

Table S1. All primer sequences used in this paper.

primers name	Sequence (5'-3')	purposes
GaBa1	ctgc <u>GGATCC</u> ATGTGTAGTTGGGCTTCGGT	forward primer for cloning 2430 bp <i>AtGCSpro</i> promoter
GaBa2	ctgc <u>GGATCC</u> AGGAAGAAACTGTGGAGGTG	forward primer for cloning 1990 bp <i>AtGCSpro</i> promoter
GaBa3	ctgc <u>GGATCC</u> CACAGCTCATATAATTAGGAG	forward primer for cloning 1640 bp <i>AtGCSpro</i> promoter
GaBa4	ctgc <u>GGATCC</u> ACCATGGTACCATACTATTCC	forward primer for cloning 1200 bp <i>AtGCSpro</i> promoter
GaBa5	ctgc <u>GGATCC</u> TGAGGTATGTGGTTGGTAG	forward primer for cloning 850 bp <i>AtGCSpro</i> promoter
GaBNR	gttc <u>GGTCTCCC</u> ATGGTATATAGCTCCTGCAATT	reverse primer for cloning <i>AtGCSpro</i> promoter
Sap401	CCGCTCTCAGCTAGGATCCATGGAATCGGCAG CAAAGGA	construction of intermediate vector pRd35SCas9
Rd4012	TGGCTCGAAGATA <u>ACCTGCAAGAGGTCTCCTGA</u> CCAATGGTGCTTTGTAG	construction of intermediate vector pRd35SCas9
Rd4013	GACCTCTGCAGGTATCTTCGAGCCA	construction of intermediate vector pRd35SCas9
Sap402	ATGCTCTCtGCTGGTACCGCTTATTGGTTATCT CATCGG	construction of intermediate vector pRd35SCas9
GaK1	ACGGTCTCAGTACATGTGTAGTTGGGCTTCGGT	construction of intermediate vector pRdGa1Cas9
GaBa4	CTGCGGATCCACC <u>ATGGTACCATACTATTCC</u>	construction of intermediate vector pRdGa4Cas9
GaK5	GTTAGGTACCTGAGGT <u>CATGTGGTTGGTAG</u>	construction of intermediate vector pRdGa5Cas9
GaX2	GCGTCTAGAGGTATATAGCTCCTGCAATT	construction of intermediate vector pRdGa1Cas9, pRdGa4Cas9, and pRdGa5Cas9
UbiKp	AGATGTGGTACCGTGACCGTGACCCGGTCGT	construction of intermediate vector pRdUbiCas9
UbiXb	ACTAGGTCTCTAGACTGCAGAACGTAACACC AAACAAACAGG	construction of intermediate vector pRdUbiCas9
YAOF18	GGCGCGCCTGCAGGTACCTCTGAATCGAGCTT TCGGAA	construction of intermediate vector pRdYCas9
PYao2	tggctctcttagTCTCTCACTCCCTCTTAG	construction of intermediate vector pRdYCas9
Ktrj71	GTCAAGGGTACAACGAGGAATT	construction of CRISPR/Cas9-mediated <i>GmNARK</i> (<i>Rj7</i>) gene knockout
Ktrj72	AAACGAATTCTCGTTGTACCCCT	construction of CRISPR/Cas9-mediated <i>GmNARK</i> (<i>Rj7</i>) gene knockout
KtLjNL1	ATTGGTTGTTCTGATGGATCCTT	construction of CRISPR/Cas9-mediated knockout for targeting <i>LjNLP4</i> with pPG35Cas9
KtLjNL2	AAACAAGGATCCATCAGAACAAAC	construction of CRISPR/Cas9-mediated knockout for targeting <i>LjNLP4</i> with pPG35Cas9, pRd35Cas9 and pRdGa1Cas9
KtLjNL3	GTCATGTTCTGATGGATCCTT	construction of CRISPR/Cas9-mediated knockout for targeting <i>LjNLP4</i> with pRd35Cas9 and pRdGa1Cas9
LjNLP4F	GTGGGATCCATATTGGCAAC	PCR amplify <i>NRSYMI/LjNLP4</i> targeted site for restriction enzyme digestion analysis
LjNLP4R	TACCGTGCTGCCTACAGATG	PCR amplify <i>NRSYMI/LjNLP4</i> targeted site for restriction enzyme digestion analysis
ktGmR11	GTGTGGTCTCGGTATAGAATT <u>CATAAAGCTTG</u> AGTTTAGAGCTAGAAAATAGCAAG	construction of p2×35Spro-Cas9- <i>Rfg1/GmNNL1</i> , pAtGCSpro ₁₂₀₀ -Cas9- <i>Rfg1/GmNNL1</i> for targeting <i>Rfg1</i>

