Identification of Calcitonin Expression in the Chicken Ovary: Influence of Follicular Maturation and Ovarian Steroids¹

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ABSTRACT

Calcitonin (CALCA), a hormone primarily known for its role in calcium homeostasis, has recently been linked to reproduction, specifically as a marker for embryo implantation in the uterus. Although CALCA expression has been documented in several tissues, there has been no report of production of CALCA in the ovary of any vertebrate species. We hypothesized that the Calca gene is expressed in the chicken ovary, and its expression will be altered by follicular maturation or gonadal steroid administration. Using RT-PCR, we detected Calca mRNA and the calcitonin receptor (Calcr) mRNA in the granulosa and theca layers of preovulatory and prehierarchial follicles. Both CALCA and Calca mRNA were localized in granulosa and thecal cells by confocal microscopy. Using quantitative PCR analysis, F1 follicle granulosa layer was found to contain significantly greater Calca mRNA and Calcr mRNA levels compared with those of any other preovulatory or prehierarchial follicle. The granulosa layer contained relatively greater Calca and Calcr mRNA levels compared with the thecal layer in both prehierarchial and preovulatory follicles. Progesterone (P4) treatment of sexually immature chickens resulted in a significantly greater abundance of ovarian Calca mRNA, whereas estradiol (E2) or P4 + E2 treatment significantly reduced ovarian Calca mRNA quantity. Treatment of prehierarchial follicular granulosa cells in vitro with CALCA significantly decreased FSH-stimulated cellular viability. Collectively, our results indicate that follicular maturation and gonadal steroids influence Calca and Calcr gene expression in the chicken ovary. We conclude that ovarian CALCA is possibly involved in regulating follicular maturation in the chicken ovary.

estradiol, follicular development, granulosa cells, progesterone, theca cells

INTRODUCTION

Calcitonin (CALCA) is a peptide hormone that is primarily associated with maintaining calcium homeostasis. Synthesized as procalcitonin in the thyroid gland of mammals or the ultimobranchial gland of avian species, CALCA is secreted in response to hypercalciemia [1]. To maintain constant levels of circulating calcium ions, CALCA inhibits osteoclastic resorption of calcium from bone but increases excretion of

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calcium ion in the kidneys [2, 3]. Recent studies have identified numerous nonthyroidal sources of CALCA, which include the thymus, jejunum, lung, urinary bladder [4], prostate [5], endometrium [6], pituitary gland [7], and mammary glands [8, 9]. Although CALCA is not well characterized in most tissues, the physiologic role of CALCA has been well documented in the pituitary gland and endometrium, wherein CALCA has been shown to inhibit prolactin secretion and favor embryo implantation, respectively [6, 10].

Numerous studies have suggested a role for CALCA in female reproduction. Circulating CALCA levels have previously been shown to rise significantly just before ovulation in the rainbow trout, eel, and rat [11-13]. More recent studies, however, have focused on CALCA's involvement in blastocyst implantation. In cycling rats, both Calca mRNA and CALCA are expressed in the glandular tubular epithelial cells of the endometrium, with significant amounts of CALCA secreted into the uterine lumen on Days 4 and 5 of gestation immediately preceding implantation [14]. An increase in endometrial Calca gene expression has been documented during the progesterone-dominant phase of the estrus or menstrual cycle, in which the uterus is most receptive to embryonic implantation in rats, baboon, and humans [6, 15, 16]. A transient expression of Calca in the preimplantation phase of the uterus appears to be critical for blastocyst implantation, since attenuation of Calca expression by administering antisense oligonucleotides is accompanied by a severe impairment in implantation of embryos. [17]. In addition, both estrogen (E_2) and progesterone (P_4) appear to play a major role in the expression of endometrial Calca. Administration of E2 or P4 was found to decrease and increase endometrial *Calca* gene expression, respectively [6,

Despite numerous studies describing Calca expression in several tissues, there is no clear evidence to suggest that *Calca* is expressed within the ovary itself. While exploring Calca expression in various chicken tissues, we serendipitously detected Calca cDNA expression in the chicken ovary. Based on this observation, we desired to further explore and characterize the expression of CALCA in the chicken ovary and determine how follicular maturation and gonadal steroids may alter ovarian Calca expression. We present novel evidence that both the Calca mRNA and CALCA peptide are expressed in the granulosa and theca cells of the ovarian follicles. We also provide novel evidence on the identification of the CALCA receptor (Calcr) mRNA within both the granulosa and theca cell layers. Furthermore, we show that follicular maturation significantly affects both Calca and Calcr mRNA expression within the granulosa layer, whereas CALCA influences the viability of prehierarchial follicle granulosa cells cultured in the presence of FSH.

MATERIALS AND METHODS

Reagents

Trizol and RNeasy kits used to isolate RNA were obtained from Invitrogen Corp. (Carlsbad, CA) and Qiagen, Inc. (Valencia, CA), respectively. M-MuLV reverse transcriptase and Taq polymerase used for RT-PCR were purchased from New England Biolabs (Beverly, MA). Additional PCR and RT materials (Platinum SYBR Green qPCR Super Mix-UDG, RNAseOut) were obtained from Invitrogen. Digoxigenin-labeled nucleotides, dinucleotide triphosphate mix, Sp6 polymerase, T7 polymerase, and antidigoxigenin antibody were purchased from Roche Applied Sciences (Indianapolis, IN). Estradiol 17-β and progesterone were purchased from Sigma-Aldrich (St. Louis, MO). Anti-eel CALCA antibody was purchased from Peninsula Laboratories (Belmont, CA). Biotinylated anti-rabbit immunoglobulin G (IgG) and streptavidin-Alexa 488 were purchased from Vector Laboratories (Burlingame, CA), and TOPRO-3 was purchased from Invitrogen, Chicken CALCA was purchased from Bachem (Torrance, CA), and recombinant human FSH was purchased from the National Hormone and Peptide Program (Torrance, CA). The CellTiter-Blue Cell Viability Assay was purchased from Promega (Madison, WI).

