

Figure S1. Motifs analysis of GmCEPs. Each block represented the position and strength of a motif. The blocks of GmCEPs motif were predicted using MEME. The motif sequence were listed in lower panel.

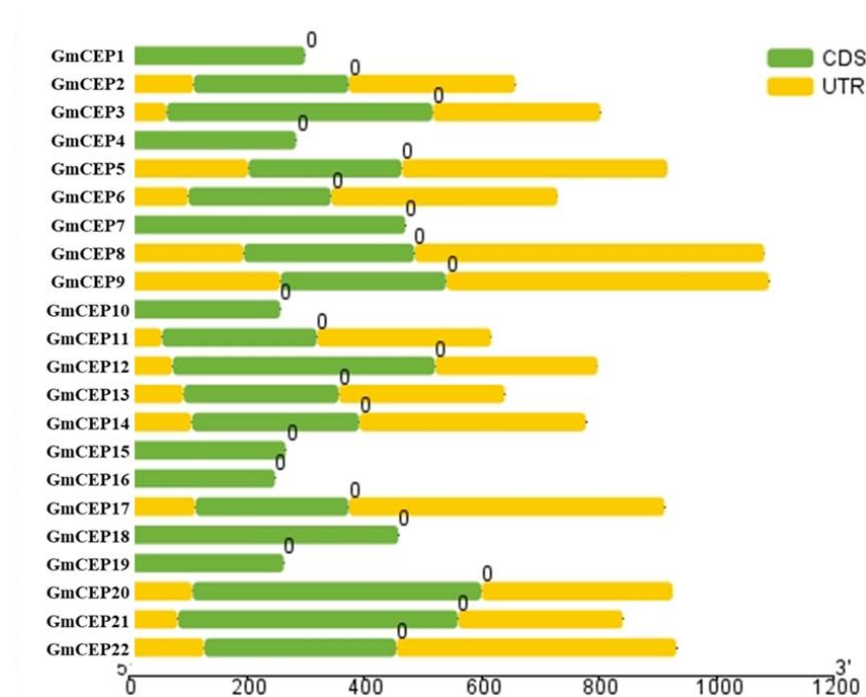


Figure S2. Gene structure of 22 *GmCEP* genes. The gene structure information of *GmCEPs* were obtained through gff file of soybean, the diagram was constructed by Tbtools software.

A

Target-1

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ATGGGCCGTCTCACCACATTTGGTTTCTCTTGCACCTCTGTTTTTGTCCCATGAAGTACTCGGTAGTGAGGGGAAGGA
ATTGAGACATACCATTCAATCACCAGATGCTACCAAAGCAATGAGCATTGCAACAAAAAGTGCTAATGTCATCCCAAC
CTACCGTAGCATTAGAAGCCTAGCGGGAGATGTTGAAGCTTTAGGCCTACAACCTCTGGCCACAGTCCTGGAGTTGGC
CATTAAT
  
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Target-2



Figure S3. Validation of the mutation of *GmCEP6*-edited hairy roots. (A) Sequence of a region of soybean *GmCEP6* with two target sites indicated. (B) Alignment of sequences of target-1 mutated alleles identified from cloned PCR fragments from *crispr cas9 GmCEP6* (KO) transgenic root lines. Highlighting blue denotes the degree of homology of the aligned fragments, and only aligned regions of interest are displayed. Each trait represents a different mutation type. The most mutation was a base shift, represented by a green triangle, with a total of 5 (n=13).

Table S1. Primer sequences used in this study.

Primer Name	Sequence (5'-3')	Purpose
GmCEP6-qRT-F	ATGGGCCGTCTCACCCACATTTG	qRT PCR
GmCEP6-qRT-R	ATGGCCAACCTCCAGGACTG	qRT PCR
GmCYP2-qRT-F	CGGGACCAGTGTGCTTCTTCA	qRT PCR
GmCYP2-qRT-R	CCCCTCCACTACAAAGGCTCG	qRT PCR
GmNIN1a-qRT-F	CATCTTGAGCCTCTACCACC	qRT PCR
GmNIN1a-qRT-R	GCTTTGACTCTAAAAGTGCCGG	qRT PCR
GmENOD40-1-qRT-F	TGGACAACACCCTCTAAACCA	qRT PCR
GmENOD40-1-qRT-R	GTGAGGGAGTGTGAGGAGTGA	qRT PCR
GmNSP1-qRT-F	GGTCTATAACTTTTGCTTCCAGC	qRT PCR
GmNSP1-qRT-R	CAGTGTCTTCGCCAAGAAGCTTG	qRT PCR
GmHAP2-1-qRT-F	CACGCCATCTACATGCGAC	qRT PCR
GmHAP2-1-qRT-R	ACTGGAATTGTCGGCCGCTG	qRT PCR
GmHAP2-2-qRT-F	GGAGTGCCTTAGGATCTCAACC	qRT PCR
GmHAP2-2-qRT-R	TACCGCTTGCTTACCGGCTG	qRT PCR
GmCEP6-Promoter-F	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCATGC	Vector construction
GmCEP6-Promoter-R	ACTGGCACGGAGATTTAG	Vector construction
GmCEP6-hairy-OE-F	GGGGACCACTTTGTACAAGAAAGCTGGGTCTGGGTG	Vector construction
GmCEP6-hairy-OE-R	TTTAGGCAAGGA	Vector construction
GmCEP6-Crispr-BsF1	GGGGACAAGTTTGTACAAAAAAGCAGGCTTCATGG	Vector construction
GmCEP6-Crispr-BsR1	GCCGTCTCACCCACATTTG	Vector construction
GmCEP6-Crispr-BsR1	GGGGACCACTTTGTACAAGAAAGCTGGGTCATGGCC	Vector construction
GmCEP6-Crispr-BsR1	AACTCCAGGACTG	Vector construction
GmCEP6-Crispr-BsR1	ATATATGGTCTCGATTGATGAACTACTCGGTAGTGA	Vector construction
GmCEP6-Crispr-BsR1	GTT	Vector construction
GmCEP6-Crispr-BsR1	ATTATTGGTCTCGAAACCTAGGCTTCTAATGCTACG	Vector construction
GmCEP6-Crispr-BsR1	CAA	Vector construction
GmCEP6-Crispr-BsR1	TGATGAACTACTCGGTAGTGAGTTTTAGAGCTAGAA	Vector construction

Crispr-F01	ATAGC	construction
GmCEP6-	AACCTAGGCTTCTAATGCTACGCAATCTCTTAGTCG	Vector
Crispr-R01	ACTCTAC	construction