Table S1 All primer sequences used in this paper (to continue).

primers name	Sequence (5'-3')	purposes
ktGmR12	GTGTGGTCTCGAAACCCATGGCATGTCCTGTTCAATCTCTT AGTCGACTCTACC	construction of p2×35Spro-Cas9- <i>Rfg1/GmNNL1</i> , pAtGCSpro ₁₂₀₀ -Cas9- <i>Rfg1/GmNNL1</i> for targeting <i>GmNNL1</i>
ktLjSNF	GTGTGGTCTCGGTCAAGTTGTTCTGATGGATCCTGTTTAGAG CTAGAAATAG	construction of p2×35Spro-Cas9- <i>LjNLP4LjSYMRK</i> , pAtGCSpro ₁₂₀₀ -Cas9- <i>LjNLP4LjSYMRK</i> for targeting <i>LjNLP4</i>
ktLjSNR	GACAGGTCTCGAAACTGCAGTT CCTCTTACTTCACAATCTCTT AGTCGACTCT	construction of p2×35Spro-Cas9- <i>LjNLP4LjSYMRK</i> , pAtGCSpro ₁₂₀₀ -Cas9- <i>LjNLP4LjSYMRK</i> for targeting <i>LjSYMRK</i>
ktSITRY1	CTGTGGTCTCGGTCAAGGTGGTGCATGAGTTGTGTGTTTAGA GCTAGAAAATAGC	construction of p2×35Spro-Cas9- <i>SITRY</i> and pAtGCSpro ₁₂₀₀ -Cas9- <i>SITRY</i>
ktSITRY2	CTGTGGTCTCGAAACACAAGTTGTGCATCCTGTCAATCTCTT AGTCGACTCTAC	construction of p2×35Spro-Cas9- <i>SITRY</i> and pAtGCSpro ₁₂₀₀ -Cas9- <i>SITRY</i>
Rj71	ACTTGAGGGCACTGCAGACT	PCR amplify <i>GmNARK</i> (<i>Rj7</i>) targeted site for restriction enzyme digestion analysis
Rj72	ACGTCTTCGCAAGGTTCATC	PCR amplify <i>GmNARK</i> (<i>Rj7</i>) targeted site for restriction enzyme digestion analysis
GmRHin1	TCTAATTGAGACACAGGCAG	PCR amplify <i>GmNNL1</i> targeted site for restriction enzyme digestion analysis
GmRHin2	CTATTGGCAGATGCTCGG	PCR amplify <i>GmNNL1</i> targeted site for restriction enzyme digestion analysis
GmRNco1	TTGTTGCACCCATTGATC	PCR amplify <i>Rfg1</i> targeted site for restriction enzyme digestion analysis
GmRNc4	GCCCCTTATATACTTCGAATCCA	PCR amplify <i>Rfg1</i> targeted site for restriction enzyme digestion analysis
LjSYF	GTGAAC TTGACTACAGGGGAAC	PCR amplify <i>LjSYMRK</i> targeted site for restriction enzyme digestion analysis
LjSYR	CCTGCATAGGGATAATGTCAG	PCR amplify <i>LjSYMRK</i> targeted site for restriction enzyme digestion analysis
SLTRY1	CCCTCCTAACACAGCAACTCTC	PCR amplify <i>SITRY</i> targeted site for restriction enzyme digestion analysis
SLTRY2	GGCCAACAAGTTCATTGATG	PCR amplify <i>SITRY</i> targeted site for restriction enzyme digestion analysis
GmActinF	GAGCTATGAATTGCCTGATGG	qRT-PCR primer for amplification <i>Actin</i> gene in soybean
GmActinR	CGTTTCATGAATTCCAGTAGC	qRT-PCR primer for amplification <i>Actin</i> gene in soybean
GUSPF	ATGGTAGATCTGAGGAACCG	qRT-PCR primer for amplification <i>GUSplus</i> gene
GUSPR	GCCAATGTCATTGTAAGTC	qRT-PCR primer for amplification <i>GUSplus</i> gene