Animals

A commercial strain of Leghorn chickens (Hyline W36 strain) was maintained at the Poultry Research and Extension Center at Pennsylvania State University (University Park, PA). The chickens were provided with 16L:8D photoperiod and were provided with water and feed ad libitum. All animal procedures were carried out in accordance with the Institutional Animal Care and Use Committee-approved protocol.

Collection of Ovary/Separation of Granulosa or Theca Layer

Approximately 4–6 h prior to ovulation, chickens (n = 6; Hyline W36 strain; 60 wk old) were killed by decapitation with a guillotine. Prior to killing, the presence of a hard-shelled egg in the shell gland was confirmed by cloacal examination. Immediately following killing, the ovarian follicles were separated from the ovary and categorized based on their diameter into preovulatory follicles (F1–F6) and prehierarchial follicles (6–8 mm and 3–5 mm). The separated follicles were placed into ice-cold 0.75% saline. The granulosa and theca cell layers from each preovulatory follicle were separated using a dissection microscope following the method described previously [18], snap frozen in liquid nitrogen, and stored at -80° C until processed. Granulosa and thecal cell layers obtained from two to three prehierarchial follicles within each category (3–5 mm and 6–8 mm) were pooled from each animal.

Calca and Calcr RT-PCR

Granulosa and thecal cell layers from preovulatory and prehierarchial follicles, as well as other tissues (adipose, liver, hypothalamus, skeletal muscle, kidney, spleen, total ovary, chondrocyte, pituitary, and blood), were collected from sexually mature female chickens. Total RNA was extracted from these tissues using Trizol (Invitrogen) and/or the RNeasy kit (Qiagen, Valencia, CA). Following on-column DNAse I (Qiagen) treatment, first-strand cDNA was synthesized by reverse transcribing 1 μg total RNA using d(T) $_{30}$ A/G/C A/G/C/T,

2U M-MLV reverse transcriptase (New England BioLabs) in a 20- to 50-μl reaction. Approximately 100 ng single-stranded cDNA was used as template to amplify a 204-bp product of Calca cDNA using CT1Fwd and CT1Rev primers (Table 1) or a 495-bp product of Calcr cDNA using CTR1Fwd and CTR1Rev primers (Table 1). A touch-down PCR was performed using the following thermocycle: 94°C for 1 min, 30 cycles of 94°C for 5 sec and 72°C-68°C for 3 min. Annealing and primer extension were done at 72°C, 70°C, and 68°C during 1-5, 6-10, and 11-30 cycles, respectively. The PCR products were subjected to agarose gel electrophoresis and ethidium bromide staining for visualization. For negative controls, reverse transcription reactions using 1 µg total RNA from each tissue/follicle with no reverse transcriptase (-RT) were used as a template in place of reverse transcription reactions that contained reverse transcriptase (+RT). In addition, total RNA extracted from blood cells was subjected to RT-PCR to rule out blood contamination as a source of Calca mRNA. Using another set of primers (CT2Fwd and CT2Rev; Table 1), a 417bp, full-length Calca cDNA was amplified with the F1 follicle granulosa cell cDNA as a template. The resultant products were sequenced (Davis Sequencing, San Diego, CA) to confirm the authenticity of Calca and Calcr cDNAs.

Cellular Localization of Calca mRNA by In Situ Hybridization Histochemistry in Ovary

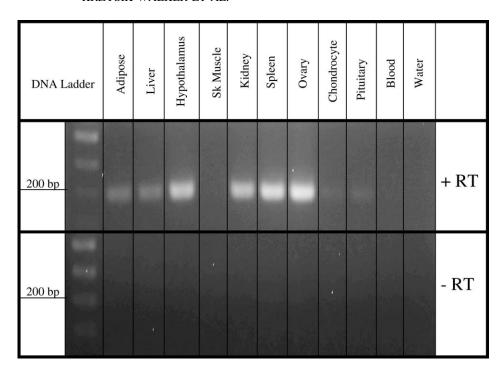
Preparation of digoxigenin-labeled Calca cRNA probe. A partial 345-bp-long chicken Calca cDNA was amplified from a chicken pituitary cDNA library and was subcloned into pGEM-T Easy vector (Promega, Madison, WI). The resultant plasmid was linearized using PstI or NcoI restriction endonucleases (New England BioLabs) and gel purified. Antisense and sense strands of Calca cDNA were transcribed using digoxigenin-labeled nucleotide (Roche Biochemicals) with T7 polymerase and Sp6 polymerase, respectively.

In situ hybridization histochemistry. A small piece of chicken ovary was fixed by immersion in 4% paraformaldehyde for 1 h at room temperature. Following fixation, the tissue was washed three times in PBS for 15 min each, dehydrated, cleared with xylene, and infiltrated with paraffin. Ovary tissue sections (10 µm) were cut using a rotary microtome (Microm, Walldorf, Germany), and serial sections were mounted on Superfrost Plus glass slides (VWR, West Chester, PA). Tissue sections were deparaffinized, hydrated, and treated with 0.2 N hydrochloric acid. Slides were rinsed in 0.3% Triton X-100 and treated with proteinase K (10 µg/ml; Roche Biochemicals) in 0.1 M Tris HCl containing 50 mM EDTA for 20 min. Tissue sections were then treated in 0.2% glycine for 5 min, fixed in 3.7% paraformaldehyde, and acetylated with 0.25% acetic anhydride and 0.1 M triethanolamine. The slides were then incubated in prehybridization buffer (50% formamide in 2× SSC) for 90 min at 37°C. Digoxigenin-labeled Calca cRNA probe (antisense, or sense strand for a negative control) were diluted in hybridization buffer (10 mM Tris-HCL, 12.5% Denhardt solution, 50% formamide, 0.5% SDS, 2× SSC, 0.73 mg/ml yeast tRNA, and 0.36 mg/ml salmon sperm) and applied to the slides under a coverslip. The slides were incubated in a humidified chamber at 45°C overnight. Afterwards, the slides were treated with RNAse (20 µg/ml; Gibco) in 10 mM Tris-HCl, 5 mM EDTA, and 0.3 M NaCl for 30 min at 37°C. Slides were then washed twice in 2× SSC for 15 min each time at room temperature, twice in 0.1× SSC for 5 min each, and twice in 0.1× SSC for 30 min each at 42°C. Next, the slides were washed in Tris-buffered saline (TBS) containing 0.1% Triton X-100 (Sigma) for 10 min, and treated with 1× blocking buffer (Roche Biochemicals) in TBS for 30 min. A monoclonal anti-digoxigenin

