

## 苹果 *MdAFS* 基因亚细胞定位表达载体的构建及分析

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**摘要:** *Malus domestica*  
Borkh cv. White Winter Pearmain) PCR  $\alpha$ - (*AFS*)  
*Rubisco* (*Ru*) (*Mit*) (*Pla*) plantCARE SignalP4.1Server  
*Ru* TATAbox CAATbox

*Ru* *AFS*  
PCR

**关键词:** ; *MdAFS* ; ;  
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## Construction and Analysis of Apple *MdAFS* Gene Expression Vector into Different Subcellular Compartments

DING Rui-rui<sup>1</sup>, WANG Lu<sup>1</sup>, WANG Ru-ru<sup>1</sup>, LIU Yu<sup>1</sup>, ZHANG Yuan-hu<sup>1\*</sup>, CHENG Ni-ni<sup>2\*</sup>

1. College of Life Sciences/Shandong Agricultural University, Tai'an 271018, China  
2. College of Life Sciences/Linyi University, Linyi 276000, China

**Abstract:** To study the change of terpene synthase subcellular localization influenced terpenoids metabolism and plant development, we used the White winter pearmain peels, Chrysanthemum morifolium, Yeast and Arabidopsis thalianas as experimental materials and cloned the strong promoter of  $\alpha$ -Farnesene synthase, *Rubisco*, transit peptide of mitochondria(*mit*) and transit peptide of chloroplast(*pla*) by RT-PCR. Then the sequence of promoter and transit peptide were predicted by PlantCARE and SignalP4.1Server. The results showed that *Ru* had typical characteristics of strong promoter elements(TATAbox, CAATbox and hormone response elements) and both of *mit* and *pla* maybe have the function of transit peptide. We have successfully constructed the  $\alpha$ -Farnesene synthase gene expression vector in different subcellular localization driven by *rubisco* strong promoter, and transformed them into tobacco. Quantitative real-time PCR showed that the gene had a higher expression and was tissue-specific in transgenic tobacco.

**Keywords:** Apple; *MdAFS* gene; subcellular; expression

[1,2]

C10

C15

[3]

[4]

/

[5]

$\beta$ -

[6]

$\beta$ -

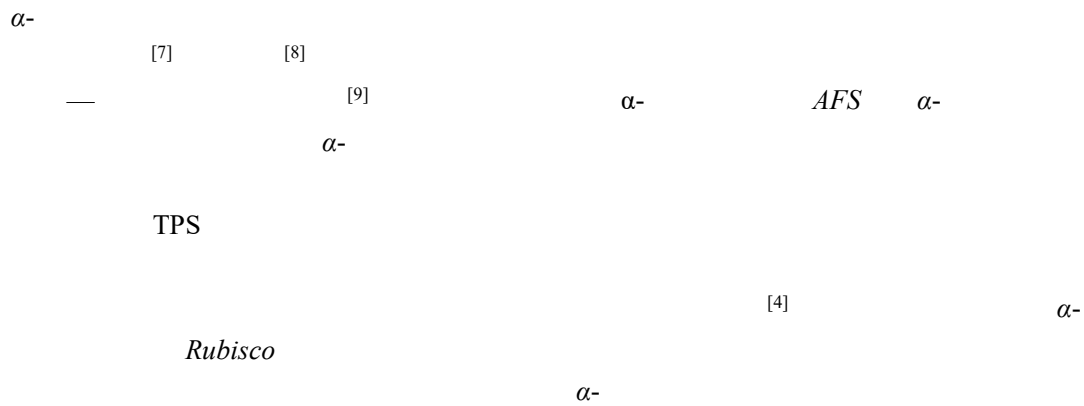
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作者简介: (1990-), , . E-mail:ruidingc@163.com

