

Identification of differentially expressed genes involved in the regression and development of the chicken Müllerian duct

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ABSTRACT Müllerian ducts of male chickens undergo regression around day 12 of incubation, but the underlying mechanisms remain unclear. The purpose of this study was to identify factors that contribute to regression of the Müllerian duct in the chicken. We first employed annealing control primer-based RT-PCR to screen candidate genes differentially expressed in the Müllerian ducts between male and female. Four differentially expressed genes (*MSX2*, *GAL10*, *VCP* and *PLCH1*) were partially sequenced. The expression of mRNA of the latter genes and *MSX1* in the male and female Müllerian ducts were compared at 7.5, 8 and 9 days of incubation using semi-quantitative RT-PCR. The results indicated that both *MSX1* and *MSX2* mRNA was highly expressed in the male Müllerian duct at day 9 of incubation, whereas, *PLCH1* mRNA was lower in the male duct at day 9 of incubation compared to that of the female duct. Although *VCP* mRNA was expressed in both left and right female Müllerian ducts, no expression was detected in the male duct. Whole mount *in situ* hybridization analysis showed that the expression of *MSX1* and *MSX2* mRNA were localized specifically in the mesenchymal cells of the male Müllerian duct at day 9 of incubation. In contrast, *VCP* mRNA expression was observed in both mesenchymal and epithelial cells of the female Müllerian duct but not detected in the male duct. These results suggest that both up-regulation of *MSX1* and *MSX2* mRNA expression is involved in the regression of the Müllerian duct in male chicken embryo, whereas *VCP* expression is involved in development of the female duct.

KEY WORDS: *chicken, embryo, Müllerian duct, annealing control primer-based RT-PCR*

Introduction

Most vertebrates have a pair of indifferent gonads at the early stages of development that later differentiate into either the testes or ovaries. Subsequently, hormones from male and female gonads induce the secondary sex characteristics, respectively. In female, due to the absence of androgens the mesonephric tubules and Wolffian ducts regress while the Müllerian ducts persist because of an absence of AMH. The Müllerian ducts give rise to the oviducts, uterus, cervix, and contribute to the upper vagina. However in the male, AMH induces the regression of the Müllerian duct by binding to the AMH receptor2 (Allard *et al.*, 2000) in the mesenchymal cells of the duct (Baarends *et al.*, 1994, di Clemente *et al.*, 1994, Roberts *et al.*, 1999). In a recent study, matrix metalloproteinase 2 (MMP2), a degrading enzyme of extracellular matrix, has been identified as one of possible target genes of AMH signal

cascade in mice (Roberts *et al.*, 2002).

In contrast to mammals, in avian species the homogametic sex is a male (ZZ) and the heterogametic sex is a female (ZW) and in most species only the left ovary and oviduct develop in females. In addition, AMH is produced by both the male and female embryonic gonads and AMH has a role in the regression of the Müllerian ducts in the male (Weniger, 1991) and the right Müllerian duct in females (Hutson *et al.*, 1981, Teng, 1987). In the chicken, the male Müllerian ducts cease development coincident with the expression of AMH on day 8 of incubation and immediately undergo regression and disappear by day 12 of incubation (Romanoff, 1960, Groenendijk-Huijbers, 1962, Carre-Eusebe *et al.*, 1996). In the female, the left Müllerian duct

Abbreviations used in this paper: MMP, matrix metalloproteinase; PLCH, phospholipase C eta; VCP, valosin containing protein .

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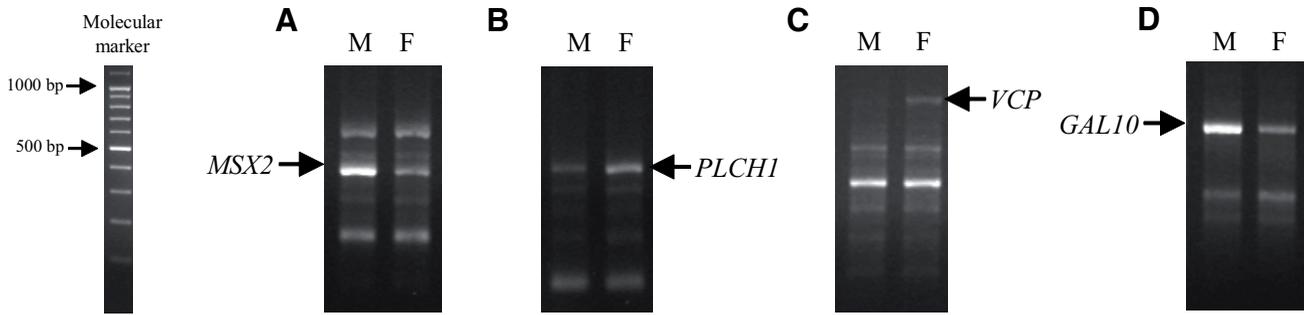