TABLE 1. The nucleotide sequences of the primers used to amplify Calca, Actb, and Calcr.

Name	Sequence	GenBank accession no.	Product length (bp)
Calca			_
CT1Fwd	5'-AGTCAGACCTGGCTTGGAGTCCATCA-3'	X03012	204
CT1Rev	5'-TGAGACAGTTTGCCCAGCACAAGT-3'		
CT2Fwd	5'-ATGGTCATGCTGAAGATTTCATCTTTCCT-3'	XM_420997	417
CT2Rev	5'-CTAGTTGTTTCCTAGGGTTTCCCCATAGTT-3'		
Actb			
BAFwd	5'-CTGGCACCTAGCACAATGAA-3'	L08165	123
BARev	5'-CTGCTTGCTGATCCACATCT-3'		
Calcr			
CTR1Fwd	5'-CATGCAGTTGCAAGAGCCAAATACTTCAAT-3'	XM_425985	495
CTR1Rev	5'-GTACAGGTAGACAGGGACCTCTGTGATGGA-3'		
CTR2Fwd	5'-TGGCAACTATATTCTGCTTCTTCA-3'	XM_425985	136
CTR2Rev	5'-GACGTTGCTGTGTAGGAGGTAGAT-3'		

FIG. 1. RT-PCR analysis of *Calca* gene expression in various tissues of the chicken. Approximately 100 ng cDNA (+RT) was used as template to amplify a 204-bp chicken *Calca* cDNA. Contamination controls consisted of RNA from each tissue without reverse transcriptase (-) or substitution of water for the cDNA template. SkM, skeletal muscle.



antibody (1:100; Roche Biochemicals) was applied to all slides and incubated at 4°C overnight. Following three washes in TBS for 5 min each, slides were treated with biotinylated anti-mouse IgG made in goat (1:400; Vector Laboratories) for 1 h at room temperature. Slides were washed three times in TBS and incubated with streptavidin-Alexa 488 (1:100; Invitrogen) for 1 h at room temperature. Following three more washes in TBS, slides were mounted with ProLong Gold Antifade Reagent and TOPRO-3 (1:500; Invitrogen). Green (Calca) fluorescent cells and their nuclei (red) were visualized by exciting the flourophores with respective lasers using an Olympus Fluoview 300 Confocal Laser Scanning Microscope (Olympus, Center Valley, PA).

Immunohistochemical Detection of Ovarian CALCA

Paraffin-embedded chicken ovary tissue sections were prepared as described above. Tissue sections were deparaffinized in Histoclear (National Diagnostics, Atlanta, GA), hydrated in decreasing concentrations of ethanol in water, and rinsed in TBS. Following a rinse in TBS containing 0.1% Triton X-100 (TBSX), slides were treated with 1% goat serum in TBSX for 30 min. Slides were then treated with affinity-purified rabbit anti-eel CALCA (Peninsula Laboratories, Belmont, CA), at a concentration of 40 µg/ml overnight at 4°C. Due to the unavailability of an anti-chicken CALCA, we selected anti-eel CALCA based on the close sequence homology (94%) as well as previous studies that used a similar antibody to identify immunoreactive cells in chicken tissue [19]. After rinsing with TBS, biotinylated goat anti-rabbit IgG (1:400; Vector Laboratories) was applied to the slides and allowed to incubate for 1 h, followed by rinsing in TBS. Sections were then incubated with streptavidin-Alexa 488 (Invitrogen) at a concentration of 1:100 for 1 h. Following three washes in TBS, sections were mounted with ProLong Gold antifade and TOPRO-3 (1:500; Invitrogen). As a negative control, nonimmune rabbit IgG (Vector Laboratories) was substituted in place of the primary antibody. Green (CALCA) fluorescent cells and their nuclei (red) were visualized by exciting the flourophores with respective lasers using an Olympus Fluoview 300 Confocal Laser Scanning Microscope.

Effect of Follicular Maturation on Calca and Calcr mRNA Quantity

Total RNA extracted from the preovulatory follicles (F1, F3, and F6) and prehierarchial follicles (3–5 mm) was subjected to quantitative PCR for determination of *Calca* and *Calcr* mRNA abundances. Total RNA (1 μg) was reverse transcribed using d(T)₃₀A/G/C A/G/C/T, 2U M-MLV reverse transcriptase in a 20-μl reaction. Chicken *Calca* mRNA, *Calcr* mRNA, and *Actb* mRNA were quantified utilizing 5–12 μl of the reverse transcription reaction (equivalent to 100–240 ng single-stranded cDNA) as template in the real-time quantitative PCR analysis. A 204-bp product for chicken *Calca* cDNA was amplified using the CT1Fwd and CT1Rev primers (Table 1),