\*通讯作者: Author for correspondence. E-mail:yhzhang9@163.com; chengnn2002@163.com



## 1 材料与amp;方法

### 1.1 材料及试剂

(*Malus domestica* Borkh. cv. white winter pearmain  
*Chrysanthemum morifolium* NC89  
*Columbia* LBA4404  
*Arabidopsis thaliana* Sall  
*E. coli* DH5α  
 pMD18-T  
 TaqDNA  
 DNAMarker  
 T4-DNA  
 MS  
 Fermentas  
 AR

### 1.2 材料处理

T1 NC89 MS 25 °C 2  
 50~60% 145 μmol·m<sup>-2</sup>·s<sup>-1</sup> 16 h /8 h  
 -80 °C

### 1.3 基因的提取

Ultrapure RNAKit RNA CWBIO RNA cDNA  
 -20 °C PCR CTAB  
 DNA

### 1.4 引物设计

PCR 1

表 1 基因克隆及引物序列  
 Table 1 Gene cloning and sequence of primers

Primer name	Genbank accession number	Primer sequence(5'-3')	Size of fragment /bp
RuFS1	AY563622	GCCCATGGAATTCAGAGTTCAC	1780
RuFS2		GCGTCGACTAGTTTACAAGAGG	
Ru1	AY163904.1	CTGAAGCTTAAAGCTTAGACAAACACCCCTTG	1100
Ru2		TCCCCCGGGGTTTCTGATACTTAGGAGGAATGG	
MitP1	X01418.1	GGATCCATGTTGTCACACTACGTCAATCTATAA	87
MitP2		CCATGGGTTTTTGCTGAAGCAGATATCTA	
PlasmidP1	CP002685.1	GGATCCATGGCGATGTCTTTCTCAGGAGCTG	156
PlasmidP2		GCCATGGCGTAGATCAACGACTTCTTGCG	

### 1.5 构建载体

1 min 15 s 1 35 PCR 72 °C 10 min 94 °C 5 min 20 μL 94 °C 1 min 55 °C 1 min 72 °C  
 °C pMD18-T 3-10:1 4  
*Pla/Mit*

PCR DNA LB Amp+  
 pMD18-T+*Pla/Mit+AFS* pBI121  
 35S *Rubisco* pBI121+*Ru* pBI121+*Ru*  
 pMD18-T+*Pla/Mit+AFS*  
 pBI121+*Ru+Pla/Mit+AFS*

### 1.6 农杆菌介导烟草转化

pBI121+*Ru+Pla/Mit+AFS* 100  
 mg·L<sup>-1</sup> Kan PCR

### 1.7 实时荧光定量 PCR 分析

T<sub>2</sub> 6 ( Ultrapure  
 RNAKit RNA CWBIO RNA cDNA -20 °C

18SF1 5'-TTCCTAGTAAGCGCGAGTCATCAGC-3' 18SR1  
 5'-GCGACGGGCGGTGTGT-3' *AFS* PaFSF 5'-AAAGCGACAATCTCGGCACAA-3'  
 PaFSR 5'-GCGCGAAATGGTGGTTCTCTAAT-3' qPCR

## 2 结果与分析

### 2.1 *AFS*、*Ru* 强启动子、*Mit* 和 *Pla* 的克隆

Ultrapure RNAKit RNA CWBIO RNA  
 CTAB DNA  
 RuFS1 RuFS2 Ru1 Ru2 PlasmidP1 PlasmidP2 MitP1 MitP2  
 PCR *AFS* *Ru* *Pla* *Mit* PCR  
 1 PCR pMD-18T  
 PCR *AFS* *Ru* *Mit* *Pla*

