

1 Microscope Model LSM510 (Carl Zeiss, Oberkochen, Germany) belonging to Central  
2 Research Laboratory, Okayama University Medical School.

3

#### 4 **Biotin labeled CCN2 protein derived from HeLa cells**

5 CCN2 protein derived from HeLa cells was biotin-labeled by a commercially available  
6 kit, following the manufacturer's instructions (Biotin Labeling Kit-NH<sub>2</sub>; Dojindo  
7 Molecular Technologies, Inc, Kamimashiki-Gun, Japan).

8

#### 9 **RNA extraction and cDNA synthesis**

10 Cells were collected, and total RNA was extracted by following the manufacturer's  
11 instructions (RNeasy kit, Qiagen). Total RNA (500 ng) was reverse-transcribed by  
12 AMV Reverse Transcriptase (Takara, Ohtsu, Japan) at 42°C for 30 min, according to the  
13 manufacturer's protocol.

14

#### 15 **Real-time PCR**

16 Real-time PCR was performed by using TOYOBO SYBR Green PCR Master Mix  
17 (TOYOBO, Osaka, Japan) in a LightCycler™ system (Roche, Basel, Switzerland).  
18 Reactions were performed in a 10- $\mu$ l reaction mixture containing 1  $\mu$ l of cDNA, 0.4  $\mu$ l  
19 of each primer (5  $\mu$ M), and 5  $\mu$ l of 1 $\times$  SYBR Green master mix. Primer sets and  
20 optimized conditions for the PCR of each target are listed in Table 1. Absence of  
21 non-specific PCR products was checked by melting curve and electrophoresis analyses.  
22 Relative copy numbers were computed based on data obtained with a serial dilution of a  
23 representative sample for each target gene.

24

#### 25 **Antisense oligonucleotides**

26 To inhibit the expression of HIF1 $\alpha$ , we prepared an antisense phosphorothioate  
27 oligonucleotide (AS-HIF) and a sense oligonucleotide (S-HIF: control) according to the

1 nucleotide sequence of the human HIF1 $\alpha$  gene (Caniggia et al., 2000). The nucleotide  
2 sequences of the AS-HIF and S-HIF were 5'-GCCGGCGCCCTCCAT-3' and  
3 5'-ATGGAGGGCGCCGGC-3', respectively. These oligonucleotides were added  
4 directly to medium in HCS-2/8 cells culture at a concentration of 10  $\mu$ M.

5

### 6 **Animals and preparation of tissue**

7 After Balbc/j mice (2 weeks of age) had been anesthetized with sodium pentobarbital  
8 (Nembutal, Abbott laboratories, North Chicago, IL; 25 mg/kg), proximal tibiae were  
9 harvested and immersed in 4% paraformaldehyde (w/v) in phosphate buffer (PB: 0.1 M  
10 NaH<sub>2</sub>PO<sub>4</sub>, 0.1 M Na<sub>2</sub>HPO<sub>4</sub>; pH 7.4) at 4°C overnight. After having been rinsed in  
11 PBS, the tibiae were decalcified in 0.5 M EDTA, pH 7.4, at 4°C and then embedded in  
12 paraffin wax. The sections were prepared at a thickness of 7  $\mu$ m and mounted on  
13 silane-coated slides. The Animal Committee of Okayama University approved all of  
14 the procedures.

15

### 16 **Immunohistochemistry**

17 Tibial sections were dewaxed in xylene and rehydrated through a graded series of  
18 ethanol to water, blocked in a blocking buffer (5% dry non-fat milk in Tris-buffered  
19 saline), and incubated overnight at 4°C with the primary anti-LRPAP1 antibody (1:100)  
20 and subsequently with an HRP-conjugated anti-rabbit IgG (1:1000) for 1 h at room  
21 temperature. Color development was performed by using 3, 3'-diaminobenzidine  
22 tetrachloride (Dojindo, Tokyo, Japan). The sections were also counterstained with  
23 hematoxylin and mounted. Control samples were processed with the omission of the  
24 primary antibody.

25

### 26 **Statistics**

27 Data were presented as means  $\pm$  standard deviations, and the statistical significance

1 of differences in mean values was assessed by performing Student's unpaired *t*-test.  
2 Differences among the mean values were considered significant at a *P* value of <0.05.

3

4 All experiments were repeated at least twice, and similar results were obtained.

5

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13

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19  
20

1 **Figure Legends**

2 **Fig. 1. Effect of *LRP1* knockdown on CCN2 association with chondrocytic cells.**

3 Recombinant CCN2 was designed to possess a Flag tag, which was captured or  
4 internalized by LRP1 (left panel). Dual-tagged recombinant CCN2 was added to  
5 control or *LRP1* knockdown HCS-2/8 cells, the medium was removed after 1 h, and the  
6 cellular protein was collected. Immunoblotting was performed by using anti-Flag or  
7 His tag antibody. Positions of molecular weight markers (35, 75 kDa) are shown at the  
8 right of the images (right panel). NC, non-silencing siRNA as a negative control;  
9 si-1163, LRP1 siRNA (target sequence position 1163); si-13157, LRP1 siRNA (target  
10 sequence position 13157).

11

12 **Fig. 2. Effect of chlorpromazine on the CCN2 association with chondrocytic cells.**

13 Dual-tagged CCN2 was added to control or chlorpromazine (A) / M $\beta$ CD (B) -treated  
14 HCS-2/8 cells and the cellular protein was collected after 1 h. Immunoblotting was  
15 performed by using anti-Flag or anti-His tag antibody. As a result, the  
16 bound/incorporated CCN2 was decreased in the chlorpromazine-treated HCS-2/8 cells  
17 (A), while it was not in M $\beta$ CD-treated ones (B). Positions of molecular weight  
18 markers (35, 75 kDa) are shown at the right of the images. (C, D) Internalization of the  
19 exogenously added rhCCN2 into HCS-2/8 cells and intracellular co-localization with  
20 endogenous LRP1. The Flag-tagged CCN2 was added and analyzed by laser-scanning  
21 confocal microscopy after 15 minutes. The distribution of LRP1 in HCS-2/8 cells was  
22 visualized with an antibody for LRP1 (H-80 for  $\alpha$ -subunit). The intracellular CCN2  
23 uptake and co-localization with LRP1 was evident. (E-L) Intracellular delivery of  
24 exogenously added rhCCN2 into certain organelle of HCS-2/8 cells. Staining of  
25 clathrin (E, F), EEA1 (G, H; a marker of early endosomes) or Rab11 (I, J; a marker of  
26 recycling endosomes) is shown. The squares in panels C, E, G and I indicate the areas  
27 enlarged in the panels D, F, H and J, respectively. Incorporated rhCCN2 was partially



