Supplementary Materials: Molecular Characterization and Biological Effects of a C-Type Lectin-Like Receptor in Large Yellow Croaker (*Larimichthys crocea*)

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										GA	AGT	TGA	AGT	CTC	TGC	CAAA	СТС	ACA	AAA	AA
1	ATGA	ATG	AAC	CTC	ATC	AAA	AAG	AAA	GGT	GCC	CTG	GGA	ATC	AGA	GGA	CAC	GCA	GTC	СТС	TTT
1	М	М	Ν	L	Ι	Κ	Κ	K	G	А	L	G	Ι	R	G	Н	А	V	L	F
61	GTT	ГТС	ATC	AGC	CTT	ATG	GTT	ТСТ	GTA	GCT	GCA	GAC	AGT	GCA	GAG	GGA	CAG	GAG	TCC	TAT
21	V	F	Ι	S	L	М	V	S	V	А	А	D	S	А	Е	G	Q	Е	S	Y
121	CTA	AAG	CTG	AGA	СТА	GAC	CTT	TTG	AAG	AAA	AGC	TAC	AGT	AAG	CTG	TGC	AGT	GAA	TAC	ACC
41	L	Κ	L	R	L	D	L	L	K	Κ	S	Y	S	Κ	L	С	S	E	Y	Т
181	AACO	CTG	GCA	CAG	AAC	TGC	TCA	GTT	CCA	GTG	CCT	AAC	TGT	TAC	GAG	TGT	ССТ	GAC	GAC	AAA
61	Ν	L	А	Q	Ν	С	S	V	Р	V	Р	Ν	С	Y	Е	С	Р	D	D	Κ
241	TGGG	CTT	СТА	GTA	GGG	GAC	CAA	TGC	CTC	CTC	CTC	GAG	ACT	GAC	AGG	AAC	GAC	TGG	CTT	AAT
81	W	L	L	V	G	D	Q	С	L	L	L	Е	Т	D	R	Ν	D	W	L	N
301	AGT	ГСА	AAA.	AAG	TGT	GAA	GAG	ATG	GGC	GCC	CAT	CTT	GCC	ATC	TTG	ACC	ACC	ACA	GAA	CAG
101	S	S	K	K	С	Е	E	М	G	A	Н	L	A	Ι	L	Т	Т	Т	E	Q
361	CATO	GAA	GCT	GTG	GAG	AAA	GAA	GGC	AGA	ATG	CTC	GGT	GGG	TTA	TAC	ACA	TTC	TAC	TGG	ATG
121	Н	E	А	V	Е	Κ	Е	G	R	М	L	G	G	L	Y	Т	F	Y	W	M
421	GGAG	CTG	ACT	GAC	ATT	GAG	AAA	GAA	GGA	GAG	TGG	AAA	TGG	GTG	GAC	AAC	TCA	ATA	GTT	AAA
141	G	L	Т	D	Ι	E	K	E	G	Е	W	K	W	V	D	Ν	S	Ι	V	K
				٠		٠														
481	AACA	ACA	CAC	TGG	AAG	GTT	GGG	ACA	TCA	GAA	CCA	GAC	AAC	AAC	CAG	TCT	GGT	GGG	GAA	GAG
161	N	Т	Н	W	K	V	G	Т	S	E	Р	D	Ν	N	Q	S	G	G	E	E
	_									\diamond		\$		•						
541	GGA	GAG	GAC	TGT	GCG	GTA	GTG	GAC	AGC	TAC	ACT	CAG	AGC	TGG	ГАС	GAT	GTT	CCC	TGC	TCC
181	G	E	D	С	A	V	V	D	S	Y	Т	Q	S	W	Y	D	V	Р	С	S
		\diamond	•												\$	\$				104757070422.0
601	TAC	ГТG	TAT	CCA	CGA	ATC	TGT	CAG	GTC	AAC	GCC	AAA	CTG	CTC	AAG	TGA	GCC	ТСТ	CCG	CAC
201	Y	L	Y	Р	R	Ι	С	Q	V	Ν	A	Κ	L	L	Κ	*				
661	CAC	ГGC.	AGG.	ACC	ACC	ACT	CAA	TAA	CAA	TAA	TTG	ATA	GCT	ATC	GAT	TCT	TTA	TTG	AAA	AAA
721	CTT(CAG	GCA	TCA	ATC	TAC	AAA	TAA	ACA	ATT	CTG	ACT	СТА	AAA	AAA	AAA	A			

Figure S1. Nucleotide and deduced amino acid sequences of LycCTLR cDNA. In the nucleotide and deduced amino acid sequences of LycCTLR cDNA, a typical mRNA polyadenylation signal (AATAAA) are shown in italic in the 3'-UTR. The transmembrane domain is boxed and the intracellular domain is marked by broken line. The C-type lectin-like domain predicted by SMART is underlined. The conserved cysteine residues are highlighted in dark grey. Ca²⁺ binding site 1 are marked with solid diamond. Ca²⁺ binding site 2 are marked with hollow diamond.



Figure S2. Production and purification of recombinant LycCTLR protein. (**a**) Lane 1: protein molecular weight marker; Lane 2: induced pET-His in *E. coli* BL21; Lane 3: non-induced pET-LycCTLR in *E. coli* BL21; Lane 4: induced pET-LycCTLR in *E. coli* BL21; Lane 5: supernatant from induced pET-LycCTLR in *E. coli* BL21; Lane 6: inclusion body from induced pET-LycCTLR in *E. coli* BL21; (**b**) Lane 1: protein molecular weight marker; Lane 2: purified rLycCTLR protein.

Primer Name	Nucleotide Sequence (5'→3')	Purpose				
5' outer	ACACTCGTAACAGTTAGGCACT					
5' inner	AGAAACCATAAGGCTGATGAAA	RACE PCR				
3' outer	GAAATGGGTGGACAACTCAATA					
3' inner	GGGACATCAGAACCAGACAACA					
CTLR-gF	GAAGTCTCTGCAAACTCACAAA	Genome cloning				
CTLR-gR	GTCAGAATTGTTTATTTGTAGAT					
CTLR-DistF	CGGAATTCGAGGGACAGGAGTCCTATCTA	Pool time DCP				
CTLR-DistR	CCCGGACTTTCACTTGAGCAGTTTGGCGT	Real-time FCK				
CTLR-RF	CCGGAATTCATGATGAACCTCATCAAAAA	Recombinant expression				
CTLR-RR	CCCAAGCTTTCACTTGAGCAGTTTGGCGT					
Actin-F	GACCTGACAGACTACCTCATG	β -actin amplification				
Actin-R	AGTTGAAGGTGGTCTCGTGGA					

Table S1. Oligonucleotide primers used for cloning and expression analyses.