

Supplementary Data

Supplementary Fig. S1



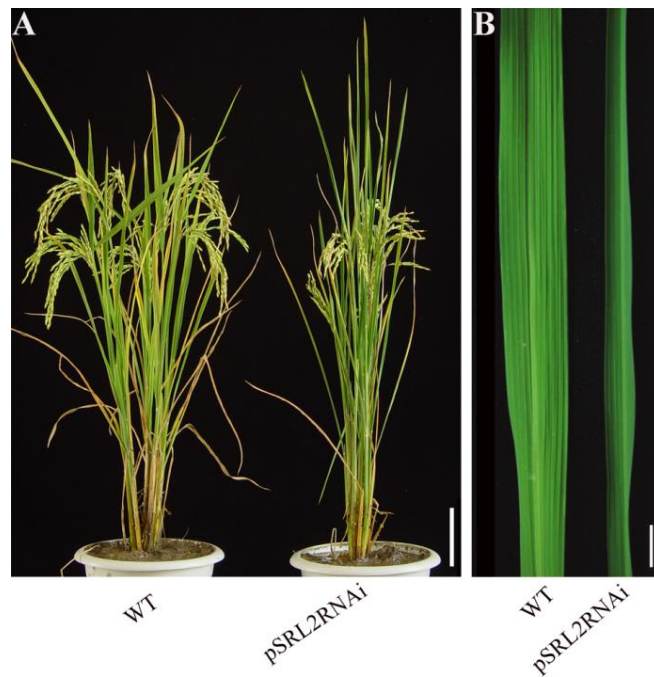
Supplementary Fig. S1. Morphological comparison of wild-type and *srl2* panicles. Scale bars = 1 cm.