whereas a 136-bp product of chicken Calcr was amplified using CTR2Fwd and CTR2Rev primers (Table 1). Similarly, a 123-bp product of chicken Actb was amplified using BAFwd and BARev primers (Table 1). The real-time quantitative PCR consisted of 1× Platinum SYBR Green qPCR Super Mix-UDG and 300 nM of forward and reverse primers. The reactions were carried out in the DNA Engine Opticon II (MJ Research) with the following thermocycle parameters: 50°C for 2 min and 95°C for 2 min, followed by 35 cycles of 95°C for 15 sec, 55°C for 30 sec, and 72°C for 30 sec. At the end of amplification, a melting curve analysis was done by heating the PCR products from 65°C-95°C, held for 15 sec at increments of 0.2°C, and the fluorescence was detected to confirm the presence of a single amplification product. Tissue samples from each animal were run in duplicate to obtain average C_T values for Calca mRNA, Calcr mRNA, and Actb mRNA. The log-linear threshold values (CT) during the exponential phase of the PCR for Calca mRNA or Calcr mRNA were subtracted from those of Actb mRNA. Calca mRNA quantity and Calcr mRNA quantity were expressed as a proportion of Actb mRNA quantity following $2^{-\Delta\Delta C}_{T}$ method for converting log-linear C_{T} values to linear term [20]. The relative amounts of Calca mRNA or Calcr mRNA in the follicles were then compared.

Effect of Ovarian Steroids on Ovarian Calca mRNA Quantity

Sexually immature female chickens (16 wk old; n = 7) were injected intramuscularly with peanut oil containing estradiol-17 β (E₂; 0.5 mg/kg body weight; four injections on alternate days [21]), progesterone (P₄; 0.17 mg/kg body weight per day for 7 consecutive days [22]), estradiol and progesterone together (E₂ + P₄) at the above dosages, or no steroids (negative control). Seven days after the first dose, the chickens were killed by decapitation, and the oviduct was isolated (infundibulum to shell gland) and weighed to confirm the efficacy of estradiol and/or progesterone treatments. Ovaries were isolated and snap frozen in liquid nitrogen. Total RNA from the ovaries of each chicken was extracted and subjected to quantitative PCR analysis for the determination of *Calca* mRNA and *Actb* mRNA, as described above. The amount of *Calca* mRNA is expressed as a proportion to *Actb* mRNA and compared among the treatment groups.

Effect of Chicken CALCA on Granulosa Cell Viability

Granulosa cell isolation from prehierarchial follicles. Approximately 2 h prior to ovulation, chickens (Hyline W36 strain; 70 wk old) were killed by decapitation with a guillotine. Prior to killing, the presence of a hard-shelled egg in the shell gland was confirmed by cloacal examination. The granulosa layers of 10–12 prehierarchial follicles (6–8 mm in size) were removed from two hens each, pooled, and dispersed in 0.3% collagenase as previously

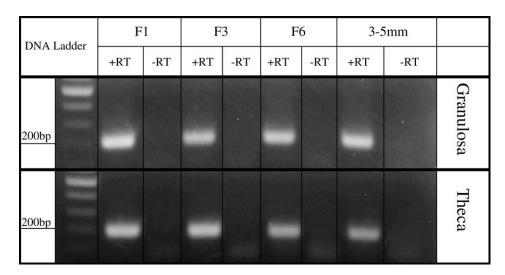


FIG. 2. RT-PCR analysis of *Calca* gene expression in granulosa and thecal cells layers of preovulatory (F1, F3, and F6) and prehierarchial (3–5 mm) follicles. Approximately 100 ng cDNA (+RT) was used as a template to amplify a 204-bp product of chicken *Calca*. Contamination controls consisted of RNA from each tissue without reverse transcriptase (-).

described [23]. Cell viability and number were ascertained by trypan blue exclusion method.

Cell viability assay. Cell viability was measured by the CellTiter-Blue Cell Viability Assay following the manufacturer's protocol. The assay determines the metabolic activity of cells based on the ability of living cells to reduce the nonfluorescent reporter compound resazurin into a fluorescent resorufin. Approximately 15 000 cells were seeded into each well of a 96-well black wall plate (PerkinElmer, Waltham, MA) coated with 0.1% gelatin. After culturing for 6 h in 100 μ l culture media (M199 with Hanks salts, 0.2% BSA, 2.5% fetal bovine serum, 0.2% α-D(+) glucose, 0.01% trypsin inhibitor (lima bean, Type II-L), 1% antibiotic-antimycotic solution) at 40°C with 5% CO₂, media was removed, and treatments applied. Chicken CALCA (0, 10^{-6} , 10^{-8} , 10^{-10} M) with or without 100 ng/ml recombinant human FSH (FSH) dissolved in 100 μl culture medium was applied, and the plates were incubated for 12 or 24 h. All treatments were done in triplicate. At the end of 12 or 24 h of incubation, 20 μ l CellTiter-Blue was added to each well, and incubation continued for an additional 1 h. Following incubation, resorufin fluorescence($544_{\rm Ex}/590_{\rm Em}$) was measured using Victor³ 1420 Multilable Counter (PerkinElmer), and cell viability was calculated as a percentage of viability recorded from vehicletreated cells. The experiment was repeated four times (n = 4).

Statistical Analyses

All analyses were completed by analysis of variance (ANOVA) using the general linear model (GLM) procedure of the Statistical Analysis System (SAS Institute Inc., Cary, NC). Differences between individual means were partitioned further by performing pair-wise comparisons of least square means. A probability level of $P \leq 0.05$ was considered statistically significant. For analysis of real-time PCR results, relative *Calca* mRNA to *Actb* mRNA or *Calcr* mRNA to *Actb* mRNA quantities were first converted from log-linear to linear terms. Data on the effect of gonadal steroid treatments on ovarian *Calca* mRNA abundance are expressed as fold differences compared with the vehicle treatment, whereas data on cell viability are expressed as percentage viable compared with the vehicle treatment for each time point. All data are represented as \pm SEM.

