

Identification of quantitative trait loci for resistance to bending-type lodging in rice (*Oryza sativa* L.)

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Abstract Bending-type lodging is one of the most important factors affecting the yield and grain quality of rice. This study identified quantitative trait loci (QTLs) for physical strength of the upper culms, and evaluated QTL effects on lodging resistance. In 2010 and 2011, QTLs for breaking strength, length, and diameter of the top three internodes were identified by analyzing chromosomal segment substitution lines (CSSLs) developed from ‘Koshihikari’ and ‘Kasalath’. The QTL analysis indicated that ‘Kasalath’ had two types of QTLs: one to strengthen specific internodes and one to simultaneously improve the physical strengths of plural internodes or the top three internodes. A QTL for breaking strengths of the top three internodes (*bsuc11*) was detected on chromosome 11 in both years. This QTL did not overlap with that for internode length. To evaluate the effects of *bsuc11* on lodging resistance, this study selected three CSSLs with *bsuc11* and analyzed the breaking strengths of the top three internodes after heading and the pushing resistance of the lower part. Internodes of ‘Koshihikari’ showed decreased breaking strengths after grain filling, while those of CSSLs with *bsuc11* did not show this decrease in breaking strength. The pushing resistance of the lower part at

the fully ripe stage was the same in ‘Koshihikari’ and CSSLs with *bsuc11*. These results suggested that *bsuc11* could be a target to improve the physical strength of the upper culms to resist bending-type lodging, and that the physical strengths of upper and lower parts are controlled by different genetic factors in rice.

Keywords Lodging resistance · Breaking strength · QTL · CSSL · Rice

Introduction

Bending-type lodging is the main type of rice lodging, which is one of the most important problems in rice production as it leads to low yields and poor grain quality (Kono 1995). Several Japanese rice cultivars, e.g. ‘Koshihikari’, are prone to bending-type lodging because of their heavy panicle weights, which make them vulnerable to environmental stresses from rain and wind at the fully ripe stage (Kashiwagi et al. 2007). This lodging-type is associated with heavier panicles, weaker upper culms, and higher plant stature, and reducing plant height has been the main target for breeding lodging-resistant varieties. When breeding for higher yield or greater biomass, an additional improvement in the physical strength of culms is one of the best traits for resistance to bending-type lodging. In fact, the New Plant Type (NPT) developed by the International Rice Research Institute showed

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thick and sturdy stems (Peng et al. 1999a), and a few second-generation NPTs produced high yields as a result of accumulating large amounts of aboveground biomass (Peng et al. 2008). The next breeding objective for lodging resistance is to target the physical strength of the plant, rather than plant height.

In previous studies, several techniques have been used to evaluate the physical strength of the plant body. Bending stress and breaking strength were measured in specific parts of cereal plants. Kokubo et al. (1989) measured maximum bending stress of the fourth internode in barley. Similarly, stiffness and the bending moment at breaking were measured for rice culms (Kashiwagi et al. 2008; Ookawa et al. 2010). Other ways to evaluate the physical strength of plant body include measuring resistance to artificial lodging by pushing the plant body. Pushing resistance has been used to evaluate lodging resistance in rice and maize (Fouéré et al. 1995; Kashiwagi and Ishimaru 2004). In their study on stem and root lodging resistance, Berry et al. (2003) used field-based measurement techniques to evaluate resistance to rotational displacement. Physical strength measurement is a useful way to evaluate lodging resistance in crops, and is required for breeding to reduce lodging damage.

The physical strength of the plant body has also been evaluated by analyzing morphological traits and chemical components. Among the morphological traits of rice and wheat, diameter and weight of stems were significantly correlated with the physical strength of the plant (Atkins 1938; Zuber et al. 1999; Kashiwagi et al. 2008). Culm wall thickness also was strongly related to lodging score and culm stiffness (Tripathi et al. 2003; Kashiwagi et al. 2008). Structural carbohydrates including cellulose and lignin are known to contribute to physical strength in plants (Kokubo et al. 1991; Jones et al. 2001; Li et al. 2003). In rice, it was reported that high re-accumulation of non-structural carbohydrates in culms at the fully ripe stage was a character of cultivars with high lodging resistance (Sato 1957; Takaya and Miyasaka 1983; Yang et al. 2001). The amount of minerals in culms is also important for lodging resistance, especially silicon, which increases culm rigidity (Idris et al. 1975). Thus, the physical strength of the plant body comprises various traits. Understanding the genetic controls of these traits is important in breeding for lodging resistance.

Certain genetic factors, including quantitative trait loci (QTLs) for lodging resistance have been studied

in several crops, e.g. rice, wheat, barley, soybean, and field pea (Lee et al. 1996; Keller et al. 1999; Spaner et al. 1999; Tar'an et al. 2003; Kashiwagi and Ishimaru 2004). The improvement of lodging resistance requires effective targets for breeding, because lodging resistance is related to complex factors including morphological traits, chemical contents, and physiological characteristics. Mulder (1954) proposed that lodging results from a disproportionate balance between the weight of upper parts and the sturdiness of the basal parts. Genetic approaches to improve these factors have been reported in rice. For example, the semidwarf gene, *sd-1*, the famous gene for the “Green revolution”, reduced plant height through the mutation of a gibberellin-biosynthesis gene (Khush 1999; Sasaki et al. 2002), and reduced the moment of upper parts. In other cereal crops, genetic factors related to gibberellin, e.g. *Rht-B1* and *Rht-D1* in wheat (Peng et al. 1999b) and *Dwf2*, *gai*, and *gal* in barley (Börner et al. 1999; Ivandic et al. 1999) contributed to dwarfing. Genetic factors related to another plant hormone, brassinosteroid, also reduced plant height in rice (Yamamuro et al. 2000; Mori et al. 2002; Hong et al. 2003). As an example of the sturdiness of basal parts, a QTL for the pushing resistance of the lower part, *pr15*, physically strengthened the lower part of the plant by delaying leaf senescence and non-structural carbohydrate accumulation in culms at the fully ripe stage (Kashiwagi and Ishimaru 2004; Kashiwagi et al. 2006). Culm strength has also been a genetic target for lodging resistance. Ookawa et al. (2010) identified *STRONG CULM2 (SCM2)*, which increased the section modulus of all internodes and the bending moment at breaking of internode IV. A QTL for stem diameter, *sdm8*, thickened the diameters of internodes II and III, and increased culm stiffness (Kashiwagi et al. 2008).