Supplementary Fig. S2

SRL2-WT	MGFMSAKLFPSCESMVCVCPALRPSSRRPVKRYKLLAEI FPKT PDGLPNERKI MKLCEYAAKNPLRI PKI AKFLECRSHKELRS AHVNF I KI I TEAYSK	100
SRL2-mutant	MGFMSAKLFPSCESMVCVCPALRPSSRRPVKRYKLLAEI FPKT PDGLPNERKI MKLCEYAAKNPLRI PKI AKFLECRSHKELRS AHVNF I KI I TEAYSK	100
Consensus	ngf ns ak l f p s c e s m v c v c p a l r p s s r r p v k r y k l l a e i f p k t p d g l p n e r k i m k l c e y a a k n p l r i p k i a k f l e c r s h k e l r s a h v n f i k i i t e a y s k	
SRL2-WT	LLFI CKEQVAFYAI SLVNVLT ELLLESKQENI HILGCOTLAKFI YSOVDNTYARNI ESLVRKVCVLSRQOGVEHSLRAASLQCLSAMVFMKEHSYI FVD	200
SRL2-mutant	LLFI CKEQVAFYAI SLVNVLT ELLLESKQENI HILGCOTLAKFI YSOVDNTYARNI ESLVRKVCVLSRQOGVEHSLRAASLQCLSAMVFMKEHSYI FVD	200
Consensus	l l f i c k e q v a f y a i s l v n v l t e l l e s k q e n i h i l g c o t l a k f i y s o v d n t y a r n i e s l v r k v c v l s r q o g v e h s l r a a s l q c l s a m v f m k e h s y i f v d	
SRL2-WT	FDEI VCSVLENYRVEES AAGDEERHAPQHNVDEI VRREGRAGLGGGNDVNCNSTAI RLRSARDSSAL TREERESPEVVAHI CVQKLAELAKESTTNRRI	300
SRL2-mutant	FDEI VCSVLENYRVEES AAGDEERHAPQHNVDEI VRREGRAGLGGGNDVNCNSTAI RLRSARDSSAL TREERESPEVVAHI CVQKLAELAKESTTNRRI	300
Consensus	f d e i v c s v l e n y r v e e s a a g d e e r h a p q h n v d e i v r r e g r a g l g g g n d v n c n s t a i r l r s a r d s s a l t r e e r e s p e v v a h i c v q k l a e l a k e s t t n r r i	
SRL2-WT	LDPMLSYFDKQKQWAPRCGLALVLSDMSYLEKSSGNEQLI LTVSIRHLDHKNVLYDPCI KSDMIQTATLLARQLRSRGI AAELVVAGLDCRHLRKTLEA	400
SRL2-mutant	LDPMLSYFDKQKQWAPRCGLALVLSDMSYLEKSSGNEQLI LTVSIRHLDHKNVLYDPCI KSDMIQTATLLARQLRSRGI AAELVVAGLDCRHLRKTLEA	400
Consensus	l d p m l s y f d k k k q w a p r c g l a l v l s d m s y l e k s s g n e q l i l t v s i r h l d h k n v l y d p c i k s d m i q t a t l l a r q l r s r g i a a e l v v a g l d c r h l r k t l e a	
SRL2-WT	NESASI EELNLNESLQNFLODCLLEVVTGI NDVRLPYDMVAI TLENLPSNPVVARASI GSLLI LSHI I SLTSM LNAPMLFPEALLQQLKSMVHPDVT	500
SRL2-mutant	NESASI EELNLNESLQNFLODCLLEVVTGI NDVRLPYDMVAI TLENLPSNPVVARASI GSLLI LSHI I SLTSM LNAPMLFPEALLQQLKSMVHPDVT	500
Consensus	n e s a s i e e l n l n e s l q n f l o d c l l e v v t g i n d v r p l y d m v a i t l e n l p s n p v v a r a s i g s l l i l s h i i s l t s m l n a p m l f p e a l l q q l k s m v h p d v t	
SRL2-WT	RVGAHHMFSAVI VOGPSRQRS ESDFLYET KKWQSRRTSVFASAT ALLEKL RREKESL GSDKT GNMDDKEKSI SEEENKHVMARKNSAYFSKLVFSFTDR	600
SRL2-mutant	RVGAHHMFSAVI VOGPSRQRS ESDFLYET KKWQSRRTSVFASAT ALLEKL RREKESL GSDKT GNMDDKEKSI SEEENKHVMARKNSAYFSKLVFSFTDR	600
Consensus	r v g a h h m f s a v i v o g p s r q r s e s d f l y e t k k w q s r t t s v f a s a t a l l e k l r r e k e s l g s d k t g n m d d e k e k s i s e e e n k h v m a r k n s a y f s k l v f s f t d r	
SRL2-WT	YAALTSAAEEANI VMLTEDQKNQLLSAFVWQAI QTDNTPFNYEAI GHSYSLTVI SSRDKSRNSNNI QFFCLPLSLRSVSLTNGVLSPCSQRSI FTLAT	700
SRL2-mutant	YAALTSAAEEANI VMLTEDQKNQLLSAFVWQAI QTDNTPFNYEAI GHSYSLTVI SSRDKSRNSNNI QFFCLPLSLRSVSLTNGVLSPCSQRSI FTLAT	700
Consensus	y a a l t s a a e e a n i v m l t e d q k n q l l s a f v w q a i q t d n t p f n y e a i g h s y s l t v i s s r l k d s r n s n n i q f f c l p l s l r s v s l t n g v l s p s c q r s i f t l a t	
SRL2-WT	SMLAFAGRVCHI TELFDVLRGFTSCNMDPYLRI GEDLQLYVRLQSDLNGYGS CS DQEI ARSVLSDCRTKVGI NDQRVL DVVACALCNLT EMDKDLVVKEL	800
SRL2-mutant	SMLALLGKS...VTLLNCLMC	818
Consensus	s m l a f a g r v c h i t e l f d v l r g f t s c n m d p y l r i g e d l q l y v r l q s d l n g y g s c s d q e i a r s v l s d c r t k v g i n d q r v l d v v a c a l c n l t e m d k d l v v k e l	
SRL2-WT	TENFTPEEVPLFGSNSAFDWFNFVQAFSDESLSFDEECRSRTSVDGGLHESPI TNTGSSI SKTTMPQSVPRVLGVCQLLESALHVAGCVAGASVSTSL	900
SRL2-mutant	918
Consensus	t e n f t p e e v p l f g s n s a f d w f n f v q a f s d e s l s f d e e c r s r t s v d g g l h e s p i t n t g s s i s k t t m p q s v p r v l g v c q l l e s a l h v a g c v a g a s v s t s l	
SRL2-WT	PYGTMTSQEALGSGTRKKLSSVLVNGHDSTPDNPAPSLPSAQHFI I PKVNSCGFESSI RTTLEPCS AVKLPASPFDNFLKAAYRAQ	988
SRL2-mutant	988
Consensus	p y g t m t s q e a l g s g t r k k l s s v l v n g h d s t p d n p a p s l p s a q h f i i p k v n s c g f e s s i r t t l e p c s a v k l p a s p f d n f l k a a y r a q	

