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**Nucleotide sequences of two genes encoding the light harvesting chlorophyll a/b binding protein of rice**


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Southern blots suggest that the rice genome contains about four genes coding for light harvesting chlorophyll a/b binding protein (LHCP). From a genomic library kindly provided by Dr. Susan Wessler at the University of Georgia, we have cloned and sequenced two of the four genes. Here we report the nucleotide sequences of the coding regions and some of 3' and 5' flanking regions of these genes, *cab1R* and *2R*. The coding regions for mature proteins are very similar to one another and also to those of LHCP genes from wheat(1) and pea(2). The presumed transit sequences and flanking sequences of the two genes differ substantially.

	1	10	20	30	40	50	60	70											
1R	GCACATCAGCTCC	<u>AA</u> T	CTCCC	GGCCC	CAGGAGG	GAATCCCC	CTCGCTT	CCCGCGCT	<u>TATT</u> AAACT	CCCGCGC									
2R	cggCc	ccatc	<u>Cag</u>	<u>CAA</u> ATC	Cgac	Ccagcaa	Aatgt	TCCC	aa	Gat	<u>CTATT</u> a	<u>T</u> Acg	CcaagGt						
									cgct	ctccacg									
1R	CAACTCCC	AACT	CAC	ACTCG	CGCTCG	CATCG	CCCATCT	CTCTCAG	GTCTCACAG	CTCACTG	GATCA	*****	*****						
2R	gttCgt	cgcca	Tg	ACA	cagcac	aca	CaCC	cag	Cagcag	Cagc	Ag	Cag	Ttgaagctgctga						
1R	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****	*****						
2R	gcgagg	tagac																	
1R	ATG	GCC	GCG	GCC	ACC	ATG	GCG	CTC	TCC	TCC	CGG	GTG	ATG	GCC	CGC	GCG	GCG	CCG	TCG
	M	A	A	A	T	M	A	L	S	S	P	V/a	M/1	A	R/g	A/k	A	P/a	S/a
2R	ATG	GCC	GCC	GCC	ACC	ATG	GCc	CTC	TCg	TCC	CCG	GcG	cTG	CCC	gGC	aaG	GCc	gcc	gCG
1R	ACC	TCC	TCC	GCG	CTC	TTC	GGC	GAG	GCG	CGG	ATC	ACC	ATG	CGC	AAG	ACC	GCC	GCG	AAG
	T	S/k	S	A	L	F	G	E	A/g	R	I	T	M	R	K	T/s	A	A	K
2R	***	aag	***	***	GtG	TTC	GGC	GAG	GgG	CGc	ATC	ACC	ATG	CGC	AAG	tCg	GCg	GCG	AAG
1R	CCC	AAG	CCG	GCG	GGC	TCG	TCG	GGG	AGC	CCG	TGG	TAC	GGC	CCC	GAC	CGC	GTC	CTC	TAC
	P	K	P	A	A	S	S	G	S	P	W	Y	G	A	D	R	V	L	Y
2R	CCC	AAG	CCc	Gcc	GCG	TCG	TCG	GGG	AGC	CCG	TGG	TAC	GGC	GCC	GAC	CGC	GTg	CTC	TAC
1R	CTC	GGC	CCG	CTC	TCG	GGC	CGC	GAG	CCG	CCG	AGC	TAC	CTC	ACC	GGC	GAG	TTC	CCG	GGC
	L	G	P	L	S	G	R	E	P	P	S	Y	L	T	G	E	F	P	G
2R	CTC	GGC	CCG	CTC	TCc	GGC	CGC	GAG	CCG	CCG	AGC	TAC	CTg	ACC	GGC	GAG	TTC	CCC	GGC
1R	GAC	TAC	GGG	TGG	GAC	ACC	GCG	GGG	CTC	TCC	GCC	GAC	CCG	GAG	ACG	TTC	GCC	AAG	AAC
	D	Y	G	W	D	T	A	G	L	S	A	D	P	E	T	F	A	K	N
2R	GAC	TAC	GGG	TGG	GAC	ACC	GCG	GGG	CTC	TCC	GCC	GAC	CCG	GAG	ACG	TTC	GCC	AAG	AAC
1R	CGG	GAG	CTG	GAG	GTG	ATC	CAC	TCC	CGG	TGG	GCG	ATG	CTG	GGC	GCG	CTC	GGC	TGC	GTC
	R	E	L	E	V	I	H	S	R	W	A	M	L	G	A/g	L	G	C	V
2R	CGG	GAG	CTG	GAG	GTG	ATC	CAC	TCC	AGG	TGG	GCG	ATG	CTc	GGC	GgG	CTg	GGC	TGC	GtG
1R	TTC	CCG	GAG	CTC	CTC	GCC	CGG	AAC	GCG	GTC	AAG	TTC	GGC	GAG	GCC	GTG	TGG	TTC	AAG
	F	P	E	L	L	A	R	N	G	V	K	F	G	E	A	V	W	F	K
2R	TTC	CCG	GAG	CTC	CTC	GCC	CGc	AAC	GGC	GTC	AAC	TTC	GGg	GAG	GCg	GTG	TGG	TTC	AAG