RESULTS

Detection of Calca and Calcr mRNA by RT-PCR

A 204-bp partial *Calca* cDNA corresponding to nucleotides 48-251 (GenBank accession no. X03012) of the ultimobranchial gland Calca gene was detected in total RNA extracted from sexually mature female chicken adipose, liver, hypothalamus, kidney, spleen, and ovary (Fig. 1). A very low level of Calca cDNA was amplified from the pituitary gland RNA, possibly reflecting the reproductive status of the hen, as we earlier found that sexually mature female chickens had lower levels of Calca mRNA in the pituitary gland compared with sexually immature chickens [24]. To further characterize chicken ovarian Calca cDNA expression, RT-PCR was performed using total RNA extracted from the granulosa and theca cell layers of preovulatory follicles (F1, F3, and F6) and prehierarchial follicles (3-5 mm). Both granulosa and theca cell layers in all of the follicles studied expressed Calca cDNA (Fig. 2). The use of either water in place of cDNA or RNA that was not reverse transcribed failed to produce any PCR product (Figs. 1 and 2), confirming the absence of genomic DNA contamination. Likewise, the blood cDNA failed to amplify any PCR product, thereby eliminating blood contamination as the source for Calca mRNA. Furthermore, RNA or cDNA sample integrity was confirmed by successfully amplifying a fragment of glyceraldehyde-3-phosphate dehydrogenase cDNA (data not shown). To determine whether ovarian Calca mRNA shares similarity with that produced by the ultimobranchial gland, the primary source of CALCA in avians, the full-length Calca cDNA was amplified from the F1 follicle granulosa cell RNA. We found that the nucleotide sequence of the 417-bp

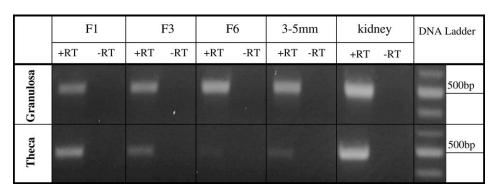


FIG. 3. RT-PCR analysis of *Calcr* gene expression in granulosa and thecal cells layers of preovulatory (F1, F3, and F6) and prehierarchial (3–5 mm) follicles, as well as from the kidney. Approximately 100 ng cDNA (+RT) was used as a template to amplify a 495-bp product of chicken *Calcr*. Contamination controls consisted of RNA from each tissue without reverse transcriptase (-).

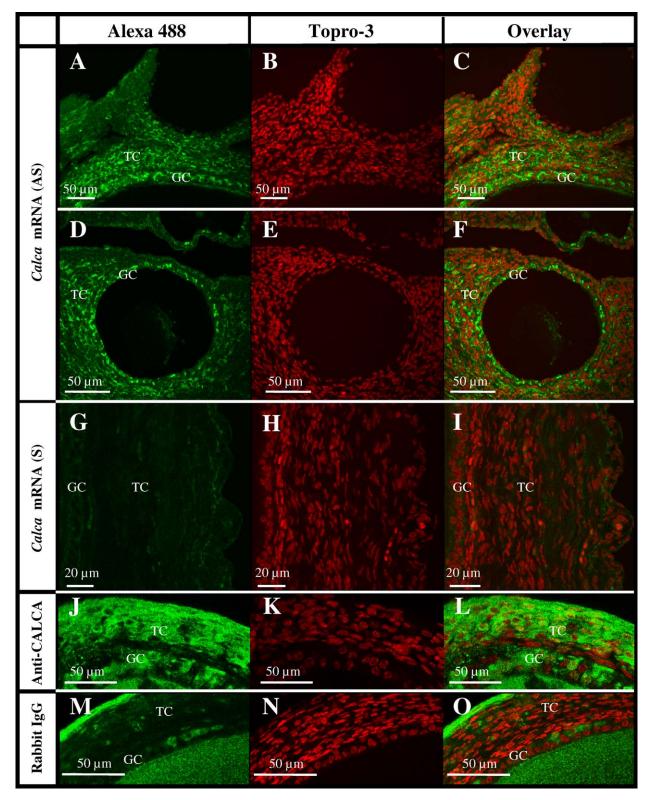


FIG. 4. Confocal photomicrographs of the chicken ovary showing *Calca* mRNA and CALCA-immunoreactive (ir) cells in the follicles. The chicken ovary tissue sections were subjected to in situ hybridization (**A–I**) or immunohistochemistry (**J–O**) to detect *Calca* mRNA and CALCA, respectively. *Calca* mRNA-expressing cells, detected by hybridizing with *Calca* antisense riboprobe, were localized to both the granulosa (GC) and theca (TC) cells (**A–F**). **G–K**) Tissue sections hybridized with sense *Calca* riboprobe as negative control. CALCA-ir cells were detected by immunohistochemistry using rabbit anti-CALCA lgG (**J–L**) or rabbit lgG (**M–O**). Green (CALCA) and red (nuclear staining) fluorescent cells were visualized using Alexa488 and TOPRO-3, respectively, using a laser confocal microscope.