Fig. 1. Examples of gel photographs of genes differentially expressed between male left and right Müllerian ducts and female left Müllerian duct of female embryo. The male (M) and female (F) samples collected between days 7.5-9 of incubation were pooled and used for electrophoresis analysis. Arrows indicate differentially expressed genes.

continues to grow despite the expression of ovarian AMH and develops into the oviduct, whereas, the right duct undergoes a relatively slow involution after day 12 of incubation (Teng, 1987) and completely disappears by the time of hatching (Romanoff, 1960). Estrogen produced by the left ovary prevents AMH from acting on the left female Müllerian duct (Hutson et al., 1985, Teng, 2001). In part, estrogen may prevent the actions of AMH on then Müllerian duct by preventing an increase in expression of *MMP2* in the mesenchymal cells of the Müllerian duct which may cause the degradation of the basement membrane in the duct and lead to cell death (Ha et al., 2004).

While, the regression of the Müllerian duct has been studied to some extent in mammals, the underlying molecular mechanism induced by AMH is still unclear in birds. Accordingly, the present study was conducted to identify factors that contribute to regression of Müllerian duct and that induce development into an oviduct in the chicken by employing the annealing control primer (ACP)-based RT-PCR analysis (Hwang et al., 2003). In this study, four differentially expressed genes, *msh homeobox homologue 2 (MSX2)*, *valosin-containing protein*

(*VCP*) *gallinacin 10 (GAL10)*, *phospholipase C eta (PLCH1)* were obtained. The mRNA expression in the Müllerian duct of these genes was quantified by RT-PCR. Due to a high sequence identity at the nucleotide level compared to *MSX2* (Coelho et al., 1992), *MSX1* mRNA expression was also analyzed and the sex-specific localization of the mRNA in the duct was verified by whole mount *in situ* hybridization.

Results

Genes which are differentially expressed in the Müllerian ducts of the male embryo and the left duct of the female embryo

Figure 1 shows that four bands were obtained using ACP-based RT-PCR analysis. Two of them showed higher expression in Müllerian ducts of male than the female, whereas, others showed higher expression in the left duct of female than in the male. Sequence analysis indicated that those 4 genes were *MSX2*, *VCP*, *GAL10* and *PLCH1*.

Analysis of differentially expressed genes by semi-quantitative RT-PCR

Figure 2 shows examples of PCR products for *MSX1*, *MSX2*, *VCP*, *GAL10*, *PLCH1* and *S17* mRNA of the Müllerian duct at 7.5, 8 and 9 days of incubation. Sizes were close to 332 bp, 277 bp, 418 bp, 385 bp, 134 bp and 351 bp, respectively as predicted by the above sequencing. Figure 3 shows changes in relative mRNA levels in the Müllerian duct at 7.5, 8 and 9 days of incubation following normalization to levels of *S17*. The expression of *MSX1* mRNA was low in both male and female Müllerian ducts at 7.5 and 8 days of incubation (Fig. 3A). However, *MSX1* mRNA levels in the male Müllerian ducts increased markedly at day 9 of incubation and the levels were significantly higher than those of the left and the right ducts of female embryo at 9 days of incubation ($p < 0.05$). *MSX2* mRNA levels in male Müllerian duct were also low at 7.5 and 8 days of incubation (Fig. 3B) but significantly higher than those of the left and the right ducts of female embryo at 9 days of incubation ($p < 0.05$). *VCP* mRNA levels were not significantly different in the left and the right Müllerian ducts of female embryo during the examined days of incubation, but it was not detectable in the ducts of male embryo (Fig. 3C). *GAL10* mRNA levels were relatively high at day 7.5 of incubation in female and male Müllerian ducts (Fig. 3D), however the levels of the left Müllerian