1R GCG GGC TCG CAG ATC TTC AGC GAG GGC GGG CTC GAC TAC CTC GGC AAC CCG ACC CTG  
A G S Q I F S E G G L D Y L G N P S L  
2R GCG GGg TCG CAG ATC TTC AGC GAG GGC GGG CTC GAC TAC CTC GGC AAC CCG AGC CTG

1R ATC CAC GCG CAG AGC ATC CTC GCC ATC TGG GCG GTG CAG GTG GTG CTC ATG GGC GCC  
I H A Q S I L A I W A V Q V V L M G A  
2R ATC CAC GCG CAG AGC ATC CTC GCC ATC TGG GCG GTG CAG GTG GTG CTC ATG GGC GCC

1R GTC GAG GGG TAC CGG ATT GCT GGC GGG CCG CTC GGC GAG GTC GTC GAC CCG CTG TAC  
V E G Y R I A G G P L G E V V D P L Y  
2R GTC GAG GGG TAC CGG ATT GCc GGC GGG CCG CTC GGC GAG GTC GTC GAC CCG CTC TAC

1R CCC GGC GGC GCC TTC GAC CCG CTC GGC CTC GCC GAT GAC CCC GAG GCG TGC GCG GAG  
P G G A F D P L G L A D D P E A C/f A E  
2R CGg GGC GGC GCC TTC GAC CCG CTC GGC CTC GCC GAT GAC CCg GAG GCc TtC GGG GAG

1R CTC AAG GTG AAG AAG ATC AAG AAC GGC CGC CTC GCC ATG TTC TCC ATG TTC GGC TTC  
L K V K K/e I K N G R L A M F S M F G F  
2R CTC AAG GTG AAG gAG ATC AAG AAC GGC CGC CTC GCC ATG TTC TCC ATG TTC GGC TTC

1R TTC GTC CAG GCC ATC GTC ACC GGC AAG GGC CCC CTC GAG AAC CTC GCC GAC CAC CTC  
F V Q A I V T G K G P L E N L A D H L  
2R TTC GTC CAG GCC ATC GTC ACC GGC AAG GGC CCC CTC GAG AAC CTC GCC GAC CAC CTC

1R GCC GAC CCC GTC AAC AAC AAC GGC TGG GCC TAC GCC ACC AAC TTC GTC CCC GGC AAG  
A D P V N N A W A Y A T N F V P G K  
2R GCC GAC CCC GTC AAC AAC AAC GGC TGG GCC TAC GCC ACC AAC TTC GTC CCC GGC AAG

1R TGA  
-  
2R TGA  
940  
1R AGTGGGGACCGTAGCTTAGCAGTGGTTAAATTGTGGTGGATGGATTGTGGCCAGCGAGTTGGTGTCTTGGGT  
2R gcgccGccgCcgcgtgctgccatGgcgAcgcatgcctcaGcTaaGctagCtagGttgaCGacGctgcccGtc  
964  
1089  
1R TGGGGAAAGATGGTTAGTGGCAGGAGATGATGATCGAGTTGGTGGTGTACACTAAGAAGATGAAGAAGAAG  
2R TctGcAgGagaGtgTgcGTgtGtgtAcgcGgtGcagtAGaTGtacGggGgaTctgGccacgtcatctacAtgcc  
1113

**Comparing cablR with 2R:** The CAAT and TATA sequences are underlined. Small letters indicate amino acid/nucleotide difference. \*\*\* represents presumed deletions.

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