In addition to reducing the moment of upper parts, genetic improvement strategies to increase lodging resistance can focus on increasing the physical strengths of internodes at mid-to-low levels or the lower parts of the plant. Greater physical strength of the plant parts at mid-to-low levels contributes to reducing lodging damage in rice. On the other hand, the upper culms, including internodes I to III, are thinner and weaker than the lower culms, e.g. the basal internode, and are easily bent by environmental stresses and panicle weight at the fully ripe stage. Identification of the genetic factors affecting the physical

strength of upper culms, therefore, would provide a new approach for improving bending-type lodging resistance. In this study, QTL for the physical strength of internodes I to III was identified using chromosomal segment substitution lines (CSSLs). This QTL was evaluated the change in the physical strength of internodes over time after heading, and the effects of the physical strength of the lower part of the plant on lodging.

Materials and methods

Plant materials

For QTL analysis, this study used 39 CSSLs of rice with chromosomal segments of ‘Kasalath’ in the genetic background of ‘Koshihikari’, developed by the Rice Genome Resource Center in Tsukuba, these substituted chromosomal segments covered most of the genome other than small regions at the distal end of the short arm on chromosome 8 and at distal end of the long arm on chromosome 12 (Ebitani et al. 2005). The seeds of these CSSLs and their parents were sown in a greenhouse in late April of 2010 and 2011. Seedlings were transplanted into paddy fields in Utsunomiya, Japan (latitude 36°N), in late May with a single plant per hill spaced at 30 × 30 cm, and were grown under natural conditions with 10 plants per line. The CSSLs and their parents were arrayed in a row, respectively, on one experimental plot in each year. In 2011 and 2012, seeds of three CSSLs with *bsuc11* and ‘Koshihikari’ were sown in late April, and these seedlings were transplanted on 22 and 16 May, respectively, at a spacing of 15 cm between hills and 30 cm between rows. Each plot of CSSL or ‘Koshihikari’ consisted of five rows with 20 hills per row, and the plot position for each line or variety in the paddy field was different in each of the 2 years. N, P₂O₅, and K₂O fertilizers were applied as basal dressing at 6 g m⁻², respectively, before transplanting in 2010–2012.

Physical strength measurements

The breaking strengths of internodes I, II, and III, (first, second, and third internode just below the neck node of the panicle, respectively) were measured a digital force gauge (SX-5, Aikoh Engineering, Osaka, Japan) on a manual test stand. The central part of the

internode was immediately measured at a distance of 6-cm between two supporting points after sampling. Breaking strength was defined as the maximum value at broken internodes. The measurement used three culms with the average culm length per plant, and six plants for each line were analyzed. The pushing resistance of the lower part was measured at 6 weeks after heading with a prostrate tester (Daiki Rika Kogyo Co., Tokyo, Japan), according to the method reported by Kashiwagi and Ishimaru (2004). For pushing resistance of the lower part, the upper part of the plant was removed at a height of 40 cm. The prostrate tester was placed vertically next to the stem at a height of 20 cm above the soil surface, and was pushed until the plants had inclined to an angle of 45° to the vertical. Pushing resistance of the lower part per one tiller (PRL/TN) was calculated using the following equation, PRL/TN = pushing resistance of the lower part/tiller no. per plant. Pushing resistances were determined using 10 plants in each line.

Morphological traits of upper culms

The length and diameter of internodes I, II, and III were measured at 6 weeks after heading. Internode diameter was measured at the major axis at the central part of each internode. As in the case of breaking strength measurement, lengths and diameters of internodes were measured for three culms per plant, and six plants in each line.

QTL analysis

QTLs for breaking strength, length, and diameter of internodes I, II, and III were determined by comparing the phenotypes and genotypes of 130 restriction fragment length polymorphism markers (Ebitani et al. 2005). QTL detection was based on the *t* test of difference between each CSSL and Koshihikari. According to the method of Madoka et al. (2008), the threshold for detection of QTLs was a probability level of 0.05. The effect of a QTL was evaluated as positive (plus) or negative (minus), compared with the ‘Koshihikari’ phenotype.

Statistical analyses

Statistical analyses were conducted with Microsoft Excel 2008 for Mac. Mean and standard error was

