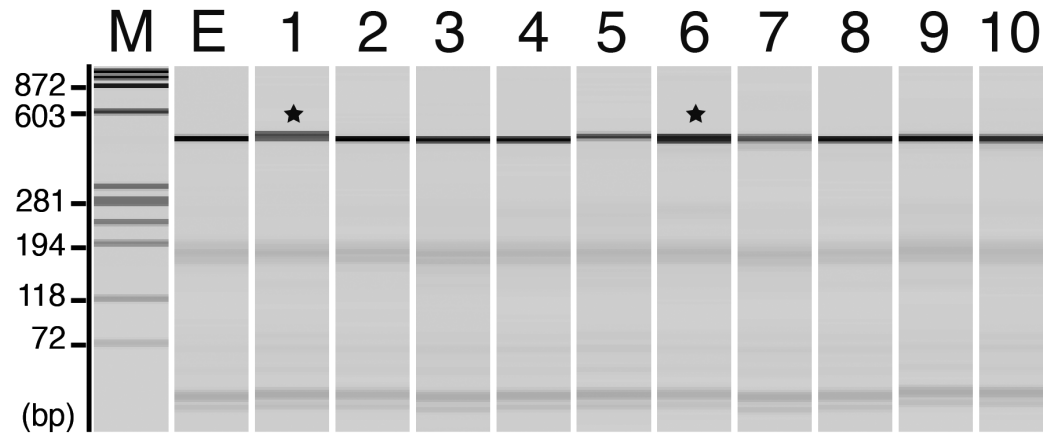
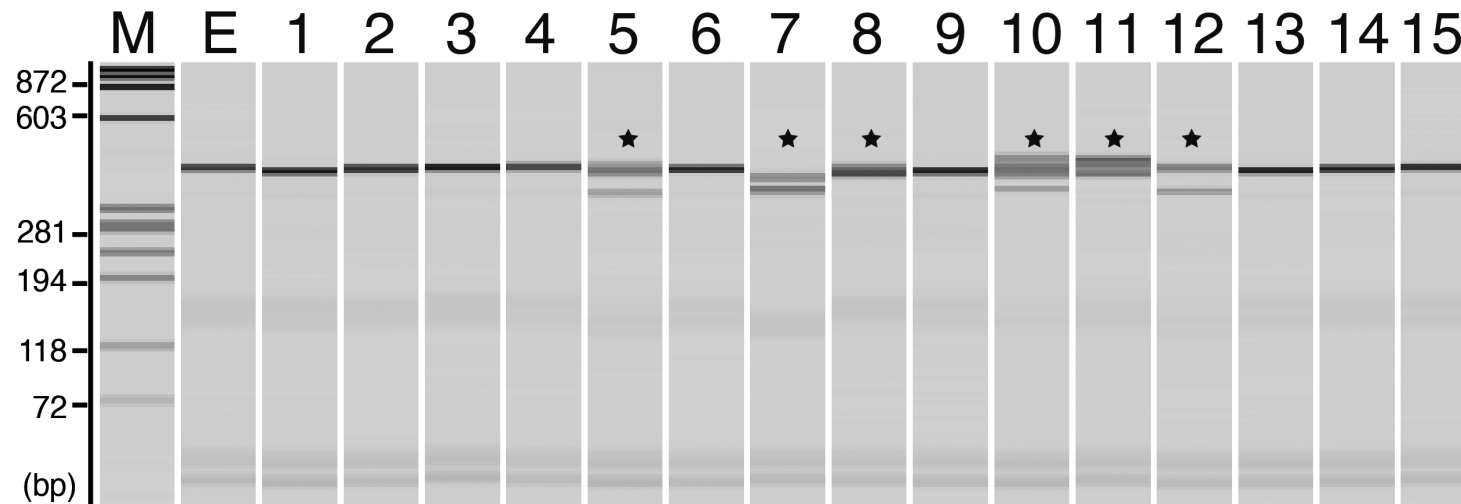


Supplementary Figure S1. Target sites for *SIS5aR1* and *SIS5aR2*. The 20-bp target sequences and protospacer adjacent motif (PAM) sequences are indicated in gray.

pMgP237_SIS5aR1_ko1



pMgP237_SIS5aR2_ko1



Supplementary Figure S2. Heteroduplex mobility assay with the MultiNA electrophoresis system. Multiple heteroduplex peaks were detected in PCR amplicons from the CRISPR/Cas9 transgenic tomato hairy roots. Black stars indicate the potential mutated lines. E: hairy roots transformed by empty pMgP237 vector. M: internal marker.