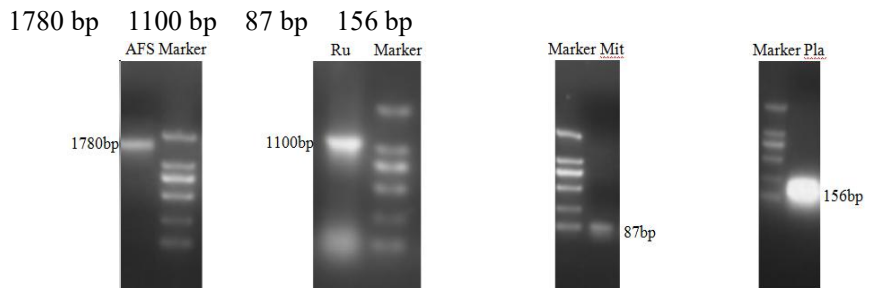


图 1 基因克隆电泳图

Fig.1 Electrophoresis results of genes cloning

Marker DL2000 ladder A *AFS* ;B *Ru* C *Mit* D *Pla*

Marker: DL2000 ladder; A: Amplified fragment of *AFS*; B: Amplified fragment of *Ru*; C: Amplified fragment of *Mit*; D: Amplified fragment of *Pla*

### 2.2 *Ru*、*Mit* 和 *Pla* 序列预测分析

DNAman GeneBank PlantCARE  
*Ru* *AFS* *Pla* *Mit* *Ru*

GeneBank PlantCARE  
*Ru* TATAbox CAATbox  
 ( 2 SignalP4.1Server(<http://www.cbs.dtu.dk/services/SignalP/>)  
 ( 3

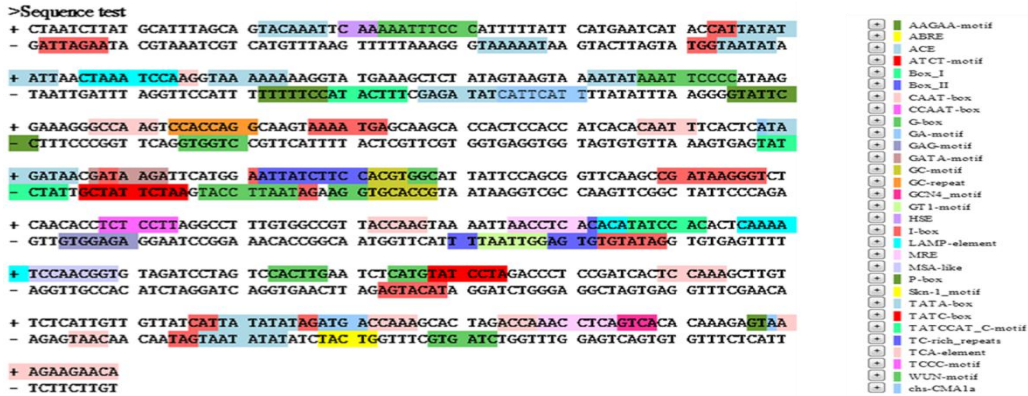


图 2 *Ru* 启动子核苷酸序列预测  
 Fig.2 Deduced nucleotide sequences of *Ru* promoter

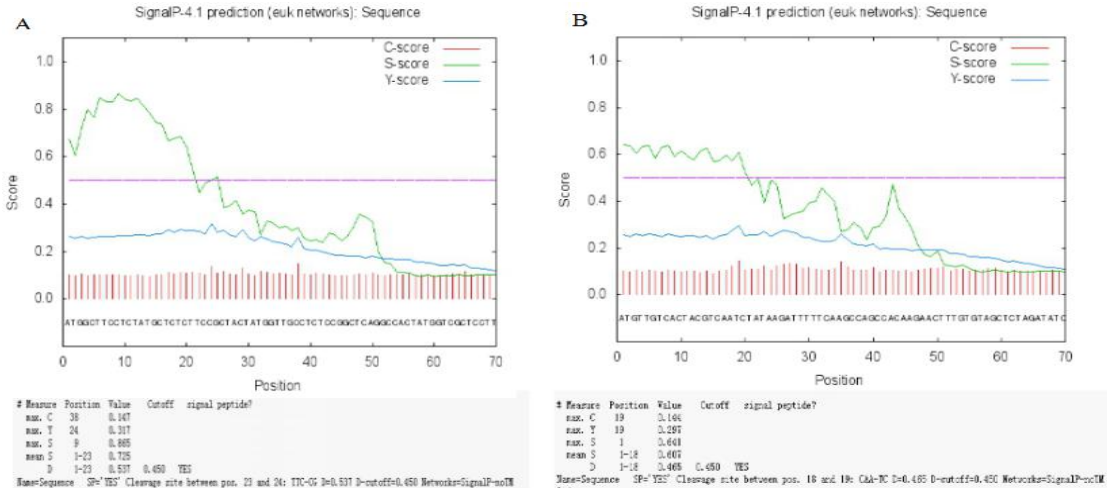


图 3 线粒体和叶绿体转运肽序列预测  
 Fig.3 Deduced sequences of Mitochondria transit peptide and chloroplast signal peptide