1 targeted to clathrin (C, D) and early endosomes (E, F). Interestingly, exogenously  
 2 added rhCCN2 and, particularly, the recycling endosome marker were predominantly  
 3 co-localized in HCS-2/8 cells (I, J). Merge: merged images of rhCCN2 with LRP1 (C,  
 4 D), clathrin (E, F), EEA1 (G, H), or Rab11 (I, J) staining. Scale bars, 5  $\mu$ m.

5

6 **Fig. 3. Effect of LRPAP1 on CCN2 transcytosis in chondrocytes.** (A) Schematic  
 7 representation of the sampling strategy is shown. *E. coli*-derived dual-tagged CCN2  
 8 (B, C) was added to control or LRPAP1-treated HCS-2/8 cells in the upper chamber of a  
 9 Transwell, the medium in the upper chamber was removed after 1 h; and the cellular  
 10 protein (B) and the medium in the lower chamber (C) were collected as illustrated.  
 11 Immunoblotting was performed by using anti-Flag or His tag antibody. As a result, the  
 12 bound/ incorporated (B) and transcytosed (C) amount of CCN2 was decreased in the  
 13 LRPAP1-treated HCS-2/8 cells. Comparable results were obtained with HeLa  
 14 cell-derived biotinylated recombinant CCN2 detected by the Streptavidin conjugate (D,  
 15 E). Positions of molecular weight markers (35, 75 kDa) are shown at the right of the  
 16 images. NC, the mixture of anti-FLAG® M2 affinity gel or Ni-NTA-agarose gel and  
 17 serum-free D-MEM without Flag or His-fusion protein as a negative control.

18

19 **Fig. 4. Effect of hypoxia on levels of LRP1 mRNA and protein in HCS-2/8 cells.**

20 (A) The *LRP1* mRNA level in HCS-2/8 cells under hypoxia. The level of mRNA was  
 21 standardized to that of *18s* mRNA. Exposure to hypoxia resulted in a time-dependent  
 22 increase in the *LRP1* mRNA level. The values represent the means  $\pm$  SD. \**P* < 0.05.  
 23 (B) LRP1, CCN2 and HIF1 $\alpha$  protein level in HCS-2/8 cells under hypoxia for 48 h.  
 24 Immunoblotting was carried out with anti-LRP1, CCN2 and HIF1 $\alpha$  antibody. Stronger  
 25 signals for the LRP1 subunit, CCN2 and HIF1 $\alpha$  were detected under hypoxia. (C)  
 26 Effect of antisense HIF1 $\alpha$  oligonucleotides on HIF1 $\alpha$ , LRP1 and CCN2 production  
 27 under hypoxic condition for 48 h. Treating the cells in 5% O<sub>2</sub> with antisense

1 oligonucleotides to HIF1 $\alpha$  (AS-HIF) for 48 h abolished LRP1, CCN2 and HIF1 $\alpha$   
2 production. S-HIF: control experiments with the sense oligonucleotide. The position  
3 of the molecular weight marker used (35 and 75 kDa) is shown at the left of the images.

4

5 **Fig. 5. Effect of hypoxia on CCN2 transcytosis in chondrocytes.** Transcytosis assay  
6 was performed as described in the legend of Figure 2. As a result, the bound/  
7 incorporated (A) and transcytosed (B) amount of CCN2 increased in the HCS-2/8 cells  
8 under hypoxia, and the increase was suppressed by LRPAP1. NC, the mixture  
9 anti-FLAG® M2 affinity gel or Ni-NTA-agarose gel and serum free D-MEM without  
10 Flag or His-fusion protein as a negative control.

11

12 **Fig. 6. LRPAP1 in HeLa, MDA-231, and HCS-2/8 cells.** (A) The mRNA level of  
13 *LRP1* and *LRPAP1* in HeLa, MDA-231, and HCS-2/8 cells. The level of each mRNA  
14 was standardized to that of *GAPDH* mRNA. *LRP1* and *LRPAP1* mRNAs were highly  
15 expressed in chondrocytic HCS-2/8 cells. The values represent the means  $\pm$  SD. \* $P$  <  
16 0.05. (B) The protein level of LRPAP1 in HeLa, MDA-231, and HCS-2/8 cells.  
17 Immunoblotting was carried out with anti-LRPAP1 antibody. The strongest signal for  
18 the LRPAP1 was detected in the HCS-2/8 cells. Positions of molecular weight  
19 markers (35kDa) are shown at the right of the images.

20

21 **Fig. 7. Difference in expression and distribution of LRPAP1 among chondrocytes**  
22 **of various differentiation stages.** (A) Immunohistochemical analysis of LRPAP1 in  
23 the growth plate. Tibial sections from mice were stained with anti-LRPAP1 antibody.  
24 The ECM in the entire growth-plate cartilage was immunopositive for LRPAP1, with  
25 the strongest signal in the resting zone. The dark gray circles in the bottom panel  
26 represent the cells that express *Ccn2* gene. The hatched circles therein represent the  
27 cells that accumulated CCN2 protein. Scale bars, 2 mm. (B) Change in the

1 expression levels of *lrpap1* mRNA and other mRNAs in chicken sternum chondrocytes  
2 of various differentiation stages. LC, USP, and USC represent resting, proliferating,  
3 and hypertrophic chondrocytes in the growth plate, respectively. The expression of  
4 *lrpap1* mRNA was the highest in the LC cells. Thus these results support the data  
5 obtained in vivo (A). The values represent the means  $\pm$  SD. \* $P < 0.05$ .