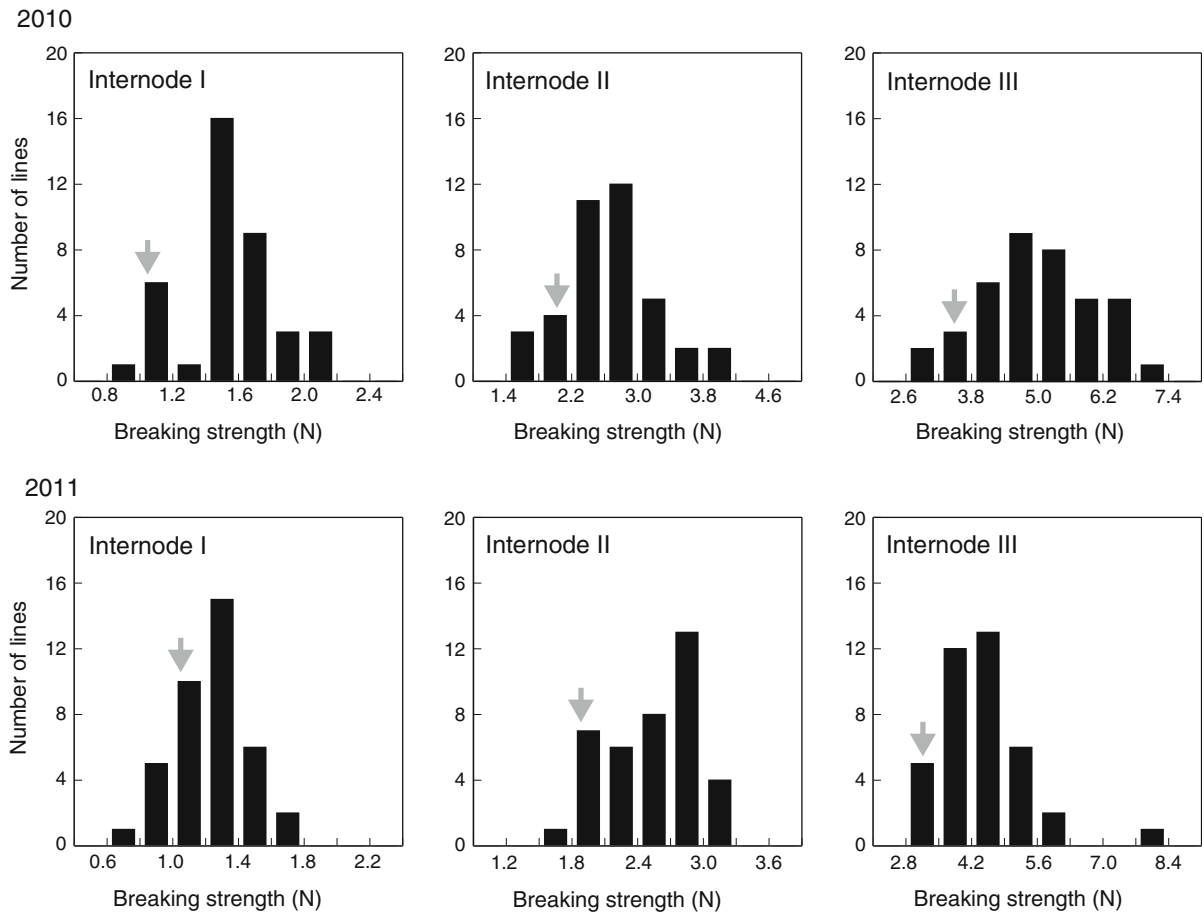


Fig. 1 Frequency distributions of breaking strengths of top three internodes in 39 CSSLs at 6 weeks after heading in 2010 and 2011. Gray arrows indicate the mean values of ‘Koshihikari’

calculated from six or ten plants. For the comparison between each CSSL and Koshihikari, means were separated using the *t* test at a probability of $P = 0.05$ only when the F-test showed significance at 0.05 probability levels.

Results

Breaking strength, length, and diameter of upper culms in CSSLs

The breaking strength of internode I, II, and III in ‘Koshihikari’ was 1.02 ± 0.06 , 2.14 ± 0.16 , and 3.53 ± 0.28 N, respectively, in 2010 (Fig. 1; Table 1). Compared with their corresponding values in ‘Koshihikari’, the breaking strength of internodes I,

II, and III showed a maximum increase of 104, 90, and 94 %, respectively, in the CSSLs. The proportion of CSSLs with a significantly higher breaking strength than that of ‘Koshihikari’ was 85, 59, and 77 % for internodes I, II, and III, respectively. A few CSSLs showed significantly decreased breaking strengths; three CSSL lines showed decreased breaking strengths of internode II, and one showed decreased breaking strength of internode III. In 2011, the breaking strength of ‘Koshihikari’ internodes I, II, and III was 1.06 ± 0.04 , 1.87 ± 0.14 , and 3.34 ± 0.18 N, respectively. The maximum breaking strength of internodes I, II, and III in CSSLs was 68, 65, and 141 % higher, respectively, than its corresponding value in ‘Koshihikari’. Compared with those of ‘Koshihikari’, 56, 74, and 69 % of CSSLs had significantly higher breaking strengths for internodes

Table 1 Breaking strength, length, and diameter of each internode for parents ‘Koshihikari’ and ‘Kasalath’, and for 39 CSSLs