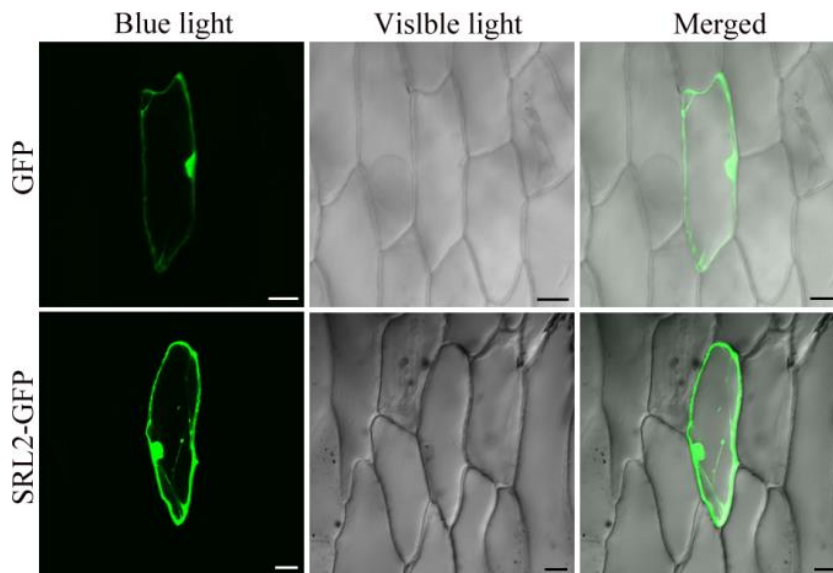
Supplementary Fig. S2. Deduced amino acid sequences of SRL2 and the changes in amino acids caused by mutations.

Supplementary Fig. S3



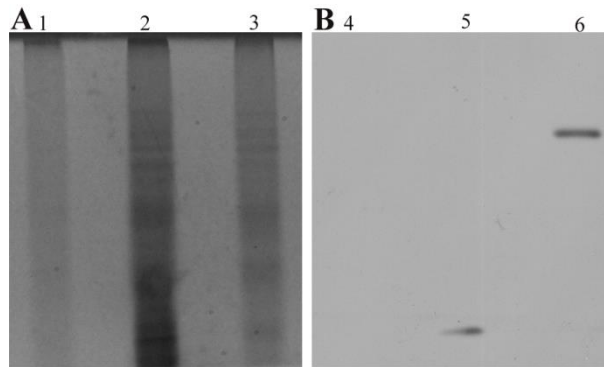
Supplementary Fig. S3. RNAi analyses. Scale bars = 10 cm (A) and 1 cm (B).

Supplementary Fig. S4



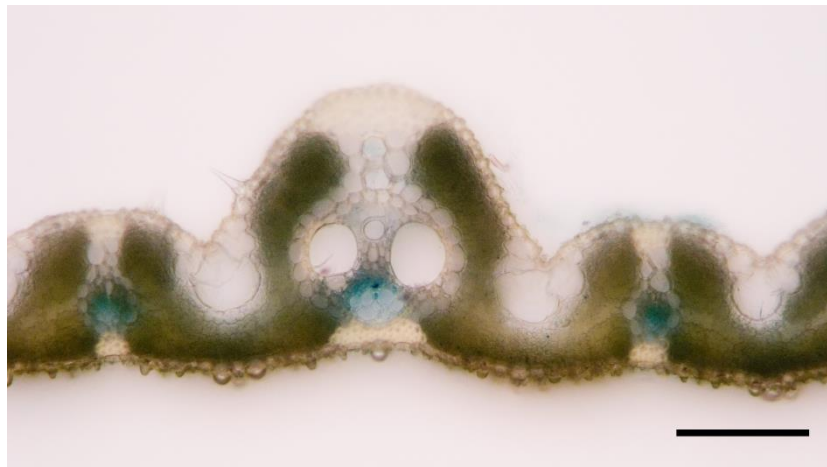
Supplementary Fig. S4. Subcellular location of the SRL2-GFP fusion protein in onion epidermal cells. Onion epidermal cell expressing free GFP showed fluorescence in the nucleus and cytoplasm. Onion epidermal cell expressing SRL2-GFP showed fluorescence in the nucleus and cytoplasm. Scale bars = 50 μm .