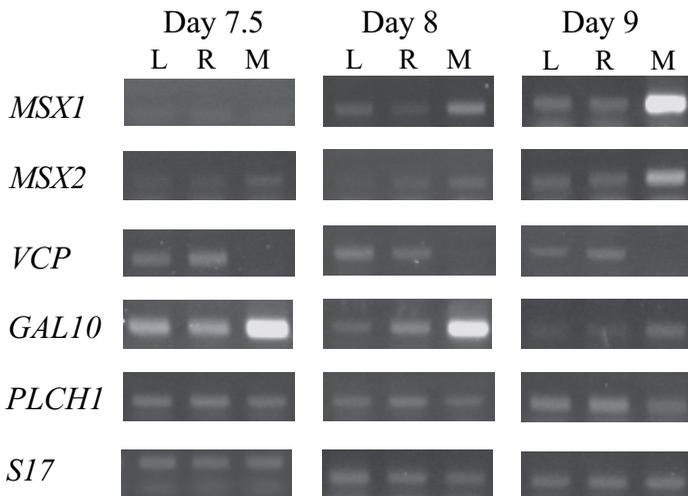


Fig. 2. Detection of mRNA transcripts of *MSX1*, *MSX2*, *VCP*, *GAL10* and *PLCH1* in the Müllerian duct at 7.5, 8 and 9 days of incubation. L, left female Müllerian duct; R, right female Müllerian duct; M, left and right male Müllerian ducts.

duct markedly dropped at day 8 of incubation and were minimal at day 9 of incubation. The levels in male ducts also decreased significantly at day 8 of incubation and remained low at day 9 of incubation. *PLCH1* mRNA levels in the female Müllerian ducts and male duct were almost the same at days 7.5 and 8 of incubation (Fig. 3E), but at day 9 of incubation, *PLCH1* mRNA levels in the left and the right Müllerian ducts of female and the male Müllerian duct increased, respectively. The levels of each duct were significantly different between days 8 and 9 of incubation, respectively ($p < 0.05$). In addition, the levels of *PLCH1* mRNA in the male duct were lower than those of the female ducts ($p < 0.05$).

Whole mount *in situ* hybridization analysis of expression of *MSX1*, *MSX2* and *VCP* in the chicken urogenital system

Figure 4 shows an example of whole mount *in situ* hybridization analysis of *MSX1* expression in the urogenital region at day 9 of incubation. *MSX1* mRNA was only expressed in the Müllerian duct of male embryo but was not detected in the Müllerian duct of female embryo (Fig. 4A). Furthermore, the

expression was not observed in either the testis or mesonephros (Fig. 4B). In the Müllerian duct of male embryo, *MSX1* was detected in the mesenchymal cells and the expression was highly condensed around the epithelial cells (Fig. 4C), but was not detected in the epithelial cells. Figure 5 shows an example of whole mount *in situ* hybridization analysis of *MSX2* expression in the urogenital region at day 9 of incubation. *MSX2* mRNA was expressed only in the Müllerian duct of male embryo but was not detected in the Müllerian duct of female embryo (Fig. 5A). The expression was not detected in either the testis or mesonephros (Fig. 5B). In the Müllerian duct of male embryo, *MSX2* expression was observed broadly within the mesenchymal cells but not detected in the epithelial cells (Fig. 5C). Figure 6 shows an example of whole mount *in situ* hybridization analysis of *VCP* expression in the urogenital region at day 9 of incubation. *VCP* mRNA was expressed in the Müllerian duct of female embryo but was not detected in the Müllerian duct of male embryo (Fig. 6A). The expression was observed in mesonephros of both male and female embryos. *VCP* expression in the gonad was observed in the left ovary but was not observed