Trait	Koshihikari	Kasalath	Mean 39 CSSLs	SD ^a	Max. ^a	Min. ^a
<i>2010</i>						
Breaking strength (N)						
Internode I	1.02	1.27 (125 % ^b)	1.53	0.28	2.08 (204 %)	0.86 (84 %)
Internode II	2.14	3.49 (163 %)	2.69	0.57	4.06 (190 %)	1.54 (72 %)
Internode III	3.53	5.97 (169 %)	4.98	1.01	6.86 (194 %)	2.94 (83 %)
Length (cm)						
Internode I	39.5	41.3 (105 %)	40.5	3.2	50.2 (127 %)	34.5 (87 %)
Internode II	20.4	33.2 (163 %)	21.4	2.3	32.1 (157 %)	18.4 (90 %)
Internode III	18.7	27.6 (148 %)	16.5	3.0	22.6 (121 %)	10.0 (53 %)
Diameter (mm)						
Internode I	1.96	2.39 (122 %)	2.13	0.19	2.41 (123 %)	1.60 (82 %)
Internode II	3.52	4.63 (132 %)	3.82	0.26	4.35 (124 %)	3.18 (90 %)
Internode III	4.15	5.69 (137 %)	4.43	0.33	5.25 (127 %)	3.47 (84 %)
<i>2011</i>						
Breaking strength (N)						
Internode I	1.06	1.32 (125 %)	1.24	0.23	1.78 (168 %)	0.73 (69 %)
Internode II	1.87	4.13 (221 %)	2.53	0.41	3.10 (166 %)	1.51 (81 %)
Internode III	3.34	7.20 (216 %)	4.45	0.95	8.05 (241 %)	3.11 (93 %)
Length (cm)						
Internode I	35.8	39.5 (110 %)	36.8	2.7	44.3 (124 %)	32.0 (89 %)
Internode II	19.5	30.1 (154 %)	21.0	1.7	24.8 (127 %)	18.6 (95 %)
Internode III	16.5	28.5 (173 %)	16.9	2.8	21.5 (130 %)	9.9 (60 %)
Diameter (mm)						
Internode I	1.89	2.29 (121 %)	1.89	0.19	2.27 (120 %)	1.44 (76 %)
Internode II	3.25	4.54 (140 %)	3.40	0.39	5.37 (165 %)	2.96 (91 %)
Internode III	3.76	5.76 (153 %)	3.91	0.30	4.88 (130 %)	3.26 (87 %)

^a SD, Max., and Min. indicate standard deviation, maximum value, and minimum value in 39 CSSLs

^b %, percentage of the value in Koshihikari

I, II, and III, respectively. The donor parent ‘Kasalath’ showed 1.3-, 1.6-, and 1.7-times higher breaking strength of internodes I, II, and III, respectively, in 2010, and 1.2-, 2.2-, and 2.2-times higher breaking strength of internodes I, II, and III, respectively, in 2011, compared with the corresponding values in ‘Koshihikari’ (Table 1).

In both 2010 and 2011, the lengths of internodes II and III in ‘Kasalath’ were significantly longer than those in ‘Koshihikari’ (Table 1). For each internode, the mean length in the CSSLs was similar to that of the corresponding internode in ‘Koshihikari’. For the CSSLs with the longest internodes, the internode lengths were at least 1.2-times longer than those of

‘Koshihikari’. In the CSSLs with the shortest length of internode III, the length of internode III was 0.6 times or less than that of ‘Koshihikari’. Compared with those of ‘Koshihikari’, the diameters of internodes of ‘Kasalath’ were significantly greater (Table 1). Comparing mean values, the diameters of internodes I, II, and III were similar between ‘Koshihikari’ and the CSSLs. In 2010 and 2011, the CSSLs with maximum internode diameters showed 1.2-, 1.2 to 1.7-, and 1.3-times greater diameters of internodes I, II, and III, respectively, compared with those of the corresponding internodes of ‘Koshihikari’. The smallest internode diameters in the CSSLs were 0.8- to 0.9-times those of the corresponding internodes of ‘Koshihikari’.

Correlations among upper culm traits

For breaking strengths, there were significant positive correlations among internodes ($P < 0.05$, Table 2). The breaking strength of internode I was very strongly correlated with that of internode III ($P < 0.001$). Breaking strength was positively correlated with the length and diameter of internode I ($P < 0.001$). Internode II showed a weak positive correlation between breaking strength and diameter ($P < 0.05$). Breaking strength of internode III negatively correlated with its length, but positively correlated with its diameter ($P < 0.001$). Only internode I showed a significant correlation between length and diameter ($P < 0.001$). The length of internode II was positively correlated with those of internodes I and III ($P < 0.01$), and there were positive correlations among the diameters of the top three internodes ($P < 0.001$).

QTL analysis

This study evaluated nine traits: breaking strengths, lengths, and diameters of the top three internodes. For the nine traits, this study detected 65 and 85 QTLs in 2010 and 2011, respectively (Table 3, 4). In 2010, nine and eight QTLs for breaking strength of internodes I and III, respectively, had positive effects from the ‘Kasalath’ allele (Table 3). With respect to the QTLs for breaking strength of internode II, eight alleles from ‘Kasalath’ showed positive effects, and two alleles on chromosomes 1 and 4 decreased the breaking strength. ‘Kasalath’ alleles for increased breaking strength of the top three internodes were detected on chromosomes 7, 10, 11, and 12, while those for increased strength of specific internodes were detected on chromosomes 1, 2, and 3. In 2011, seven, eight, and eight QTLs for breaking strength of internodes I, II, and III, respectively, showed positive effects from the ‘Kasalath’ allele (Table 4). Three ‘Kasalath’ alleles on chromosomes 3, 4, and 5 contributed a decrease in the breaking strength of internode I. On chromosomes 1, 7, and 11, there were regions associated with increased breaking strength of the top three internodes. QTLs on chromosomes 5, 9, and 10 contributed to strength of specific internodes.

Comparing the QTLs detected in the 2 years, 21 corresponding QTLs were found for breaking strengths, lengths, and diameters of the top three internodes (Fig. 2). On chromosome 11 (marker

Table 2 Correlation coefficients between breaking strength, length, and diameter of internodes I, II, and III in CSSLs

	Internode I BS	Internode II BS	Internode III BS	Internode I L	Internode II L	Internode III L	Internode I D	Internode II D
Internode II BS ^a	0.274*							
Internode III BS	0.682***	0.236*						
Internode I L	0.585***	0.318**	0.405***					
Internode II L	0.024	-0.030	0.123	0.414***				
Internode III L	-0.358**	-0.328**	-0.548***	-0.194	0.317**			
Internode I D	0.872***	0.266*	0.550***	0.650***	0.163	-0.245*		
Internode II D	0.683***	0.271*	0.518***	0.541***	0.203	-0.142	0.766***	
Internode III D	0.693***	0.127	0.570***	0.543***	0.310**	-0.041	0.816***	0.813***