Supplementary Fig. S5



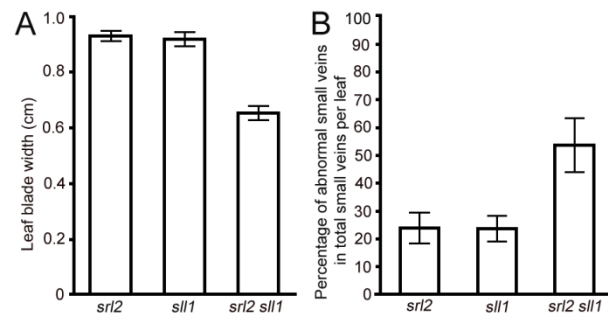
Supplementary Fig. S5. Western blot assays. (A) An SDS-PAGE gel stained with Coomassie Brilliant Blue. (B) Western blotting of the GFP and SRL2-GFP using anti-GFP antibody. Line 1, 4: Total protein from onion epidermis as a negative control; Line 2, 5: Total protein from onion epidermis expressing free GFP; Line 3, 6: Total protein from onion epidermis expressing SRL2-GFP.

Supplementary Fig. S6



Supplementary Fig. S6. Hand-cut section of *ProSRL2-GUS* transgenic rice leaf. Scale bars = 100 μ m.

Supplementary Fig. S7



Supplementary Fig. S7. Characteristics of the *srl2 sll1* double mutant. (A) The leaf width of leaf blade in *srl2*, *sll1* and *srl2 sll1* double mutant. (B) The percentage of abnormal small veins in *srl2*, *sll1* and *srl2 sll1* double mutant.

Table S1: Primers used in map-based cloning of *SRL2*

Marker	Primers	Fragment size (bp)
S1	F 5'-TGTATGAAGAGGAGTGGAGC-3'	100/115
	R 5'-GGTTAATTGGTTTGGTTGGC-3'	
S2	F 5'-CTCCAATTGGTGGGTATG-3'	120/100
	R 5'-ACTTAACACCATTACCCTGC-3'	
S3	F 5'-CGAAACCTCGTCAGTTCGAG-3'	130/115
	R 5'-TTGGAGGGCAATACGGGGAT-3'	
S4	F 5'-AGGAAGGTGCCTCTCTCGA -3'	140/120
	R 5'-CCTGGTTTAGGAGCTTCTCC-3'	
S5	F 5'-CATGAGTATGTCAATGCTGC-3'	100/120
	R 5'-TCTCTTGCTAATATCTAGCC -3'	
S6	F 5'-TACGGTTGCACGGTCAAAGA -3'	100/120
	R 5'-TCGTGTGCATCAAGTTGAGG-3'	

Table S2: Primers used for in situ hybridization analysis

Primer name	Primer pairs
SRL2 T7	GAAGCTCTCAAGCTGGCTTG
SRL2 sp6	CAATATGGTCATACATGGTC

Table S3: Primers used in real-time qRT-PCR analysis

Gene	Primer pairs
<i>SRL2</i>	TCCATCTGCGCAGCATTTC CTACTGGGCACGATATGCAG
<i>YABBY1</i>	TGGTGAATGTGCCAAACAAT GCTTGGTGAAAAGGAGCAAG
<i>YABBY2</i>	AATTTTCGCGGTCAGTGTT AGTTTTTCGCGGAAGCTCATA
<i>YABBY3</i>	ATCAAGGACGAAATCCAACG GGCATCAGTCCAAAGTGGAT
<i>YABBY4</i>	ATCACATCAAAGGGGACCAA CGAGTACGCAATGGCCTTAT
<i>YABBY5</i>	GAGCCTAATGACCGAACAGG CTCTGCCGCTTCTCTGAAGT
<i>YABBY6</i>	TCTGTTCATCACTGGCTTCG CGTGTTGCAGAAGTTGCAGT
<i>YABBY7</i>	CACAGCTGCTAAAACTGGG CTGGACAGCTACAACCTGGA
<i>Ub</i>	GTCTGATCTTCGCTGGCAAGCAGC GCATACTGCTGTCCCACAGGAACTG