6

7 **Fig. 8. Schematic representation of the molecular mechanism determining the**  
8 **polarity of CCN2 distribution.** CCN2 protein is transported from prehypertrophic  
9 chondrocytes, where it is produced, through transcytosis mediated by LRP1 to the  
10 hypertrophic chondrocytes. The high levels of LRPAP1 in the resting zone interfere  
11 with this transportation, whereas this interference is presumably attenuated by  
12 down-regulation of LRPAP1 in the hypertrophic layer and down-regulation of LRP1 by  
13 oxygen in the late hypertrophic layer. As a result of the down-regulation of LRP1 by  
14 oxygen in the late hypertrophic layer, CCN2 is accumulated in the hypertrophic layer.

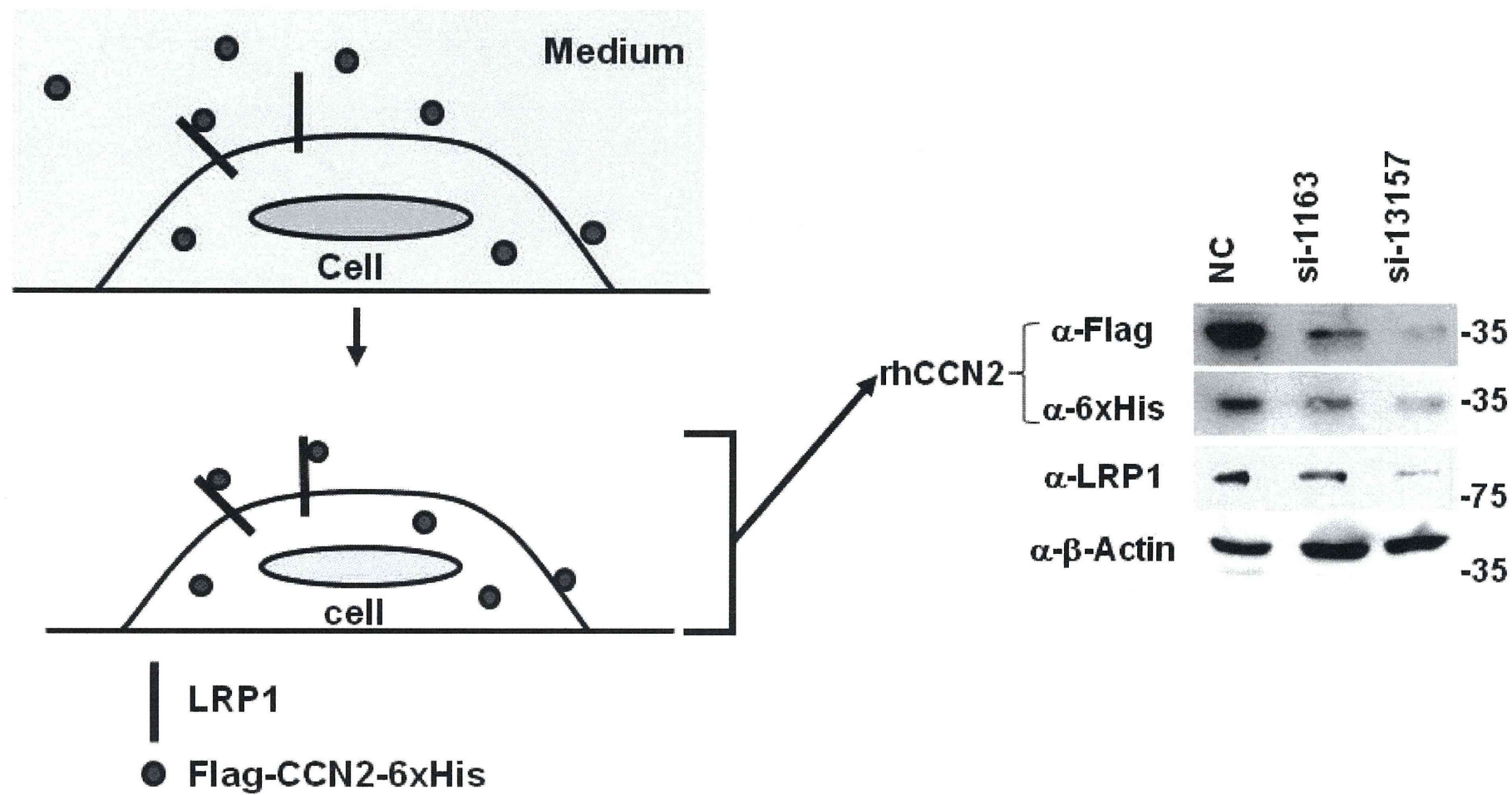