***, **, * significant at $P < 0.001$, 0.01, and 0.05, respectively

^a BS, L, and D indicate breaking strength, length, and diameter, respectively

Table 3 QTLs for breaking strength, length, and diameter of internodes I, II, and III at the fully ripe stage in 2010

Traits	Number of QTLs detected	Chromosome number	Nearest markers	Effect from 'Kasalath' allele ^a		
Breaking strength of internode I	9	1	S13849-R2635	+		
		2	C499-C747	+		
		3	C515-C25	+		
		5	R2232	+		
		7	R1357-C596	+		
		9	R1751-C1263	+		
		10	C1286	+		
		11	C535-G257	+		
		12	G24B	+		
		Breaking strength of internode II	10	1	R607-C178	+
				1	C112	–
				2	C1419-C1470	+
3	S1513-R663			+		
4	R374-C1016			–		
6	R2171-R2549			+		
7	R1357-C597			+		
10	C1286			+		
11	C535-G257			+		
12	G24B			+		
Breaking strength of internode III	8			3	S1513-R663	+
				5	R2232	+
		6	R2171-R2549	+		
		7	R1357-C596	+		
		9	R1751-C1263	+		
		10	C1286	+		
		11	C535-G257	+		
		12	G24B	+		
		Length of internode I	6	1	R607-C178	–
				3	G332-S1571	–
				4	R374-C1016	–
				5	C246-C1230	–
10	C1286			+		
12	S10637A			+		
Length of internode II	7	1	R2417-C472	+		
		2	C499-C747	+		
		3	G322-S1571	–		
		4	R2373-C891	+		
		5	R566-C128	–		
		8	R1943	+		
		12	S10637A	+		

Table 3 continued

Traits	Number of QTLs detected	Chromosome number	Nearest markers	Effect from 'Kasalath' allele ^a
Length of internode III	8	2	C1357-C132	–
		3	G332-S1571	–
		5	R2232	–
		7	R1357-C596	–
		9	R1751-C1263	–
		10	C1286	–
		11	C535-G257	–
		12	R1709-R1684	+
Diameter of internode I	5	3	G332-S1571	–
		7	R1357-C596	+
		8	C347-S11322	+
		9	C1454-R79	+
		10	C1286	+
Diameter of internode II	8	3	G332-S1571	–
		5	C597-R3166	+
		7	R1357-C596	+
		8	C347-S11322	+
		9	R1751-C1263	+
		10	C1286	+
		11	C535-G257	+
		12	S10637A	+
Diameter of internode III	4	3	G332-S1571	–
		8	C347-S11322	+
		9	C1454-R79	+
		12	S2572-R3375	+

^a + and – indicate positive and negative effects, respectively, from 'Kasalath' allele

C535–G257), there was the only chromosomal region (*bsuc11*) associated with a simultaneous increase in the breaking strengths of the top three internodes. QTLs for breaking strength of specific internodes were detected on chromosomes 2 and 9. Two chromosomal regions on chromosomes 10 and 12 improved breaking strengths of internode II and III, and of internode I and II, respectively. A QTL for elongation of internode I was detected on chromosome 10, and three QTLs for elongation of internode II were detected on chromosomes 1, 2, and 12. A chromosomal region associated with simultaneous shortening of internodes I and III was found on chromosome 3. Three QTLs on chromosomes 7, 9, and 11 were related to an increase in the diameter of internode II.

Effects of *bsuc11* on lodging resistance characteristics

In 2011 and 2012, this study analyzed three CSSLs containing *bsuc11*; CSSL233, 234, and 235, and compared the breaking strengths of their top three internodes with those of 'Koshihikari' after heading (Fig. 3). Internode I in 'Koshihikari' showed an increase in breaking strength from heading to 4 weeks after heading (WAH) in 2011, and the breaking strengths of internode II and III increased until 2 WAH. After 6 WAH, the top three internodes of 'Koshihikari' showed a decrease in breaking strength. CSSLs with *bsuc11* showed increased breaking strengths of internode I at 6 WAH; their breaking

Table 4 QTLs for breaking strength, length, and diameter of internodes I, II, and III at the fully ripe stage in 2011

Trait	Number of QTLs detected	Chromosome number	Nearest markers	Effect from 'Kasalath' allele ^a		
Breaking strength of internode I	10	1	C122-C1370	+		
		2	C499-C747	+		
		3	S1513-R663	–		
		4	R374-C1016	–		
		5	C246-C1230	–		
		7	C213	+		
		9	S2676	+		
		10	C701-R2174	+		
		11	C535-G257	+		
		12	G24B	+		
		Breaking strength of internode II	8	1	C122-C1370	+
				3	G332-S1571	+
5	R2232			+		
7	C213			+		
9	R1751-C1263			+		
10	C1286			+		
11	C535-G257			+		
12	G24B			+		
Breaking strength of internode III	8	1	C122-C1370	+		
		2	C499-C747	+		
		3	G332-S1571	+		
		5	R556-C128	+		
		7	C213	+		
		9	R1751-C1263	+		
		10	C1286	+		
		11	C535-G257	+		
		Length of internode I	11	1	R607-C178	–
				1	R2417-C472	+
				2	C1419-C1470	+
3	C515-C25			+		
3	G332-S1571			–		
4	C445			–		
5	C246-C1230			–		
7	C213			+		
10	C1286			+		
11	R77-G320			–		
12	R1709-R1684			+		
Length of internode II	9			1	R2417-C472	+
		2	C499-C747	+		
		3	S1513-R663	+		
		4	C513	+		
		6	C556-R1167	+		
		7	R1357-C596	+		
		8	C390-C1121	+		

