

Figure S1

A

BstYI
GATCC CAATTTTCCTCAATTTGTAGTCGACAATGGCACCTCTTCCAGCACCATACTCAGGAACCA
 GCACTCTTGCTCTGGTGGCGCGGCATCAGCATTTACCTTTGGACTCGCTTATGGAAGC **GTTAA**
 MseI

SlAucsia-1 ATCGGCCGAATTGAGGCGTGTGCGCACATTTGGTTGCAGCTTCCCCGATTCTCGTCATCA
SlAucsia-2 -----

SlAucsia-1 CAAAACTGATATTGACGGGCTGAGTGTGGTGAATTCTAGGGCTTGTAAGGAGATCTAG
SlAucsia-2 -----

SlAucsia-1 CAGCTTCTTTCTCAGATCCCAATT**TTCCTCAATT-TGTAGTCGACAATGGCACCTCTTCC**
SlAucsia-2 -----AGAGATCTTGAAGCTCTCTGCTTGTGAAGTCAACA**ATGGCACCTCTTCC**
 ***** * ** ** * * ***** *****

SlAucsia-1 **AGCACCATACTCAGGAACCAGCACTCTTGCTCTGGTGGCGCGGCATCAGCATTACCTT**
SlAucsia-2 **TGGACCTTACTCTGGAACCAGCACCTTCTGCTCTGGTGGCTCGTACTTCAGCCTTACATT**
 * *** ***** ***** * * ***** ** ** *

SlAucsia-1 **TGGACTCGCTTATGGAAGCGTTAAGCTCAAGTATCTCAAGGCAAAGGCCAAGTCTCAAAA**
SlAucsia-2 **TGGACTTGTCTATGGTAGCATGAAGCTCAAGTATCTCAAGGCTAAAGCCAAGTCTCACCA**
 ***** * ***** ** * ***** ***** ** ***** *

SlAucsia-1 **GAAGGCTGAAGCTAAGGCACGTCAC**TGAAAGATATCT-CATGGTGG**ATTGCTCTTTGCT**
SlAucsia-2 **GAAAGCTGAAGCTAAGGCACATCAC**TGAAAGAGAGCAGTATGCCGCATTGCTGTCTGTT
 *** ***** ***** ***** * * *** * ***** * ** *

SlAucsia-1 GATAGAATAAGATGTGGCTGATTAAACTGTATCGCGTTTCACCTGATGCTTATAG--GAT
SlAucsia-2 AA--AAATAATATATGAC-AATCACATTATATCAGTTTTATTATGGCCCTGTTGCTGAC
 * ***** ** * * * * * * * * * * ***** * * * * *

SlAucsia-1 AATTAGATTCTGACTTGTAGCTGTTCAATATTTTGGT----AGTTTTGTGGT-TTTTATG
SlAucsia-2 CATCAT-TTTCGACTTGTAACTGTTCAATTTTGTGTT**CACGAGTTATTAATGTTTATG**
 ** * * * ***** ***** ** * * * * * * * * * * *****

SlAucsia-1 GTTCAGTGGGGTTCTCCACGATCAAATATTATTATATTTATAGTACAAGCCCTCTCA---
SlAucsia-2 **GTCCAATGAGTACAATGGTGGTCCAATTTTGGAGATTATATTGGACAAACCGGGGAAAAG**
 ** ** *

SlAucsia-1 TTAAAATTCAA--ACAACATATGGT-----
SlAucsia-2 **TTAGCATTGATCTGCTAGCATAATAATCTTCTTGTTACATCA**
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B

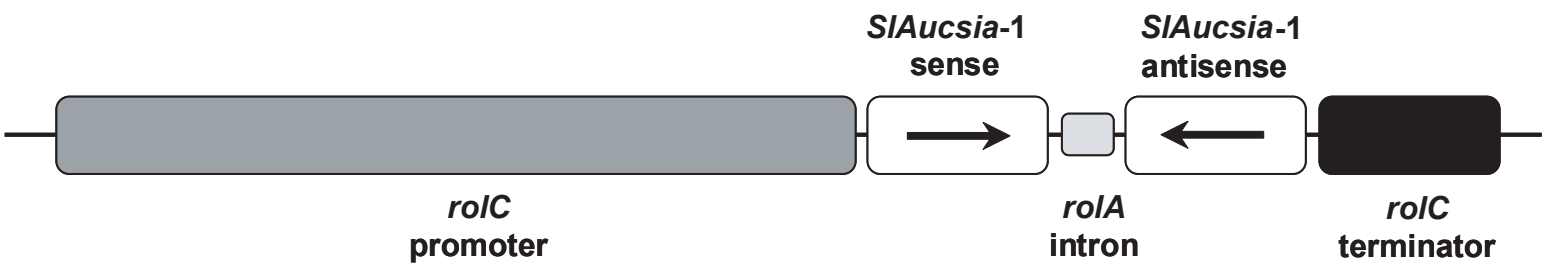


Figure S1. Sequence of cDNA-AFLP clone, *SlAucsia-1* and *SlAucsia-2* transcripts and schematic drawing of rolC-hpAucsia construct. A, (Upper) Sequence of differentially-expressed cDNA fragment. Restriction sites are underlined and selective nucleotides used in the primer combination are indicated in green (BstYC-MseC). (Bottom) Alignment of *SlAucsia-1* (AK224828) and *SlAucsia-2* (AK224647) transcripts. The boxed region is the sequence used to design the sense and antisense arms of rolC-hpAucsia construct. DNA fragments corresponding to the sequences shown in red, blue and magenta were used to produce the three RNA probes for siRNA analysis. B, Schematic drawing of rolC-hpAucsia construct.