Fig. 1

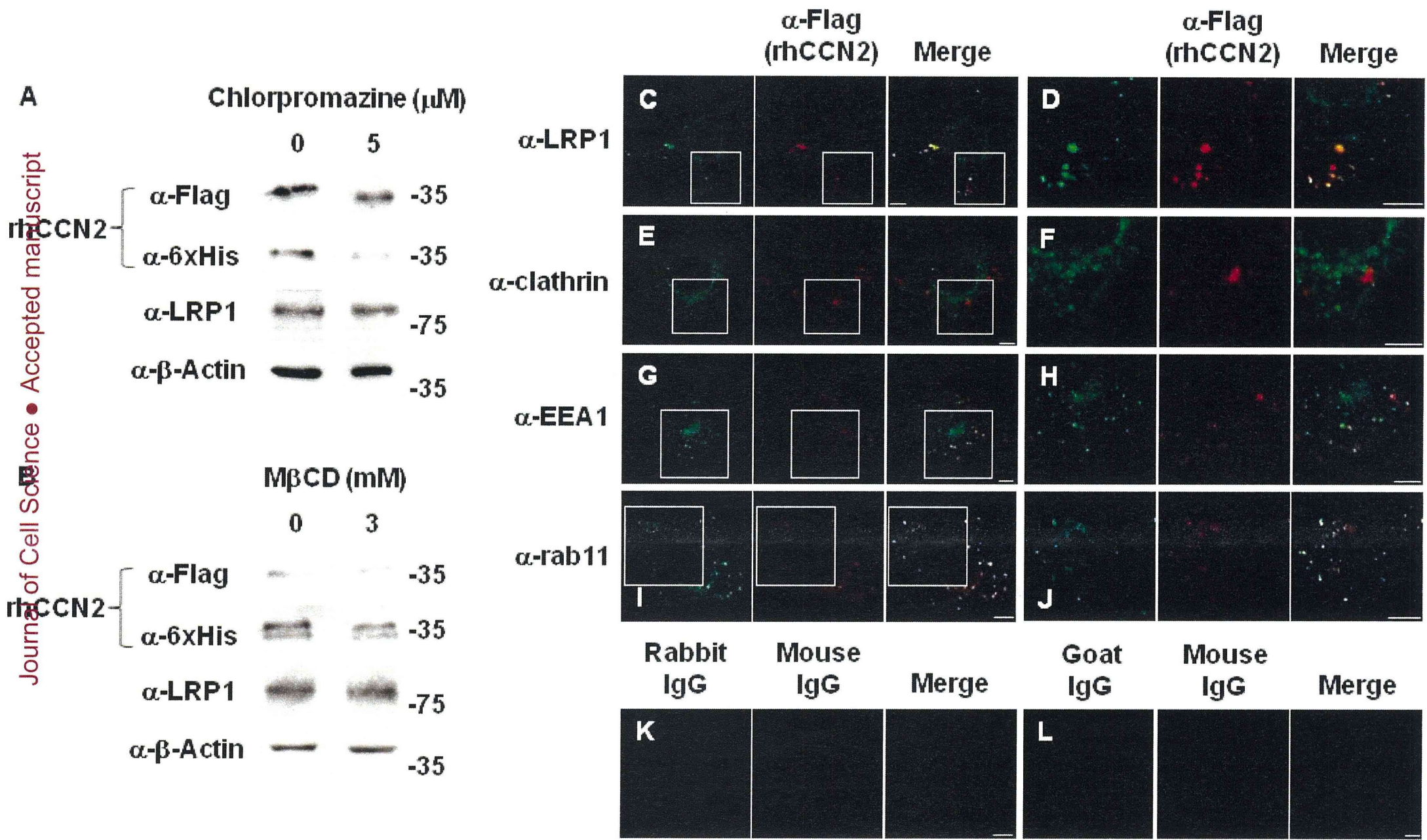


Fig. 2

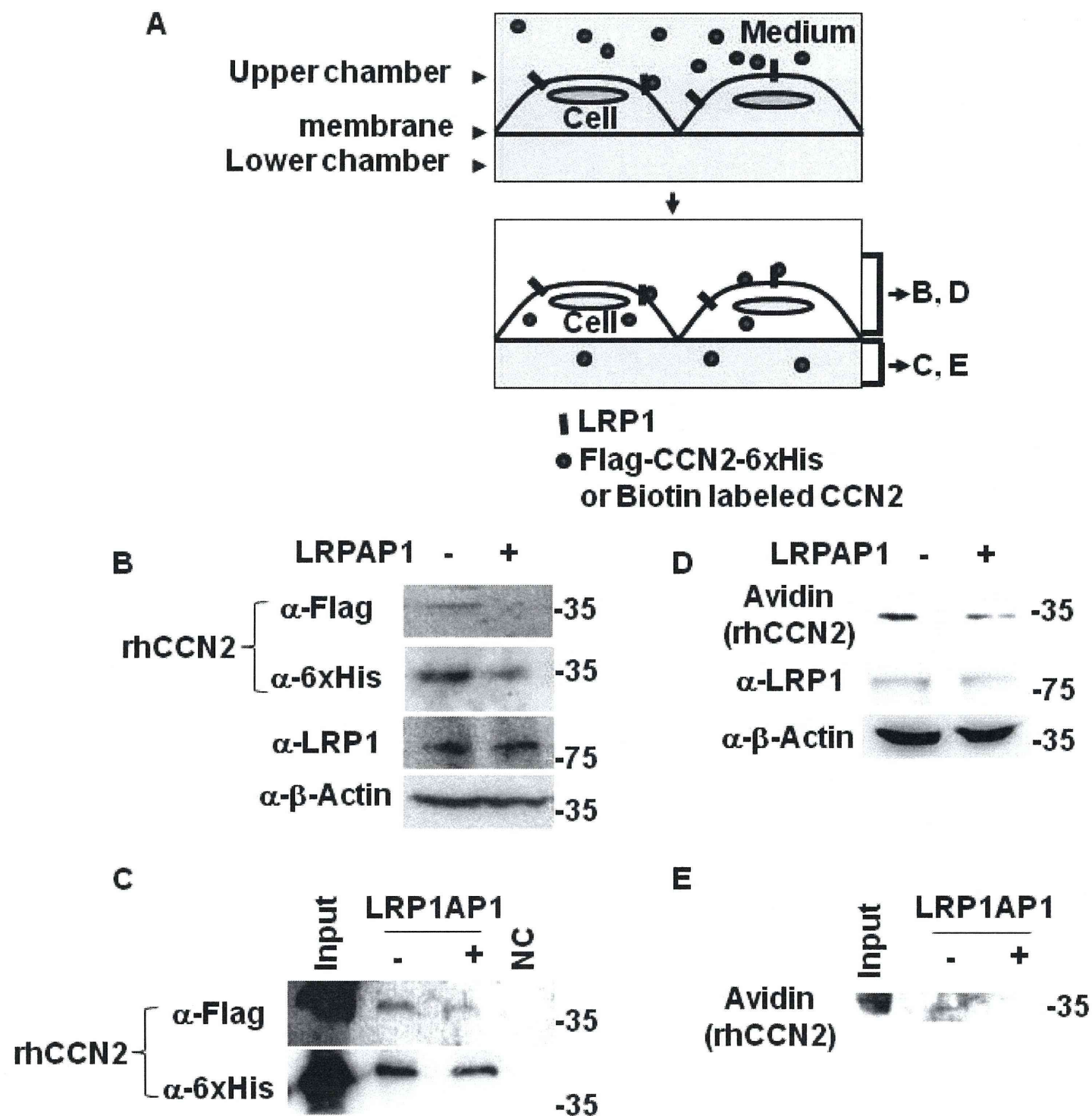


Fig. 3

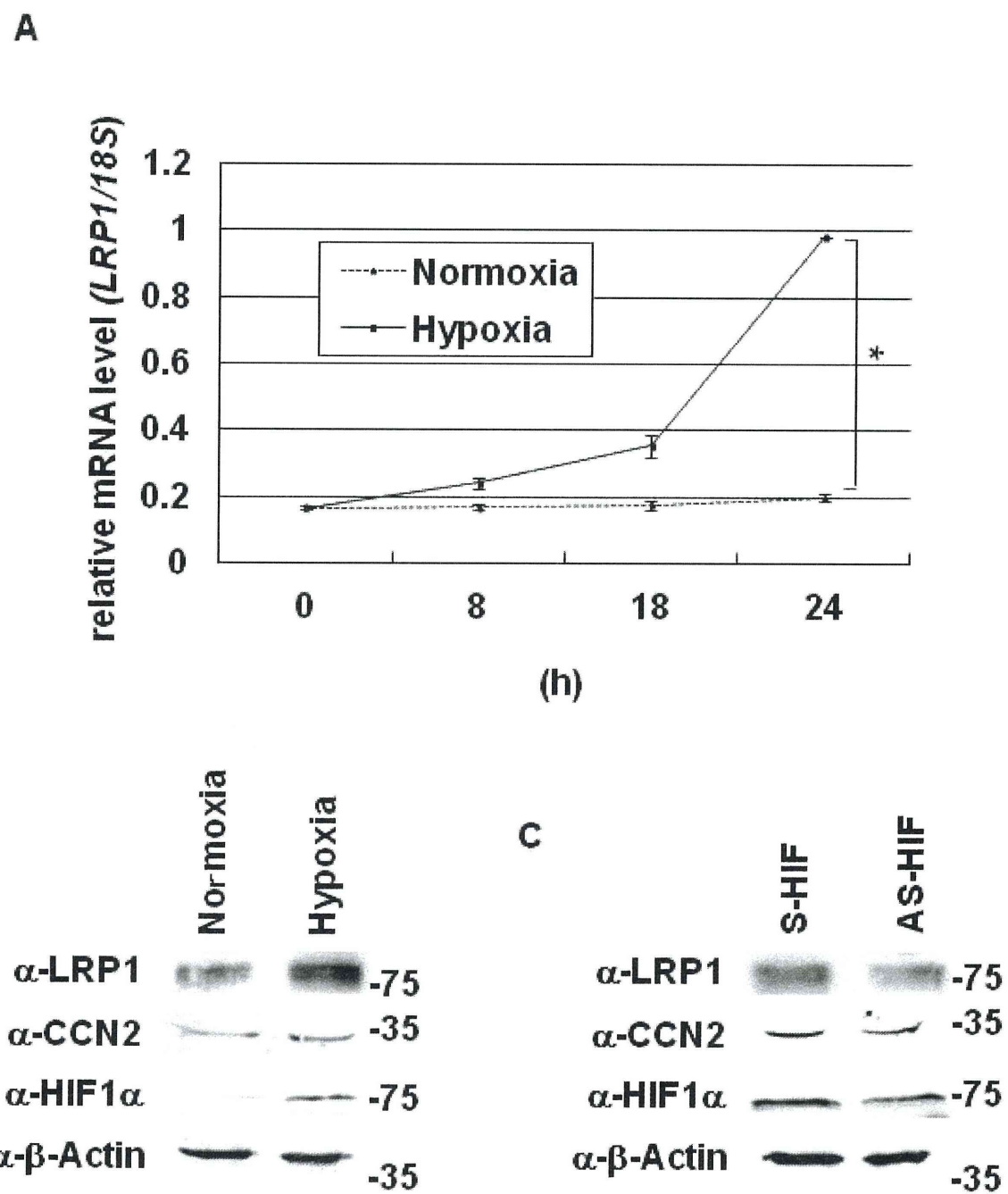


Fig. 4

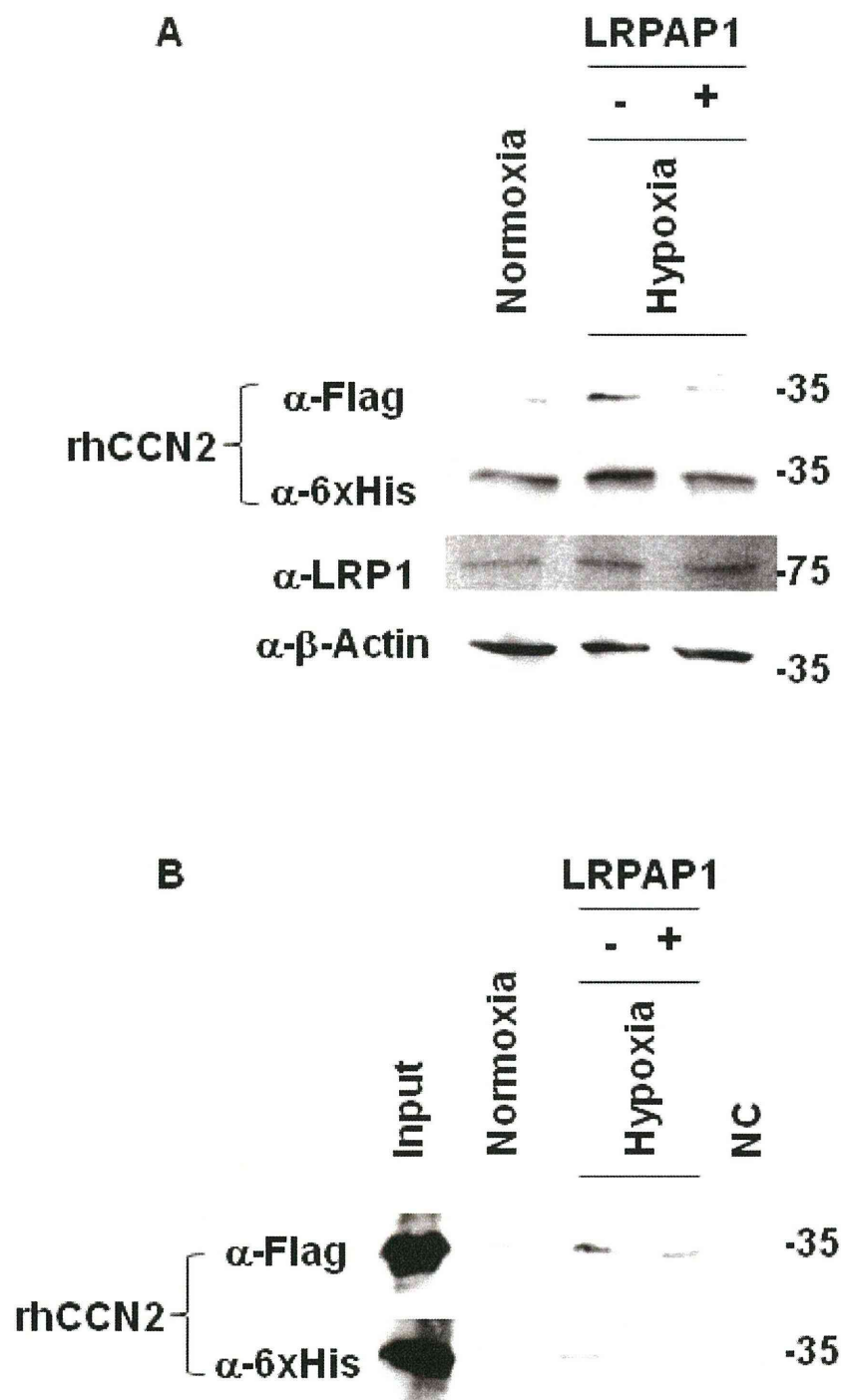
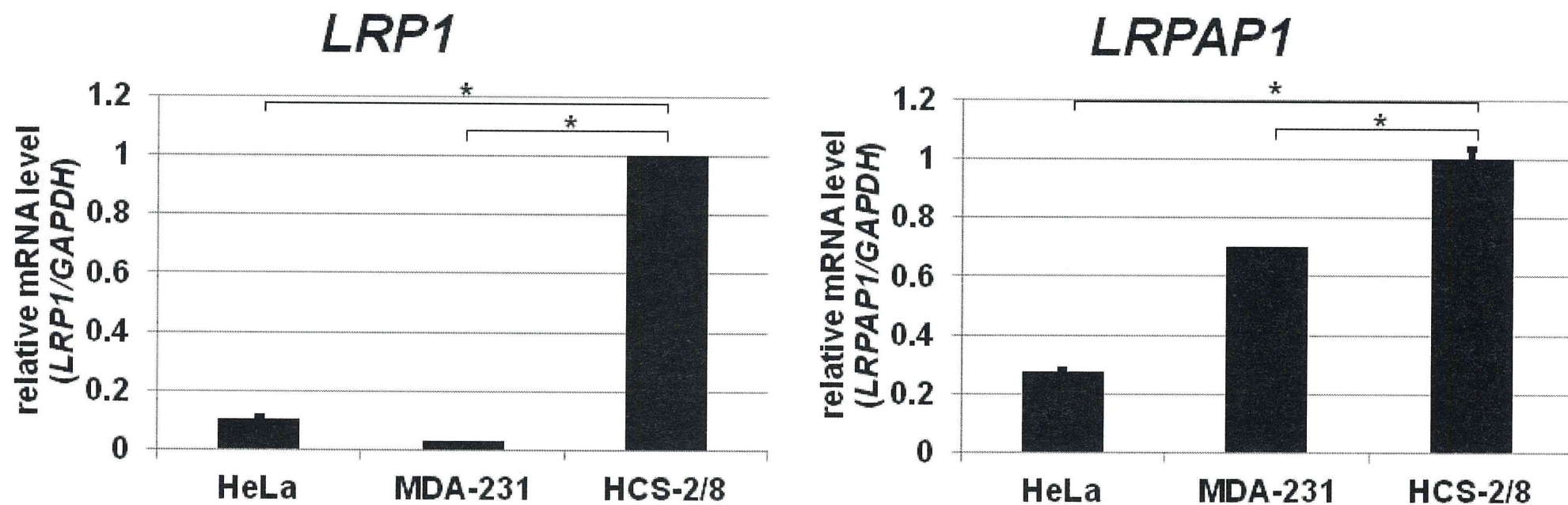