Table 4 continued

Trait	Number of QTLs detected	Chromosome number	Nearest markers	Effect from 'Kasalath' allele ^a
Length of internode III	14	11	R77-G320	–
		12	S10637A	+
		1	R607-C178	+
		1	C112	+
		2	C499-C747	+
		3	C515-C25	+
		3	G332-S1571	–
		4	C513	+
		5	C246-C1230	+
		6	C10038-S1520	+
		6	C556-R1167	+
		7	C213	–
		8	C347-S11322	+
		9	R79-C1263	–
Diameter of internode I	8	10	C223-G127	–
		12	S2572-R3375	+
		1	R2417-C472	+
		2	C1357-R712	–
		3	S1513-R663	–
		4	C513	–
		5	C246-C1230	–
		6	C556-R1167	+
Diameter of internode II	10	8	R1943	–
		11	C535-G257	+
		1	C122-C1370	+
		2	C1357-R712	+
		3	S1513-R663	–
		4	C513	–
		6	R2171-R2549	+
		7	R1357-C596	+
		8	R1943	–
		9	R1751-C1263	+
		10	C223-G127	+
Diameter of internode III	7	11	C535-G257	+
		1	R2417-C472	+
		2	C1357-R712	–
		3	S1513-R663	–
		8	R1943	–
		9	R1751-C1263	+
		11	R77-G320	+
		12	G24B	+

^a + and – indicate positive and negative effect, respectively, from 'Kasalath' allele

strengths were 1.61- to 1.78- times higher than that of internode I of 'Koshihikari' at corresponding time points ($P < 0.01$). In CSSLs with *bsuc11*, internode II

showed significantly higher breaking strengths at 4 and 6 WAH than those of internode II of 'Koshihikari' at corresponding time points ($P < 0.05$, 1.19–1.46

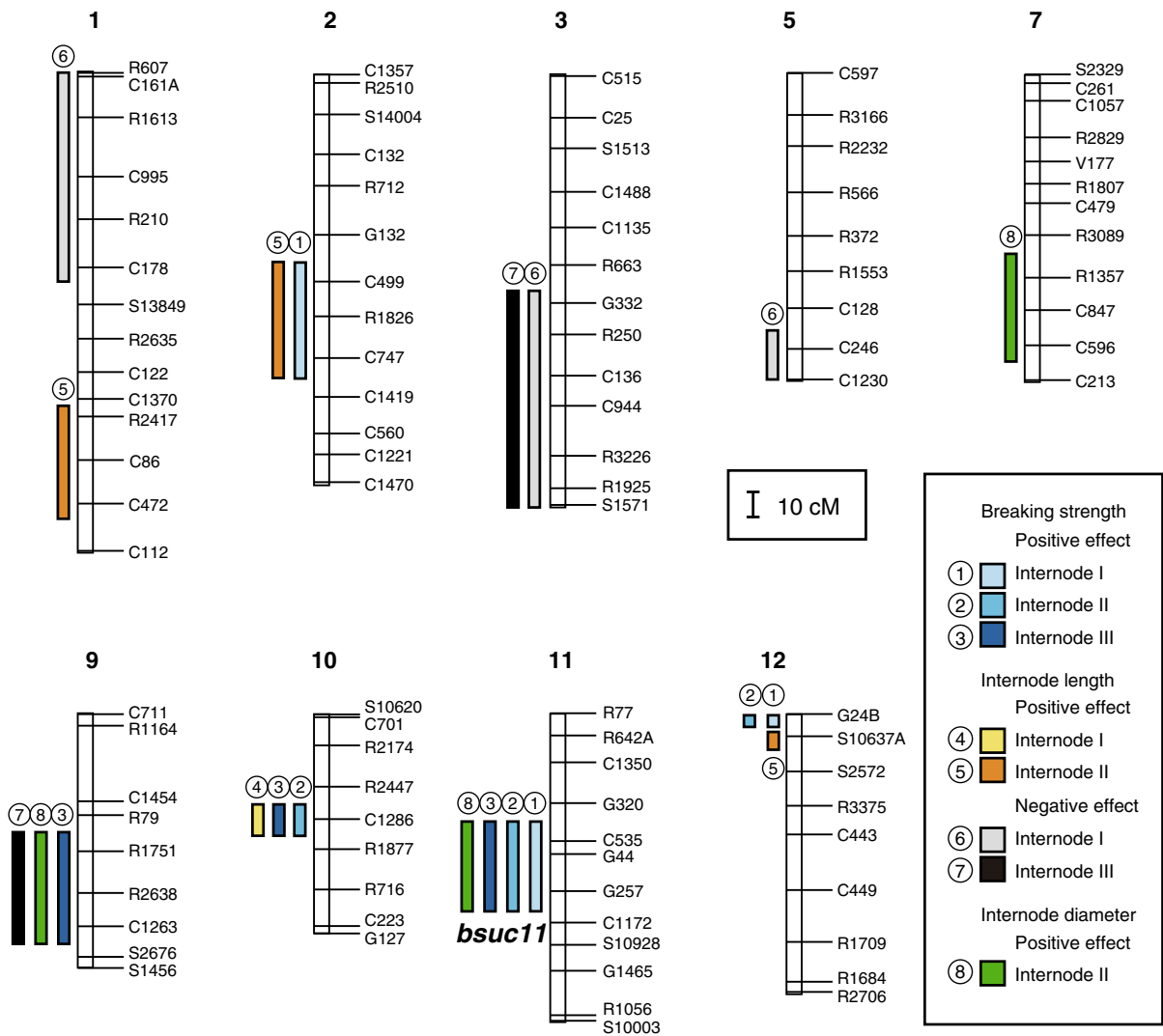


Fig. 2 Positions of QTLs for breaking strengths, lengths, diameters of top three internodes on the rice genetic map. *Data* indicate corresponding QTLs between 2010 and 2011. RFLP markers were defined by Ebitani et al. (2005)