A: Targeting peptide of chloroplast; B: Targeting peptide of mitochondria

2.3 真核表达载体构建

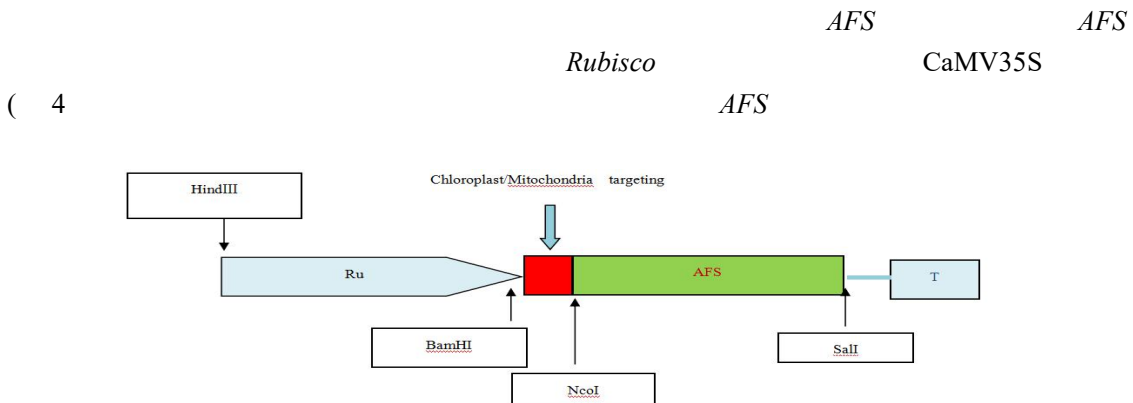
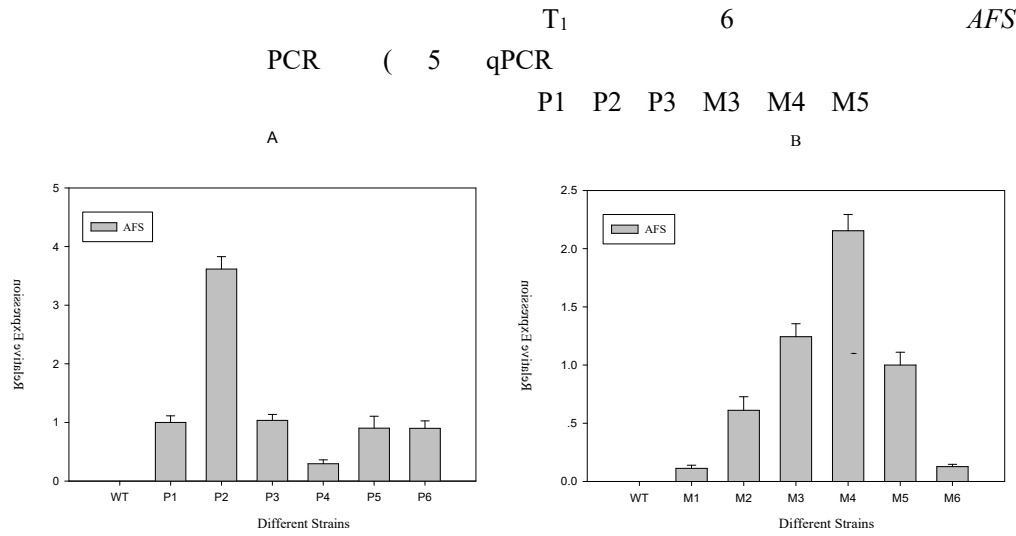