Table S4: The width of leaf blade in both wild type and *srl2* (mm)

	WT		<i>srl2</i>	
	before stretched	after stretched	before stretched	after stretched
1	11	11.8	6	8
2	11.5	12	4	8
3	12	12	6	8.5
4	13	13	6	9
5	12.5	12.5	4	7
6	12	12	6	9
7	12	12	5.5	8
8	12	12	6	8
9	12.5	13	4	8.2
10	12	12	4	8
11	13	13	5	8
12	11.5	11.5	5	8.5
13	12.5	13	5	8
14	12	12.8	4	8
15	12	12	3.5	7
16	13	14	5	9
17	11.8	12	5	8.5
18	14	14	5	9
19	12.5	13	5	8
20	11.5	12.5	5	8
21	11	12	5	8
22	12	13	4	7
23	13	13	3	8
24	11	11.5	4	8
25	12	12.5	4	7.5
26	12	13	4	7
27	12.5	13.5	5	8.5
28	12	13	5	8
29	11.5	12	5	8.5

1-29: different rice plants.

Table S5: The height of plants in both wild type and *srl2* (cm)

	1	2	3	4	5	6	7	8	9	10
WT	78.2	79.5	81.3	78.6	78.4	81.2	78.1	79.2	82.1	79.2
<i>srl2</i>	56.1	55.2	57.2	54.3	56.5	56.7	57.4	55.1	55.2	54.3

1-10: different rice plants.

Table S6: The length of internode in both wild type and *srl2* (cm)

	WT					<i>srl2</i>				
	I	II	III	IV	V	I	II	III	IV	V
1	27.7	12.8	8.5	6.7	2.2	17.8	7.5	4.1	2.5	1.5
2	27	12	8.2	6.8	2.1	19	7	4	3.1	1.3
3	27.6	13	8.5	6.4	2.4	17.5	7	4.2	3	1.2
4	26.1	11	7.2	5.8	2.3	16.5	7.2	4	3	1.2
5	26.8	12	8.8	6	2.2	17	8	3.8	2.5	1.1
6	27.5	12.1	8.1	6.5	1.8	20	8.5	5.5	3.5	1.5
7	27	13.2	8.5	6.5	2.2	18	8	5.6	2.5	1.1
8	28.5	13.2	8.5	6.2	2.4	17.5	7.2	4.2	2	0.9
9	26.8	12.5	7.2	6.4	2.5	18	7.8	3	1.5	1.2
10	28.5	12.4	7.8	6.5	2.5	17.5	7.2	5.5	3.5	0.9

1-10: different rice plants. I-V: the 1st to 5th internodes.

Table S7: The number of abnormal small vascular bundles in *srl2*

	No. of abnormal SV	No. of normal SV	No. of total SV
A	5	17	22
B	5	23	28
C	5	18	23
D	6	22	28
E	5	18	23
F	5	23	28
G	5	21	26

A-G: different rice plants. SV: small vascular bundles

Table S8: The number of cell in each internode in both wild type and *srl2*

	WT					<i>srl2</i>				
	I	II	III	IV	V	I	II	III	IV	V
1	3523	1991	1545	924	560	2591	1385	1045	555	240
2	3796	1845	1436	899	768	2456	1503	1080	600	280
3	3633	1874	1508	927	731	2501	1488	1150	515	300
4	3881	2008	1568	1001	677	2602	1550	1050	650	326

1-4: different rice plants. I-V: the 1st to 5th internodes.

Table S9: The number of large vascular bundles and small vascular bundles in both wild type and *srl2*

	WT		<i>srl2</i>	
	No. of LV	No. of SV	No. of LV	No. of SV
1	12	38	10	22
2	11	33	10	28
3	12	34	9	23
4	11	34	10	28
5	11	33	10	23
6	11	28	10	28

1-6: different rice plants; LV: large vascular bundles; SV: small vascular bundles.