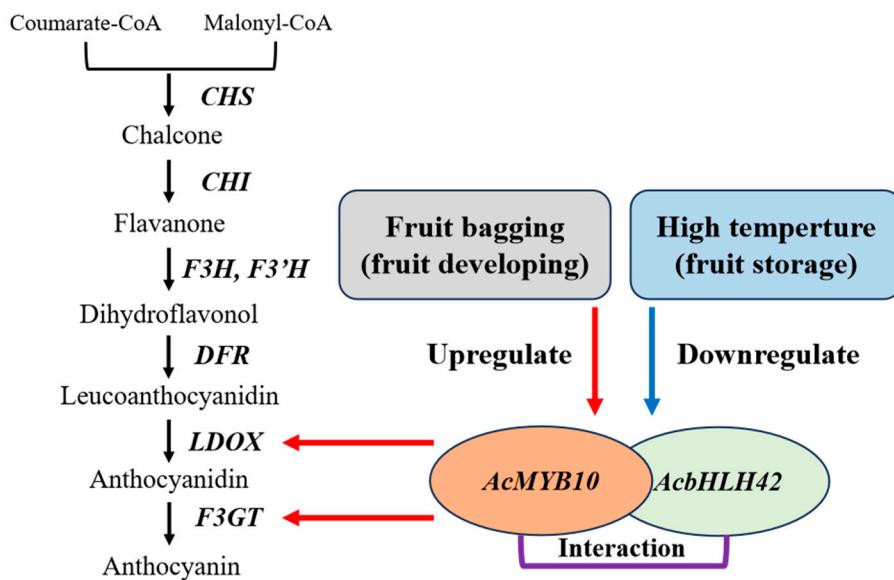


Figure S3. Regulation model of anthocyanin accumulation pathway in bagging and storage of kiwifruit.