

Supplementary Materials

Table S1. The primers were used for the quantitative real time PCR.

Gene	Primer	Sequence
<i>MaWRKY24</i>	QMaWRKY24-F	GAAGACTTGGAGTTGGCGG
	QMaWRKY24-R	TCATGAGAAGAAGCTGGAGA
<i>MaATG8f</i>	QMaATG8f-F	ATGGCGAAGAGTTCTCAA
	QMaATG8f-R	TTAACACAGATCAAATGTGT
<i>MaATG8g</i>	QMaATG8g-F	ATGGCGAAGAATTCTCAA
	QMaATG8g-R	CTACGCAGACCCAAATGTGT
<i>MaRPS2</i>	QMaRPS2-F	TAGGGATTCCGACGATTGTTT
	QMaRPS2-R	TAGCGTCATCATGGCTGGGA
<i>MaATG8f</i>	QChIP-MaATG8f-1F	TCCATCTTGTATTAGTATC
	QChIP-MaATG8f-1R	AAACAAAAGTCAAATCC
<i>MaATG8f</i>	QChIP-MaATG8f-2F	GCAGAAGTTGCTTGGATA
	QChIP-MaATG8f-2R	ATTAGTGGAGCTTAGTT
<i>MaATG8g</i>	QChIP-MaATG8g-1F	TTCGCTGGCAATAACGGACTT
	QChIP-MaATG8g-1R	TTCTTGCTCGCGGTGG
<i>MaACT</i>	QChIP-MaACT-F	AAGGTAGTCGACTTCGTCTC
	QChIP-MaACT-R	AGTTACCAAATGTTCGCTCGAT

Table S2. The primers were used for the vector construction.

Gene	Primer	Sequence	Construction
<i>MaWRK</i> <i>Y24</i>	MaWRKY 24-F	TCCCCCCC <u>GGG</u> ATGGAGAAAGGCAAAGCAAG	MaWRKY24-pEGAD
	MaWRKY 24-R	CGCG <u>GATC</u> TCATGAGAAGAAGCTGGAGA	
<i>MaWRK</i> <i>Y24</i>	MaWRKY 24-F	CGCG <u>GATC</u> CATGGAGAAAGGCAAAGCAAG	MaWRKY24-pET28a
	MaWRKY 24-R	AAGGAAAAAA <u>AGCGGCCG</u> CTGAGAAGAAGCTGGAG AAGT	
<i>MaATG8</i> <i>f</i>	pMaATG8 f-F	CGGG <u>GTACC</u> ATTTGTAGTGTGTAAT	pMaATG8f-pGreen II 0800-LUC
	pMaATG8 f-R	TCCCCCCC <u>GGG</u> AAGGATTGATAATATAAG	
<i>MaATG8</i> <i>g</i>	pMaATG8 g-F	CGGG <u>GTACC</u> AACCATGCGGAGGTAGTCG	pMaATG8g-pGreen II 0800-LUC
	pMaATG8 g-R	TCCCCCCC <u>GGG</u> CTCGACCAAGATCGCG	
<i>W-box-</i> <i>mini35S</i>	W-box- mini35S-F	TCCCCCCC <u>GGG</u> TTGACCTGACTTGACCTGACTTTG ACCTTGACGATATCTCCACTGACGTAAGGG	pW-box-mini35S- pGreen II 0800-LUC
	W-box- mini35S-R	CGCG <u>GATC</u> GTGTTCTCTCAAATGAAATG	
<i>mW-box-</i> <i>mini35S</i>	mW-box- mini35S-F	TCCCCCCC <u>GGG</u> TTCTGGTTCTGTTCTGGTTCTGTTCT GGTTCTGGATATCTCCACTGACGTAAGGG	pmW-box-mini35S- pGreen II 0800-LUC
	mW-box- mini35S-R	CGCG <u>GATC</u> GTGTTCTCTCAAATGAAATG	

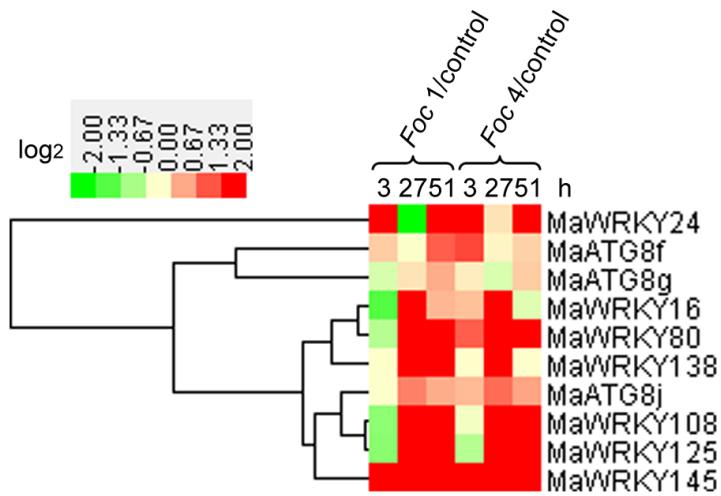


Figure S1. Cluster analysis of correlation between *MaWRKYs* and *MaATG8s* expression. The transcriptomic data were downloaded from Li et al. (2013). For the assay, banana roots were inoculated by *Foc* 1 or *Foc* 4, respectively. The heatmap of gene expression was constructed using CLUSTER program and Java Treeview.