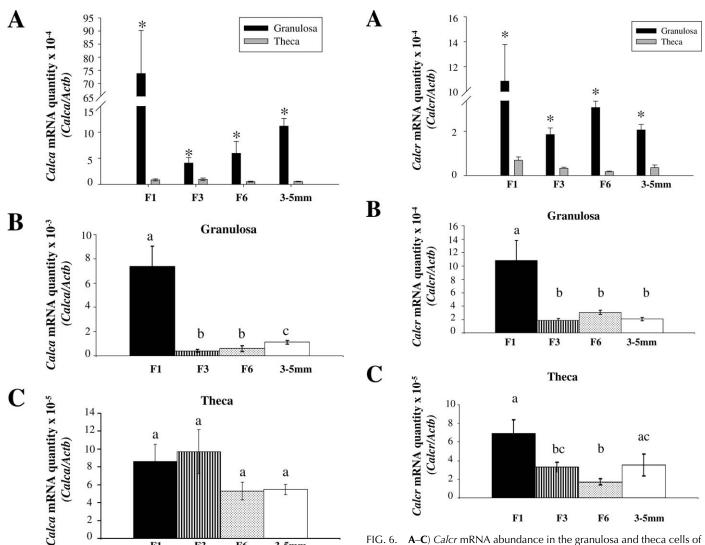


FIG. 5. A-C) Calca mRNA abundance in the granulosa and theca cells of preovulatory (F1, F3, and F6) and prehierarchial (3-5 mm) follicles as measured by real-time quantitative PCR. A) A comparison of Calca mRNA abundance between granulosa and thecal cell layers of preovulatory and prehierarchial follicles. An asterisk above each set of bars indicates significant differences at P < 0.05. **B**, **C**) A comparison of *Calca* mRNA abundance within the granulosa and theca layers. Different letters above each bar indicate significant difference at P < 0.05. Data in **A–C** represent mean \pm standard error of the mean (n = 5 to 6).

F3

F6

3-5mm

F1

ovarian Calca mRNA revealed significant homology (>99%) with that of ultimobranchial gland as well as a predicted chicken Calca cDNA (GenBank accession no. XM_420997).

To determine the presence of the CALCA receptor (Calcr) gene in the ovarian follicles, an RT-PCR was performed using total RNA from the granulosa and thecal layers at various developmental stages (F1, F3, F6, and 3–5 mm) as well as from the kidney (as positive control). The 495-bp product corresponding to Calcr cDNA was amplified in the granulosa cell layers at all stages of development, as well as in the thecal cell layers from the F1 and F3 follicles (Fig. 3). A very low level of Calcr cDNA was amplified from the thecal cell layers of F6 and 3- to 5-mm follicles. The RNA or cDNA sample integrity was confirmed by successfully amplifying a fragment of glyceraldehyde-3-phosphate dehydrogenase cDNA for both granulosa and thecal cell layers at all developmental stages (data not shown). Furthermore, sequencing of the ovarian

FIG. 6. A-C) Calcr mRNA abundance in the granulosa and theca cells of preovulatory (F1, F3, and F6) and prehierarchial (3-5 mm) follicles as measured by real-time quantitative PCR. A) A comparison of Calcr mRNA abundance between granulosa and thecal cell layers of preovulatory and prehierarchial follicles. An asterisk above each set of bars indicates significant differences at P < 0.05. **B**, **C**) A comparison of *Calcr* mRNA abundance within the granulosa and theca layers, respectively. Different letters above each bar indicate significant difference at P < 0.05. Data in **A–C** represent mean \pm standard error of the mean (n = 5 to 6).

Calcr cDNA revealed significant homology (>99%) to that of kidney as well a predicted chicken Calcr cDNA (GenBank accession no. XM_425985).

Cellular Localization of Calca mRNA and CALCA in the Chicken Ovary

Calca mRNA-expressing cells were localized by in situ hybridization histochemistry in both granulosa and thecal cells of the ovary follicle (Fig. 4, A–F). No *Calca* mRNA-specific staining was observed when a sense riboprobe was used in place of antisense cRNA probe (Fig. 4, G-I). Similar to Calca mRNA-containing cells, CALCA-immunoreactive cells, detected by immunohistochemistry, were found distributed in both the granulosa and thecal cells (Fig. 4, J-L). Both Calca mRNA and CALCA were located in the cytoplasm of granulosa or thecal cells. In addition, when rabbit IgG was used in place of the anti-CALCA antibody, no CALCAspecific staining was evident in these cells (Fig. 4, M-O).

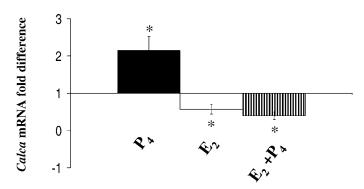


FIG. 7. Effect of P_4 and/or P_2 on Calca mRNA quantity in the chicken ovary. Calca mRNA abundance in response to P_2 and/or P_4 treatment is represented as fold difference compared with vehicle control. An asterisk above each bar indicates significant difference at P < 0.05 compared with vehicle control. The data represent mean P_2 standard error of the mean P_3 (n = 7).

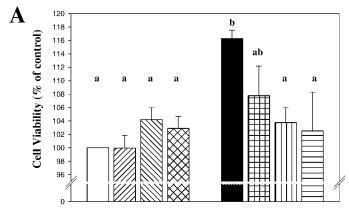
Effect of Follicular Development on Calca and Calcr mRNA Quantity

The diameters of preovulatory and prehierarchial follicles (n = 6 chickens) included in this experiment are as follows: 36.2 ± 0.65 mm for F1 follicles; 27.0 ± 0.84 mm for F3 follicles; 10.5 ± 0.43 mm for F6 follicles; and 3–5 mm for prehierarchial follicles. The granulosa cell layers of all preovulatory follicles (F1, F3, and F6) and prehierarchial follicles (3–5 mm) contained significantly greater abundance of Calca mRNA versus the thecal cell layer (P < 0.05; Fig. 5A). Within the granulosa cell layer, the quantity of Calca mRNA in F1 follicles was significantly higher (P < 0.05; Fig. 5B) compared with that in the granulosa cell layer of other follicles studied (F3, F6, and 3-5 mm). Relative to F3 and F6 follicles, the granulosa cell layers of the prehierarchial follicle (3–5 mm) contained significantly higher Calca mRNA quantity (P < 0.05). In the thecal cell layer, Calca mRNA quantity was not significantly different among preovulatory or prehierarchial follicles (P > 0.05; Fig. 5C). Melting curve analyses showed the presence of a single PCR product for Calca mRNA or Actb mRNA, confirming the specificity of the reaction (data not shown).