times higher at 4 WAH, and 1.58–1.67 times higher at 6 WAH). For internode III, CSSLs with *bsuc11* showed higher breaking strengths at heading and 6 WAH, compared with those of ‘Koshihikari’ at corresponding time points ($P < 0.05$, 1.39–1.49 times higher at heading, 1.49–1.65 times higher at 6 WAH). In 2012, ‘Koshihikari’ showed increased breaking strength of only internode I until 2 WAH, and then the breaking strength of the top three internodes decreased after 2 WAH. At heading and 6 WAH, CSSLs with *bsuc11* showed 1.42–1.69 times higher and 1.21–1.44 times higher breaking strength of internode I,

compared with those of ‘Koshihikari’ at corresponding time points. The breaking strength of internode II gradually decreased after heading in CSSLs with *bsuc11*, but these lines showed higher breaking strengths than those of ‘Koshihikari’ from 4 WAH to 6 WAH (1.20–1.26 times higher and 1.22–1.46 times higher, respectively). Internode III showed the same breaking strength in CSSLs with *bsuc11* and ‘Koshihikari’ from heading until 4 WAH, but the internode breaking strengths in these CSSLs were 1.10–1.31 times higher than that of internode III of ‘Koshihikari’ at 6 WAH.

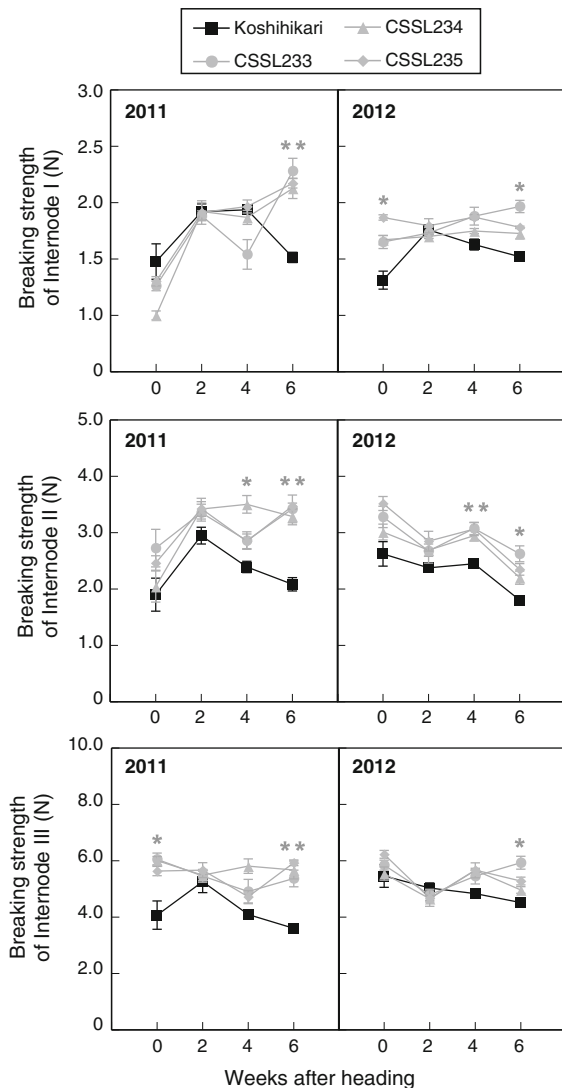


Fig. 3 Changes in breaking strengths of top three internodes in ‘Koshihikari’ (black line) and CSSLs with *bsuc11* (gray lines) after heading. Data are means of six plants in each line. Vertical bars indicate standard errors. Double asterisk and single asterisk indicate significant differences at 0.01 and 0.05 probability levels, respectively, between ‘Koshihikari’ and three CSSLs with *bsuc11*

The pushing resistance of the lower part (PRL) in ‘Koshihikari’ was $7.15 \pm 0.57 \text{ kPa cm}^{-2}$, and the PRL per one tiller (PRL/TN) was $0.56 \pm 0.06 \text{ kPa cm}^{-2}$ in 2011 (Table 5). Two CSSLs (CSSL234 and 235) showed significant decreases in PRL compared with that of ‘Koshihikari’ ($P < 0.05$), while the values for PRL/TN were the same in ‘Koshihikari’ and CSSLs with *bsuc11*. For ‘Koshihikari’ in 2011, the values of PRL and PRL/TN were 6.10 ± 0.31 and

$0.53 \pm 0.03 \text{ kPa cm}^{-2}$, respectively. Only CSSL233 showed a significantly lower PRL than that of ‘Koshihikari’ ($P < 0.05$), but three CSSLs showed similar PRL/TN values to that of ‘Koshihikari’.

Discussion

This study identified one QTL, *bsuc11*, for breaking strengths of the top three internodes at the fully ripe stage over the 2 years, which did not relate to the internodes length (Fig. 2). The donor parent ‘Kasalath’ had greater diameters and breaking strengths of the top three internodes, but these internodes were long (Table 2). These properties in ‘Kasalath’ were inherited by the CSSLs in the genetic background of ‘Koshihikari’, and the CSSLs with maximum breaking strengths of top three internodes showed greater strengths than those of ‘Kasalath’. Thus, ‘Kasalath’ would be a valuable genetic resource for physical strength of the upper culms, at least in the Koshihikari background. From the results of QTL analysis, ‘Kasalath’ had QTLs to strengthen specific internodes and to simultaneously improve the physical strengths of plural internodes or the top three internodes (Tables 3, 4). To improve lodging resistance, it is important that culm strength is genetically controlled by two types of QTL. The most useful QTLs for breeding will be those that improve the physical strengths of plural internodes. *bsuc11* physically strengthened the top three internodes, and did not overlap with QTLs for internode elongation (Fig. 2). Therefore, *bsuc11* would efficiently improve resistance to bending in upper culms and could be a useful genetic target to reduce bending-type lodging.

