Supplemental Data. Ueguchi-Tanaka et al. (2008). Release of the Repressive Activity of Rice DELLA Protein SLR1 by Gibberellin Does Not Require SLR1 Degradation in the *gid2* mutant.

Supplemental Table 1. Comparison of the 2nd and 3rd Leaf Sheath Length among Various GA-Deficient and GA-Insensitive Rice Mutants

Genotype	2nd leaf sheath length (mm)	3rd leaf sheath length (mm)
WT	30.0 ± 1.5	62.3 ± 6.0
d35	23.8 ± 0.9	41.9 ± 1.2
d18 (WC)	10.9 ± 1.3	18.1 ± 1.6
cps1-1	7.4 ± 1.1	12.0 ± 0.7
gid1-8	20.9 ± 1.5	40.0 ± 1.7
gid1-7	13.4 ± 1.8	20.9 ± 2.3
gid1-3	5.6 ± 0.6	7.1 ± 1.0
gid2-1	9.7 ± 0.9	15.3 ± 1.0
gid2-2	9.3 ± 0.9	15.2 ± 0.9
gid2-5	10.6 ± 0.8	15.4 ± 0.3

All plants were grown for 2 weeks.

The measurements are the means ± SE (n = 8 to 14).

Supplemental Table 2. Comparison of the 4th Leaf Sheath Length among WT, *gid2* and *gid1*, Treated with/without Uniconazole and GA_3

Genotype	uniconazole	GA_3	4th leaf sheath length (mm)
	-	-	110.6 ± 6.9
WT	+	-	29.1 ± 2.7
	+	+	90.3 ± 23.1
	-	-	19.7 ± 0.9
gid2-2	+	-	8.3 ± 0.8
	+	+	15.0 ± 1.5
	-	-	8.8 ± 0.9
gid1-3	+	-	8.1 ± 1.0
	+	+	8.1 ± 0.3

All plants were grown for 3 weeks.

The measurements are the means ± SE (n = 8 to 14).

Supplemental Table 3. Comparison of the 4th and 6th Leaf Sheath Length among WT, *gid2*, *cps1* and *gid2 cps1*

Genotype	4th leaf sheath length (mm)	6th leaf sheath length (mm)
WT	96.9 ± 11.0	134.3 ± 8.8
gid2-2	20.1 ± 1.0	30.5 ± 2.0
cps1-1	7.5 ± 0.5	9.2 ± 0.8
gid2-2 cps1-1	8.3 ± 1.0	9.5 ± 0.5

All plants were grown for 4 weeks.

The measurements are the means ± SE (n = 6 to 10).

Supplemental Table 4. Comparison of the 4th and 6th Leaf Sheath Length among WT, and Plants with Mutant Alleles at the *GID1* and/or *GID2* Loci

Genotype		4th leaf sheath length (mm)	6th leaf sheath length (mm)
GID1	*GID2*		
+/+	+/+	104.6 ± 4.7	142.8 ± 8.8
+/+	-/-	19.9 ± 2.1	31.1 ± 1.7
+/-	-/-	14.6 ± 2.1	18.4 ± 0.9
-/-	-/-	8.0 ± 0.7	9.8 ± 1.3
-/-	-/+	8.0 ± 0.6	9.1 ± 1.1
-/-	+/+	7.7 ± 1.0	10.0 ± 1.1

All plants were grown for 4 weeks.

The measurements are the means ± SE (n = 6 to 10).

Supplemental Table 5. Primers Used in This Study

Primer	Sequence (5' to 3')	Note
gid1–26U	TAGTAGAGGGTCGCTCACTG	genotyping for *gid1–3* mutant
gid1–26L	GGACACCACGACGCCCTTGC	
gid2–2U	AGGTTCAGCTCTCGCTGTCA	genotyping for *gid2–2* mutant
gid2–2L	TTGGAGTGAAGCTTATTCCA	
cpsTos1U	TCCAATGCAGTGATGGATCC	genotyping for *cps1–1* mutant
LTR4A	ACTGTATAGTTGGCCCATGTCCAG	(insertion of Tos17)
cpsTos1U	TCCAATGCAGTGATGGATCC	genotyping for *cps1–1* mutant
cpsTos1L	TGGAAGTACCGCGATATCCC	(no insertion of Tos17)
HA–SLR1–U	GGTCTAGAT AAT ACG ACTCAC TAT AGG GC	cloning for HA–SLR1
HA–SLR1–L	CGCCCGGGTGGAATTCACGCCGCGGCGACG	