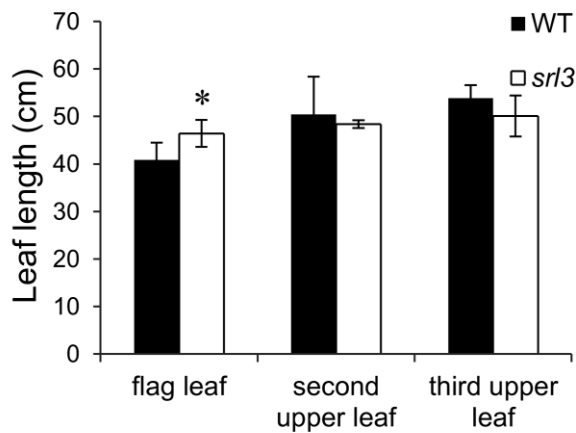


Supplemental data

Appendix A



Appendix A Statistical analysis of leaf length in the wild type and *srl3* mutant. Values are the mean \pm SD of ten samples. *, significant difference at $P < 0.05$ (Student's *t*-test).

Appendix B Primers used in the study

Purpose	Primer name	Sequence
Mapping	RM13633-F	CTCTTCCGACCGGAGATAATGC
	RM13633-R	GTTCACTCGCTGACGCTATCTGC
	M1-F	GTTAGGGGCTTGGTACGTCA
	M1-R	CAAAGAGGAGTCATGGATGGA
	M4-F	AGGAAATTTTCGTCTGCTTGG
	M4-R	ACCGAACGCATTTAGTGGAC
	M7-F	CCTATGTCGACCGACTCCAT
	M7-R	CAATCCCCGTTGAACAGTTT
	M8-F	AAGGGTTTGGAGGCCTATGT
	M8-R	CGGGGATAAGCAGAGAAGC
	M19-F	TTTATGACAAACCTACTG
	M19-R	AACTCAAACATAGCATT
	M10-F	GGGGAAGTGAAGTCCAACCT
	M10-R	GGAACCGAATTCAAATGCAA
M14-F	CATTGTGTACCGCGAAATTG	
M14-R	TACGCACTCTTAGGCCCACT	

	RM13840-F	CGGTCTTTAGTAATGGTGCTTTGC
	RM13840-R	GAGGCAGGTGTTTGTCTAGC
qRT-PCR	SLL1-F	CAGCTCGCAGGTGTCCAA
	SLL1-R	CCTCCCTAGAGTGA ACTCGAGACT
	SLL2-F	TAGGTTGCCTTCGGTTTTGA
	SLL2-R	TTGGCTCCTCCACCAGTTTG
	REL1-F	GTGGACGACGACGAGGACGA
	REL1-R	GGCGAGGATGAGCAGGAACA
	REL2-F	CTTGCTATCGTCGCTGTCGT
	REL2-R	GCTCTTCCCTTCCATTGACT
	ACL1-F	TCGTGCCTGTCGTCGGAGAA
	ACL1-R	CGTTGTTGTTGCCGTTGTCC
	ACL2-F	CTCGTGCCTGTCGTCGGAG
	ACL2-R	TCGGCGAGCATGACGTAGAG
	ADL1-F	ATCTGGCTTCTTCTTTGGGG
	ADL1-R	TGCCCTTCCTTCGAAAACC
	SRL1-F	ACTGCTTTGCGTCCTTCG
	SRL1-R	CGCCCTGGCTCAACAATA
	NAL1-F	TGCAGTGTCCGCTCAATAGC
	NAL1-R	GACCAGAGCTTCTGCCAACTTT
	NAL7-F	CAGGAGGACGGTGAGTTGTTC
	NAL7-R	GCAGTAGAGCCCGTTTGGAC
	OsZHD1-F	ACCAAGTTCACCCAGGAGCAGA
	OsZHD1-R	CGCCGACCTCGTCACAGAA
	OsZHD2-F	CCGTGCAGCAGTTCTGCGA
	OsZHD2-R	CAGGGTGTGCTTGTTGTTGTG
	Roc5-F	AGGTGGAAATGTGAAGTGTAGG
	Roc5-R	AGGCGAGACCAGACCGAAG
	RL14-F	CTATCCACCGTGCTTGAA
	RL14-R	AGAACCATTTGCCATCCT
	OsAGO7-F	CCGCATCCCCTTGATGATT
	OsAGO7-R	GGCCAATTCATGCTTGCAA