Culm diameter is one of the traits that contributes to physical strength, and thus to lodging resistance (Atkins 1938; Zuber et al. 1999; Kashiwagi et al. 2008). In this study, internode diameter was correlated with breaking strength (Table 2), and QTLs for diameter partly overlapped with those for breaking strength of each internode (Tables 3, 4). These results indicate that some QTLs for internode diameter would contribute to significant correlations between diameters and breaking strengths of internodes, but all QTLs to increase in culm diameter do not necessarily improve lodging resistance. In a previous study, a QTL for stem diameter in ‘Kasalath’, *sdm8*, increased the culm wall thickness and physical strength of internodes in ‘Nipponbare’ (Kashiwagi et al. 2008). In

Table 5 Pushing resistances of the lower part of ‘Koshihikari’ and CSSLs with *bsuc11*

	Koshihikari	CSSL233	CSSL234	CSSL235
2011				
PRL (kPa cm ⁻²)	7.15 ± 0.57	5.80 ± 0.86	5.12 ± 0.71*	5.29 ± 0.44**
PRL/TN (kPa cm ⁻²)	0.56 ± 0.06	0.55 ± 0.07	0.51 ± 0.05	0.50 ± 0.02
2012				
PRL (kPa cm ⁻²)	6.10 ± 0.31	5.13 ± 0.44*	6.35 ± 0.46	5.29 ± 0.52
PRL/TN (kPa cm ⁻²)	0.53 ± 0.03	0.49 ± 0.03	0.58 ± 0.03	0.51 ± 0.04

Data are mean ± SE ($n = 10$ plants)

PRL, pushing resistance of the lower part; PRL/TN, PRL per one tiller

** and * indicate significant differences at 0.01 and 0.05 probability levels, respectively

this study, *bsuc11* overlapped with a QTL to increase the diameter of internode II in the QTL analyses in both years (Fig. 2). Therefore, the improvement in the breaking strengths of the top three internodes by *bsuc11* might be partly related to morphological changes in culm diameter and culm wall thickness.

In terms of properties associated with higher lodging resistance, *bsuc11* prevented the decrease in the breaking strengths of top three internodes after grain filling, rather than increasing the strengths of internodes from heading (Fig. 3). Takaya and Miyasaka (1983) reported that breaking strengths of culms decreased after heading. The top three internodes of ‘Koshihikari’ showed decreases in breaking strength after 2 WAH. However, lines with *bsuc11* did not show this decrease in internode strength after 2 WAH, and therefore showed greater physical strength than that of ‘Koshihikari’ at the fully ripe stage (Fig. 3). These results indicated that the decrease in culm strength after heading contributes to bending-type lodging; thus, preventing this decrease in strength would be a good target to improve lodging resistance. This study evaluated the effects of *bsuc11* on the physical strength of the lower part of the plant. This study found that *bsuc11* increased the physical strength of the upper culms but did not improve the pushing resistance of the lower part (Table 5). This result implies that the physical strengths of upper and lower parts of rice plants are controlled by different genetic factors, and the improvement of both strengths could develop cultivars with greater lodging resistance.

A previous study on the genetic factors associated with lodging resistance identified some QTLs and genes in rice. Ookawa et al. (2010) revealed that *SCM2*

on chromosome 6, which was identical to *ABERRANT PANICLE ORGANIZATION (APO1)*, enhanced culm strength at 20 days after heading and increased the diameters of the minor and major axes of all internodes. From this result, *SCM2* could be expected to increase the maximum strength of culms after heading. This effect differs from the improvement in culm strength associated with *bsuc11*, because the CSSLs with *bsuc11* had similar internode breaking strengths to those of ‘Koshihikari’, when ‘Koshihikari’ showed maximum culm strength (Fig. 3). Therefore, there are two approaches to improve culm strength; to increase the maximum strength of the culm, or to prevent the decrease in culm strength after heading. The QTL for lodging resistance in a typhoon on chromosome 5, *lrt5*, increased young’s modulus of internode I and II, but the stem diameter was the same as that of the control (Ishimaru et al. 2008). This QTL physically strengthened the upper culms but did not increase the PRL, an indicator of the physical strength of the lower parts. This study found that ‘Koshihikari’ and the CSSLs with *bsuc11* had the same PRL at the fully ripe stage (Table 5), suggesting that *bsuc11* would be similar to *lrt5* in terms of its effect on lodging resistance. *lrt5* delayed senescence in the upper leaves and resulted in high starch content in the upper culms. Non-structural carbohydrate (NSC) re-accumulation is an indicator of breaking-type lodging resistance in rice (Yagi 1983). Other components of culms, e.g., cellulose, lignin, and silicon, also contribute to the physical strength of the plant body (Kokubo et al. 1991; Jones et al. 2001; Li et al. 2003; Idris et al. 1975). Because *bsuc11* did not affect the diameters of all internodes, the greater culm strength associated with *bsuc11* might result from differences in culm components.

This study showed that *bsuc11* prevented the decrease in strengths of the upper culms after heading. The improved culm strength includes the thick culm diameter or culm wall and the changes in culm components. Genetic information about culm strength will provide a new approach in breeding for lodging resistance. To clarify the function of *bsuc11*, further research is required to reveal the morphological characters and components of culms, and to identify the gene locus. These would be key findings to improving the lodging resistance of rice.

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