Fig. 5



A



B

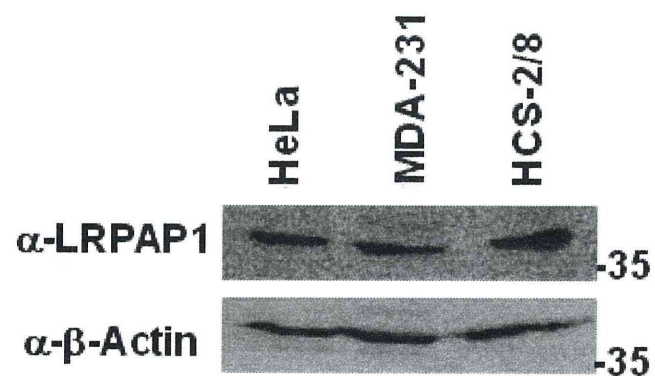
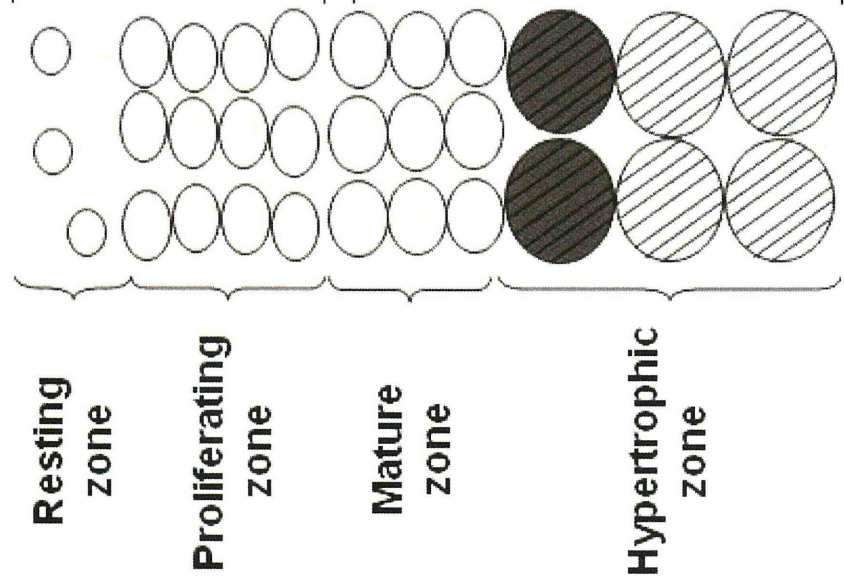
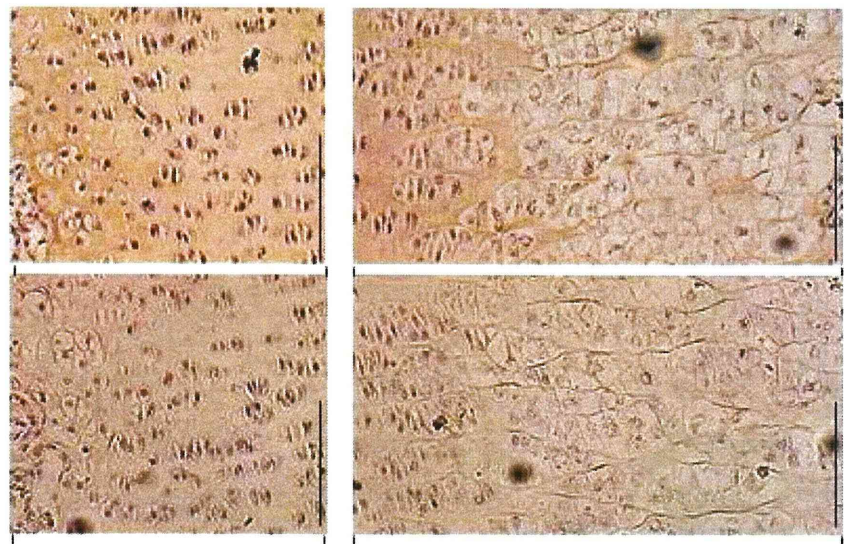