In addition, the granulosa cell layers also contained significantly greater levels of Calcr mRNA versus the thecal cell layers (P < 0.05; Fig. 6A) for both preovulatory and prehierarchial follicles. Furthermore, the granulosa layer of the F1 follicle contained significantly higher Calcr mRNA quantity compared with that in the granulosa layer of all other follicles (P < 0.05; Fig. 6B). Within the thecal cell layer, Calcr mRNA quantity was significantly higher in the F1 follicle as well as in the prehierarchial follicles (3-5 mm) relative to F3 and F6 follicles (P < 0.05; Fig. 6C). Melting curve analyses showed the presence of a single PCR product for Calcr mRNA or Actb mRNA, confirming the specificity of the reaction (data not shown).

Effect of Ovarian Steroids on Ovarian Calca mRNA Quantity

 $\rm E_2$ and/or $\rm P_4$ treatment for 7 days resulted in a significant increase in oviduct weights (data not shown). Ovarian *Calca* mRNA quantity was significantly elevated (2.2-fold higher) in response to $\rm P_4$ treatment (P < 0.05; Fig. 7) versus vehicle treatment. $\rm E_2$ or a combination of $\rm E_2$ and $\rm P_4$, however, caused a



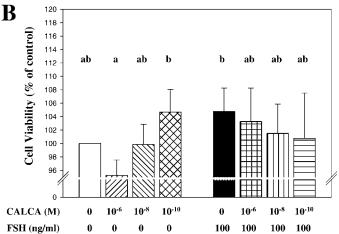


FIG. 8. Effect of chicken CALCA and/or FSH on granulosa cell viability. Granulosa cells were dispersed from the 6- to 8-mm follicles of two hens, pooled, and treated in triplicate with or without FSH (100 ng/ml) and chicken CALCA (0, 10^{-10} , 10^{-8} , 10^{-6} M) for 12 h (A) and 24 h (B). Cell viability was measured using the CellTiter-Blue assay, and viability is expressed as a percentage of vehicle control \pm SEM (n = 4 replicates of two animals each). Different letters above each bar indicate significant difference at P < 0.05.

significant decrease (0.4- or 0.5-fold, respectively) in ovarian Calca mRNA abundance relative to vehicle treatment (P < 0.05, Fig. 7). Melting curve analyses showed the presence of a single PCR product for Calca mRNA or Actb mRNA, confirming the specificity of the reactions (data not shown).

Effect of Chicken CALCA and/or FSH on Granulosa Cell Viability

To determine the role of chicken CALCA on granulosa cell viability in preheirarchial follicles (6–8 mm), granulosa cells were cultured for 12 and 24 h with various concentrations of chicken CALCA and/or FSH, with viability determined by the CellTiter-Blue assay. At 12 h, FSH treatment significantly increased cell viability compared with vehicle treatments (P < 0.05; Fig. 8A), indicating that the cells were responsive to FSH treatment. This stimulation was significantly inhibited by a combination of FSH and CALCA at 10^{-8} and 10^{-10} M (P < 0.05; Fig. 8A). However, no significant changes in cell viability were observed at 12 h in response to CALCA treatment alone. There was no difference in cell viability in response to either CALCA or a combination of FSH and CALCA following 24 h of incubation (Fig. 8B).

DISCUSSION

The present study is the first to characterize the expression of Calca in the ovary of any vertebrate species. By RT-PCR, we identified *Calca* mRNA in the chicken adipose, liver, hypothalamus, kidney, spleen, and ovary. We found that the nucleotide sequence of ovarian *Calca* cDNA was homologous (>99% identities) to the ultimobranchial gland-derived *Calca* [25], indicating that ovarian- and ultimobranchial glandderived CALCA may share biologic properties. We confirmed the expression of Calca in the chicken ovary by in situ hybridization and immunohistochemistry. Both Calca mRNA and CALCA were localized in the granulosa and theca cell layers within the chicken ovarian follicles at various maturation stages. A recent study by Hidaka et al. [26] hints at the presence of Calca mRNA in the ovary, in which Calca cDNA was amplified as a faint band from trout and rat ovarian RNA by RT-PCR. However, in the same study, the presence of Calca mRNA in the ovary could not be confirmed by in situ hybridization. Nevertheless, Calca mRNA has been documented in the testis of rainbow trout [26], as well as endometrium of rats [14], baboons [15], and humans [16], suggesting a role for CALCA in reproduction.

In the present study, *Calcr* gene expression was identified in both the granulosa and theca cell layers of follicles at various maturation stages. Previous studies using receptor binding assays have demonstrated CALCA binding sites in the shell gland endometrium of chicken [27] and guinea fowl [28], as well as in rat Leydig cells [29, 30], suggesting the presence of a specific plasma membrane receptor for CALCA within these reproductive tissues. Moreover, calcitonin receptor-like molecules have also been found in the testis and ovary of flounder [31]. Taken together, these findings further support a possible role for CALCA and its receptor in both male and female reproduction.

While there are no other reports available on the possible functions of ovarian CALCA or related peptides, earlier studies have found a temporal relationship between circulating CALCA levels and ovarian development. In trout and eel, circulating CALCA levels were found to fluctuate with respect to ovarian activity, with a peak observed at the time of ovulation [11, 12]. Plasma CALCA levels were also elevated in the Japanese quail at the time preceding eggshell deposition. However, plasma CALCA levels remained constant throughout the ovulatory cycle in egg-laying hens [32]. In addition, removal of the ultimobranchial gland, the primary source of CALCA in chickens, did not appear to affect egg production [33]. It is possible that the ovarian CALCA found in the present study may possibly compensate for the ultimobranchial gland-derived CALCA at the ovarian level.