Supplementary Table S1. Candidate S5αRs from tomatoes were selected based on the BLASTP search in the Sol Genomics Network (<http://solgenomics.net>). RPKM values were obtained from the Tomato Functional Genomics Database (<http://ted.bti.cornell.edu>).

Clone ID	Bud	Flower	Leaf	Root	1cm_Fruit	2cm_Fruit	3cm_Fruit	Mature green	Breaker
<i>SIS5aR1</i>	34.5	21.4	16.4	15.0	41.9	27.5	38.0	55.3	43.5
<i>SIS5aR2</i>	74.8	12.8	30.2	22.9	105.3	32.3	7.4	3.8	1.2

Supplementary Table S2. Oligonucleotides used in the present study.

Primer No.	Description	Sequences
1	SIS5aR1qPCR Fw	5`-TGTGGTTTTGGGTTTCGATTTGCCG-3`
2	SIS5aR1qPCR Rv	5`-TCACTCTTCAGGCCCACTAACACC-3`
3	SIS5aR2 qPCR Fw	5`-TTCACTTCTCCTTACGGCAAACACTACA-3`
4	SIS5aR2 qPCR Rv	5`-CGAAATACGATAATTGTGAGCCAAAGCGT-3`
5	ubiquitin qPCR Fw	5`-CACCAAGCCAAAGAAGATCAAGC-3`
6	ubiquitin qPCR Rv	5`-TCAGCATTAGGGCATCCCTTACG-3`
7	SIS5aR1 ORF BamHI Fw	5`-GGATCCATGTTTTTCAGATCAAACCTTGTATCAC-3`
8	SIS5aR1 ORF Sall Rv	5`-GTCGACTCAATACAAAAATGGAATAACAGCTTTT-3`
9	SIS5aR2 ORF BamHI Fw	5`-GGATCCATGTTCTCTTCAGATGAAAATCTCTTC-3`
10	SIS5aR2 ORF Sall Rv	5`-GTCGACTCAATACAAAAGGGGAAAACAGC-3`
11	SI3βHSD1 ORF BamHI-NdeI Fw	5`-GAATTCCATATGGCAAATAAGCTCAGGTTGGAGGGC-3`
12	SI3βHSD2 ORF Sall Rv	5`-GTCGACTTATTGTAGCTTCAAATAGAACTTAGTCCACC-3`.
13	F2_tgRNA_SIS5R1_492-511	5`-TTGGGTCTCGTGCAGATGCTTATGTACAAGCTAGGGTTTTAGAGCTAGAAATAGCA-3`
14	R2_tgRNA_SIS5R1_552-571	5`-TTGGGTCTCCAAACGCCGGCAAATCGAACCCAACTGCACCAGCCGGGAATCGAA-3`
15	F2_tgRNA_SIS5R1_449-468	5`-TTGGGTCTCGTGCAGTATTGGGCTGGTTATATTCGTTTTAGAGCTAGAAATAGCA-3`
16	R2_tgRNA_SIS5R1_511-530	5`-TTGGGTCTCCAAACCCTTGACTCTTCAGGCCCAACTGCACCAGCCGGGAATCGAA-3`
17	AtU6-pMgP237 Fw	5`-GGCCCCTGGGAATCTGAAAG-3`
18	AtU6-pMgP237 Rv	5`-GGAAGAAGAAATCGATCTGGAAAATTTTGC-3`
19	SIS5aR1_g43 Fw	5`-CTCTTCCTCATCGGACCACC-3`
20	SIS5aR1_g501 Rv	5`-CCAAACATTCACCGCCATCC-3`
21	SIS5aR2_g296 Fw	5`-TTCGCCTCCGTAGCAGAAAC-3`
22	SIS5aR2_g678 Rv	5`-TCGAAACAGCTCGAGGAACC-3`