Fig. 6

**A**

$\alpha$ -IRPAP1  
NC



**B**

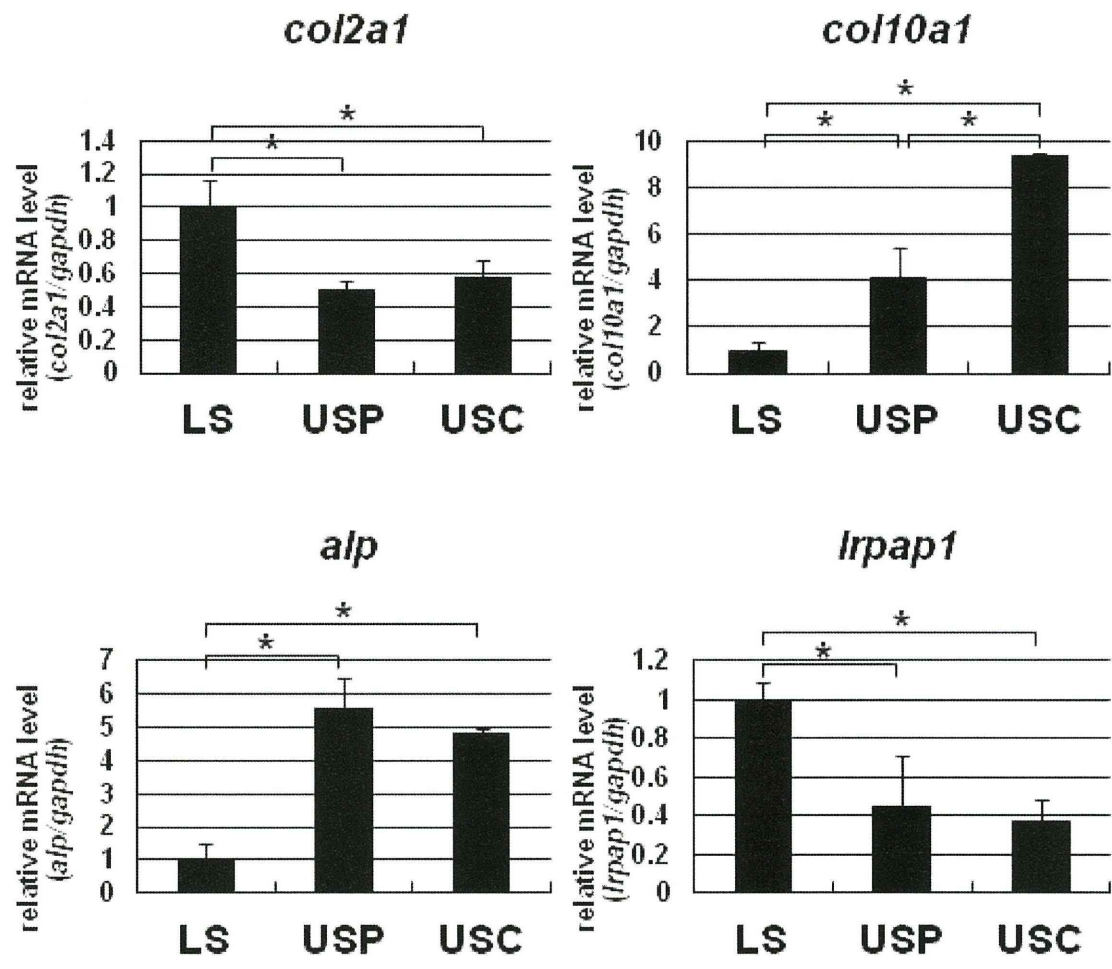


Fig. 7

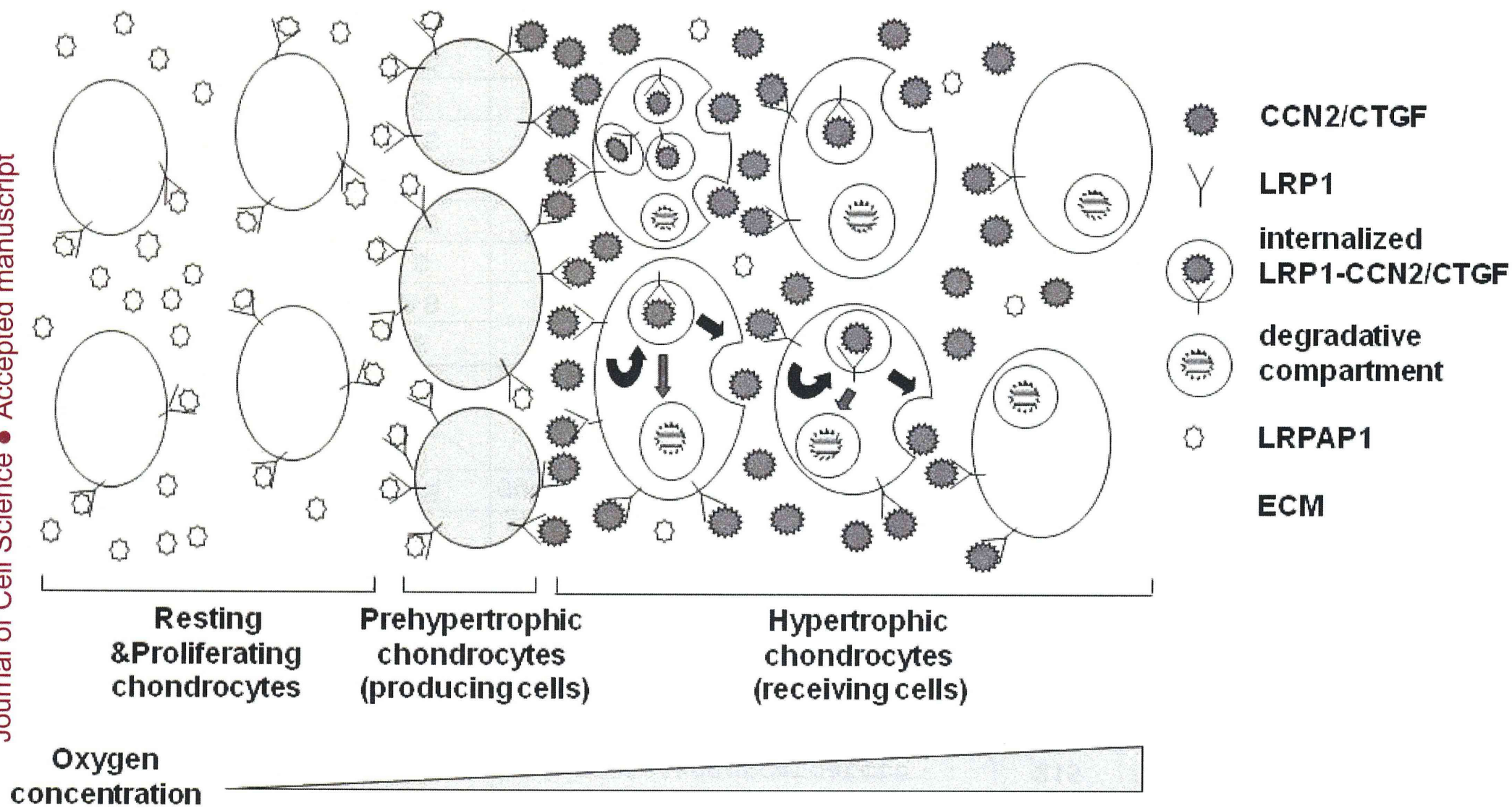


Fig. 8

Table 1.

## Primers and experimental conditions for real-time PCR

<i>target gene</i> (human)	Primer direction	Sequence(5'→3')	Length of PCR product	Annealing temperature (°C)
<b><i>GAPDH</i></b>	<b>S</b>	gccaaaagggtcatcatctc	<b>215</b>	<b>65</b>
	<b>AS</b>	gtcttctgggtggcagtgat		
<b><i>LRP1</i></b>	<b>S</b>	acatatagcctccatcctaadc	<b>152</b>	<b>65</b>
	<b>AS</b>	ttccaatctccacgttcat		
<b><i>LRPAP1</i></b>	<b>S</b>	ctgaggctgagttcgaggag	<b>150</b>	<b>65</b>
	<b>AS</b>	gctgcttctggtagtggttg		
<b><i>18S</i></b>	<b>S</b>	gcgaattcctgccagtagcatatgcttg	<b>140</b>	<b>60</b>
	<b>AS</b>	ggaagcttagaggagcgagcgaccaaagg		
<b>Target gene (chicken)</b>				
<b><i>gapdh</i></b>	<b>S</b>	aggctgtggggaaagtca	<b>202</b>	<b>65</b>
	<b>AS</b>	gacaacctggtcctctgtgtat		
<b><i>col2a1</i></b>	<b>S</b>	agaaaggaa tccagcccaat	<b>236</b>	<b>65</b>
	<b>AS</b>	acacctgccagattgattcc		
<b><i>col10a1</i></b>	<b>S</b>	acatgcatttacaatatcgttac	<b>160</b>	<b>60</b>
	<b>AS</b>	aaaatagtagacgttaccttgactc		
<b><i>alp</i></b>	<b>S</b>	aacggccctggctataagat	<b>186</b>	<b>60</b>
	<b>AS</b>	tgggggatgtagttctgctc		
<b><i>lrpap1</i></b>	<b>S</b>	acccggtgaaagaggaagtc	<b>164</b>	<b>65</b>
	<b>AS</b>	tgccatgtcccacaaatc		

S, sense; AS, anti-sense.