图 4 表达载体的构建图  
 Fig.4 Construction of expression vector

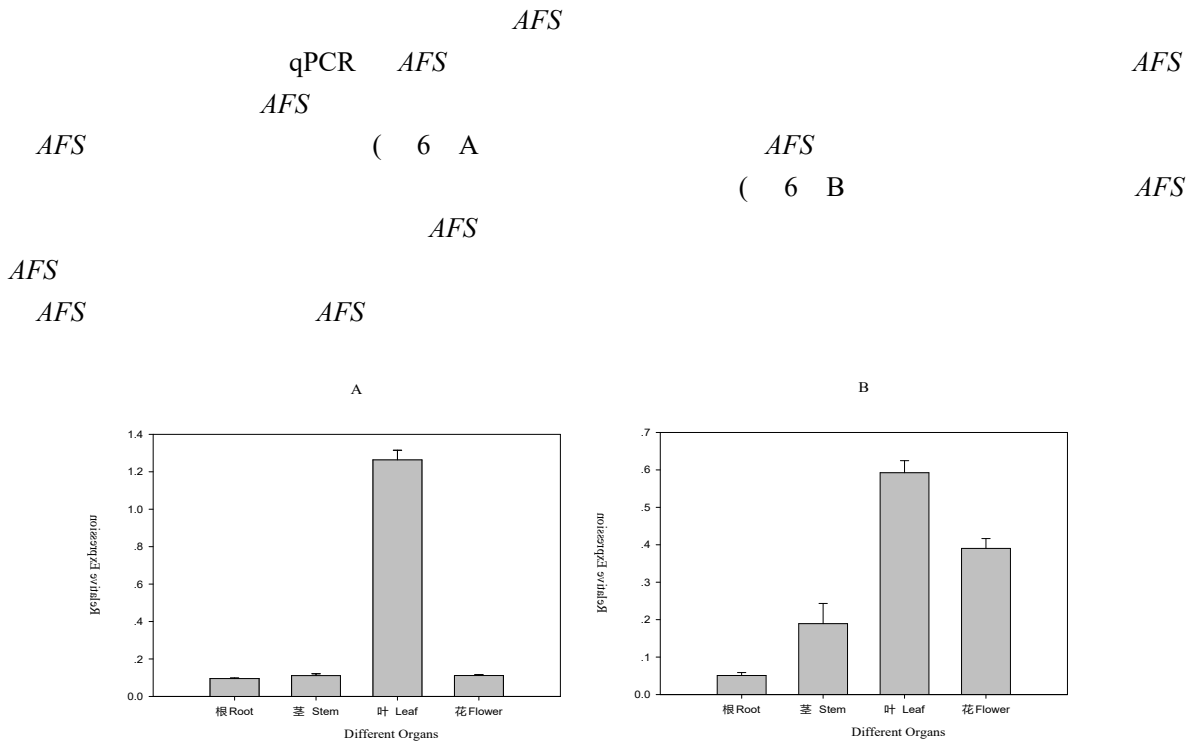
2.4 qPCR 分析不同转基因烟草株系中 AFS 的表达



**图 5 qPCR 分析不同转基因烟草株系中 AFS 的表达**  
**Fig.5 Expression of AFS in different transgenic tobacco strains by qPCR**

A: Chloroplast over-expression plants; B: Mitochondria over-expression plants

### 2.5 转基因植株 AFS 表达的空间特异性



**图 6 AFS 在不同器官中的表达**  
**Fig.6 Expression of AFS in different organs**

A: Chloroplast over-expression plants; B: Mitochondria over-expression plants

### 3 讨论

FPP C15 [10]  
 FaNES1 FaNES1

FPP [11,12]  
 [4] Rubisco 35S 7~8 [13]

*Rubisco*

35S

[14]

*Ru*

,

PCR

 $\alpha$ - $\alpha$ -

[15,16]

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