OsBAK1-F	GAGTTGATCTTGGGAATGCTGC
OsBAK1-R	CACTAGGTATCGTTCCGCTTATGTT
LC2-F	ATCATCTGGTGGTGGCGTTGT
LC2-R	TCTTTCGCTGCCGTTTCG
CYCA2.1-F	AGGTTGTCAAGATGGAGAGCGA
CYCA2.1-R	CGCTTTTTGTCTTCCTGGCA
CYCA2.3-F	GTTTCGGTTGACGAGACGATGT
CYCA2.3-R	CGCTGCAAGGAACCTAGAACTG
CYCB2.2-F	CTCAAGGCTGCACAATCTGACA
CYCB2.2-R	GCATTGACGGCTGGAATTTG
CYCD4-F	GCCATGGAGTTGATACATCCAA
CYCD4-R	CCAGTAGGGCTCCGTGGAAT
CAK1A-F	GACCGACAAGGGTTTCAGCAT
CAK1A-R	CCAGCATGTTTCAGGAAGATACAAT
CDKB-F	AAGTTTGGCCAGGAGTGAGCA
CDKB-R	TCAAGAGCATCAGCGTCGAGA
CDKA1-F	GGTTTGGACCTTCTCTCTAAAATGC
CDKA1-R	AGAGCCTGTCTAGCTGTGATCCTT
CDKA2-F	CGAGATTTGAAGCCCCAGAA
CDKA2-R	TCCGCGAGCTTCAATGAGTT
CYCT1-F	GCATTTGTTGCAGCTCAAG
CYCT1-R	TCACCACTTCGCTGACTTATTG
CAK1-F	GACGGTCAGATTAGACGCAAGA
CAK1-R	TCCAAAGGATGTCCACA
OsEXPA2-F	TTTGGCTATTCTGAGGCTGCT
OsEXPA2-R	TGGTCCCAAAGCACAAGAGT
OsEXPA32-F	GCAGTTCGGCGTCACCTACCAG
OsEXPA32-R	GGTTCGATCCAGATTTGCAGTAGTCA
OsEXPB9-F	TGCCAGACGCCATCTACGTATCAA
OsEXPB9-R	TGGCGCAGATATAACAAGCGCAACT
OsEXPB10-F	ATTGGAAGGCCAACGCTCTCTACA
OsEXPB10-R	TTCTGCTCACTCCACAAACCTAGC

MCM2-F	AAGTTGGCAAAAGATCCACGG
MCM2-R	CCCCCAAACATAGCTAGTGCAA
MAPK-F	ACAGAGCAGCCGAATTTTGAGA
MAPK-R	TTCAGCGAAGCTCACACTTGG
YUCCA1-F	TCATCGGACGCCCTCAACGTCGC
YUCCA1-R	GGCAGAGCAAGATTATCAGTC
YUCCA2-F	GTCCAAAGGGAGGAGTCGTCCAG
YUCCA2-R	GCATGATGTTTACACCCGGCCTT
YUCCA3-F	GTGAGAACGGGCTCTACTCGGTTCG
YUCCA3-R	GCTTATGCATGACCGATGAACACG
YUCCA4-F	GCAGAATGGCCTGTACGCTGTTGG
YUCCA4-R	CAGACCAGCACATGACGTGTCTAC
YUCCA5-F	ACCTCCTACGACGCCGCCATGATC
YUCCA5-R	CTCCCAACACAGCGACGACAGAAC
YUCCA6-F	CCATTCCCAGATGGTTGGAAGG
YUCCA6-R	CATGTTGCGCCTCAAGATATTTG
YUCCA7-F	CACTGCTGTGTCCTACAATATCAC
YUCCA7-R	GGAGGTGCATCTCCGTCATCTTC
GH3.6-F	CACTAGCATCTGTCTCATTGTGTCA
GH3.6-R	ACCTTGTCAGTGCCGGAATT
GH3.8-F	CGCGACGGTCCGAATAGT
GH3.8-R	GGAGGCATCCGGCTTCTACT
ARF25-F	CCACAACCTTCTCTAACCAGG
ARF25-R	TGTGGCTTGCAGAATAGTCG