In the present study, we found that follicular maturation influenced both Calca and Calcr gene expression. The F1 preovulatory follicle granulosa layer showed a dramatic and abrupt increase in Calca mRNA abundance compared with that of any other preovulatory follicle. Similarly, the granulosa layer of the F1 follicle also had the greatest level of Calcr gene expression, suggesting an important role of CALCA signaling in the most mature preovulatory follicle destined for ovulation. In contrast, relatively low levels of both Calca mRNA and Calcr mRNA were found in the smaller preovulatory and prehierarchial follicle granulosa cells. Since the granulosa cells of the prehierarchial follicles remain undifferentiated compared with the fully mature granulosa cells found in the F1 follicle [34], ovarian CALCA may function as a local regulator of cell differentiation and therefore influence follicular maturation. Likewise, a closely related peptide belonging to Calca gene

family, adrenomedullin, has previously been identified in rat granulosa cells, where it is believed to function as an autocrine regulator of cellular differentiation [35].

To further investigate the role of CALCA on cell differentiation, we treated cultured granulosa cells obtained from prehierarchial follicles with chicken CALCA and/or FSH, and we quantified cell viability using a fluorescence-based detection method. Despite previous reports of CALCA altering proliferation in various cell types (i.e., lactotrophes [36], mammary epithelial cells [37], prostate cancer cell lines [38], HEK cell line [39]), the present study failed to find any significant changes in granulosa cell viability when CALCA was applied alone. However, a combination of CALCA and FSH did cause a significant decrease in cellular viability compared with cells treated with FSH alone following 12 h of treatment. This decreased viability may represent the ability of CALCA to inhibit FSH-induced proliferation [40], as previous reports have correlated cell viability with mitochondrial activity [41] and cell proliferation [42, 43]. Alternatively, CALCA may have inhibited other FSH-induced functions in granulosa cells, such as promoting cell survival, increasing steroidogenic potential, or inhibiting apoptosis [44–46]. Following 24 h of CALCA treatment, however, this inhibitory effect was not observed, possibly due to degradation of CALCA in culture media over time.

Based on CALCA's ability to decrease viability of FSHstimulated granulosa cells, we propose that intraovarian CALCA may be functioning as a local regulator of granulosa cell proliferation in the preheirarchial follicles, thereby prohibiting the entry of numerous follicles into the preovulatory hierarchy. In the viability assay, the greatest inhibition was observed with the lowest concentration of CALCA (10^{-10} M), whereas the highest CALCA concentration (10^{-6} M) did not further exaggerate this response, possibly indicating receptor saturation at such supraphysiologic levels. Interestingly, the preheirarchial follicles were also found to have the lowest level of Calca mRNA expression and contained the greatest amount of FSH receptors [47, 48], and therefore may be the most susceptible to this inhibitory effect. In addition, these prehierarchial follicles are the main producers of estradiol [49], which was shown to decrease total ovarian Calca mRNA expression (Fig. 7). In this manner, estradiol may be acting to maintain the low CALCA concentrations within the prehierarchial follicles so as to enable the granulosa cells to be maximally responsive to the inhibitory effect of CALCA.

The role for CALCA in follicular maturation is further supported by the ability of gonadal steroids, which fluctuate throughout the ovulatory cycle, to alter ovarian Calca expression. In the present study, treatment of sexually immature chickens with estradiol caused a significant reduction in ovarian Calca mRNA levels. Previous studies have shown that estradiol inhibits Calca expression in the endometrium [14] and pituitary [50]. In contrast, estradiol administration to goldfish stimulated ultimobranchial CALCA secretion [51], and it had similar effects in vitro on CALCA secretion from rat thyroid glands [52]. This variable response may further substantiate the various roles for CALCA in different organs. We also found that P₄ administration to sexually immature chickens caused a significant increase in Calca mRNA levels in the ovary compared with vehicle treatment. As the follicle matures in the chicken ovary, the granulosa cell layer undergoes differentiation to produce large quantities of P, [53]. The largest, most mature follicle (F1), which produces the highest amount of P4, was found in this study to express the highest levels of Calca mRNA. In support of our data, P. treatment has been shown to increase Calca gene expression in

rat thyroid glands [52] and in the endometrium of humans and baboons [6, 15, 16].

Whereas P₄ administration influences Calca mRNA expression in the chicken ovary, CALCA treatment has been shown to significantly decrease basal and hCG-induced P₄ secretion from rat granulosa cells in vitro in a post-cAMP-dependent pathway [54]. However, we did not observe any change in P_4 secretion from the granulosa cells of F1 follicles cultured for 2 h with chicken CALCA at 10^{-6} , 10^{-8} , and 10^{-10} M \pm 5 ng/ml oLH (data not shown). Such a lack of response to CALCA treatment may be due to differences in the source of ovarian P₄ production between mammalian and avian species. Alternatively, the granulosa cells from F1 follicles are possibly resistant to the inhibitory effect of CALCA on P₄ production in the chicken. Although not immediately involved in steroidogenesis regulation, the large increase in both Calca and Calcr mRNA observed in the granulosa layer of the F1 follicle may facilitate ovulation through increased activation of plasminogen activators. As the follicle matures, the activity of plasminogen activator, a protease believed to be involved in follicular rupture and ovulation, drastically increases in both the granulosa and theca cell layers [55]. Interestingly, CALCA has been shown to increase plasminogen activator production up to 1000-fold in porcine renal tubular cells in vitro [56]. It is therefore possible that increased CALCA within the F1 follicle increases plasminogen activator, thereby aiding in follicular rupture and ovulation.

In conclusion, we found that *Calca* mRNA, CALCA, and *Calcar* mRNA are expressed by the granulosa and thecal cells in the chicken ovarian follicles. A dramatic increase in both *Calca* mRNA and *Calcr* mRNA abundances in the F1 follicle granulosa cell layer suggests its possible role in follicular maturation and granulosa cell differentiation. In addition, both E₂ and P₄ were found to influence *Calca* mRNA abundance in the chicken ovary. Furthermore, we found that chicken CALCA significantly decreased the FSH-stimulated viability of prehierarchial follicular granulosa cells. Further studies are therefore required to characterize additional role(s) of CALCA and CALCR in the ovary.

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