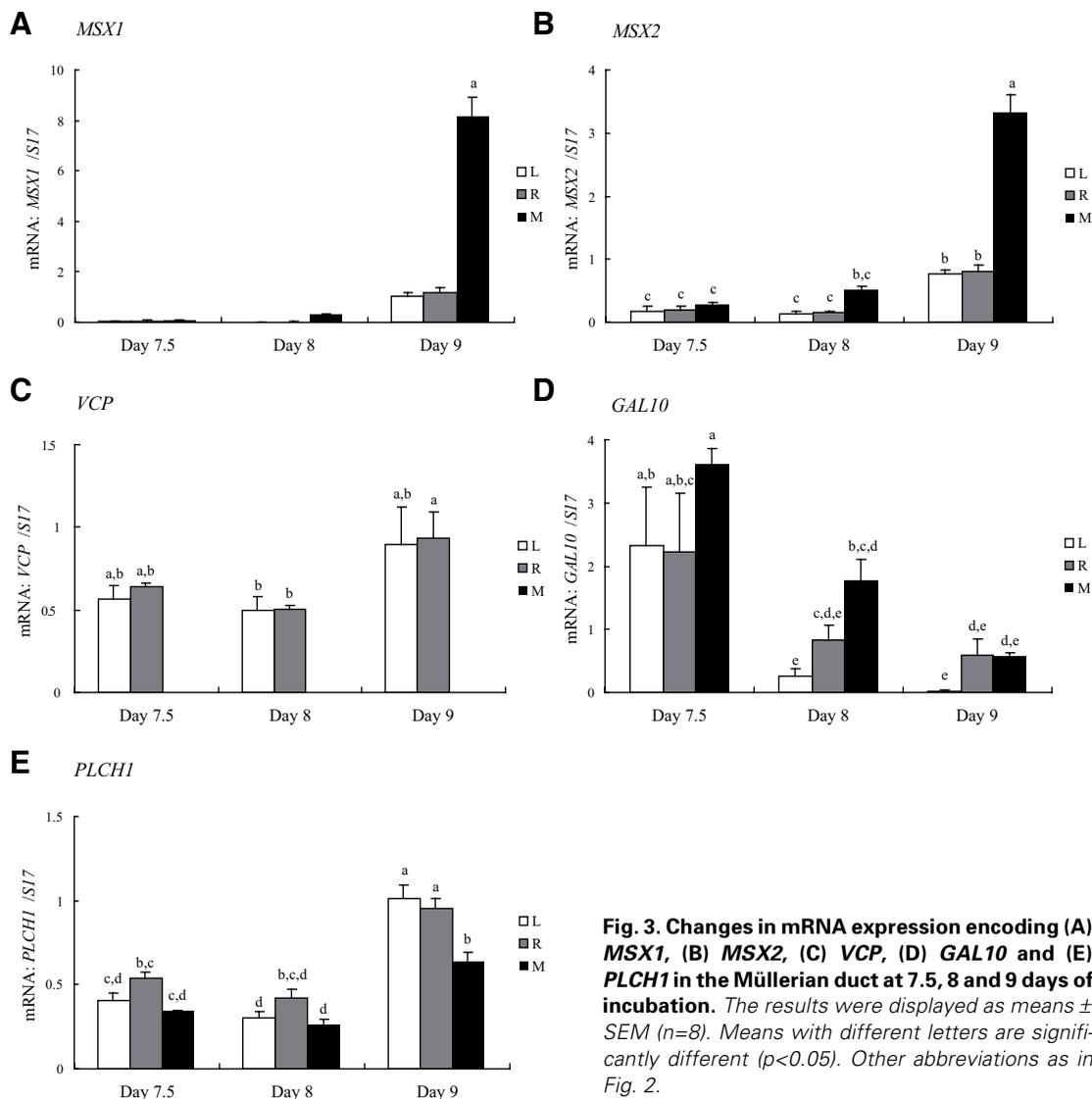


Fig. 3. Changes in mRNA expression encoding (A) *MSX1*, (B) *MSX2*, (C) *VCP*, (D) *GAL10* and (E) *PLCH1* in the Müllerian duct at 7.5, 8 and 9 days of incubation. The results were displayed as means \pm SEM ($n=8$). Means with different letters are significantly different ($p < 0.05$). Other abbreviations as in Fig. 2.

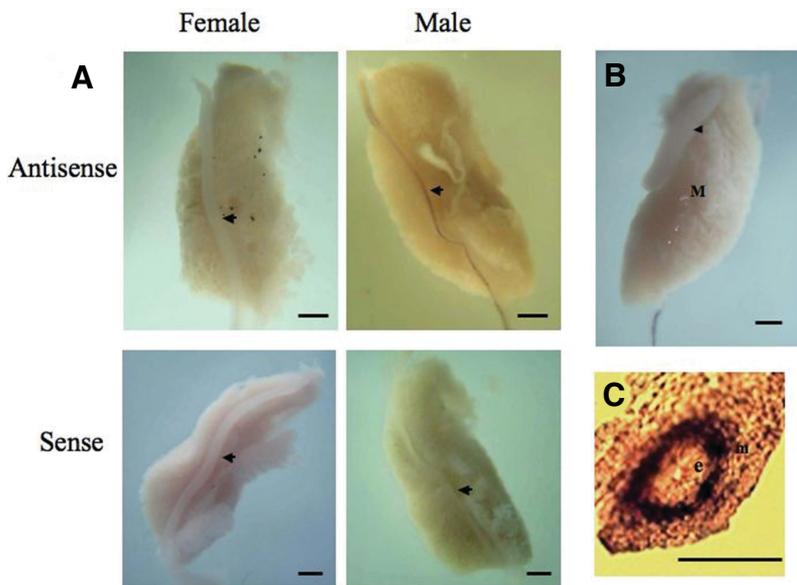


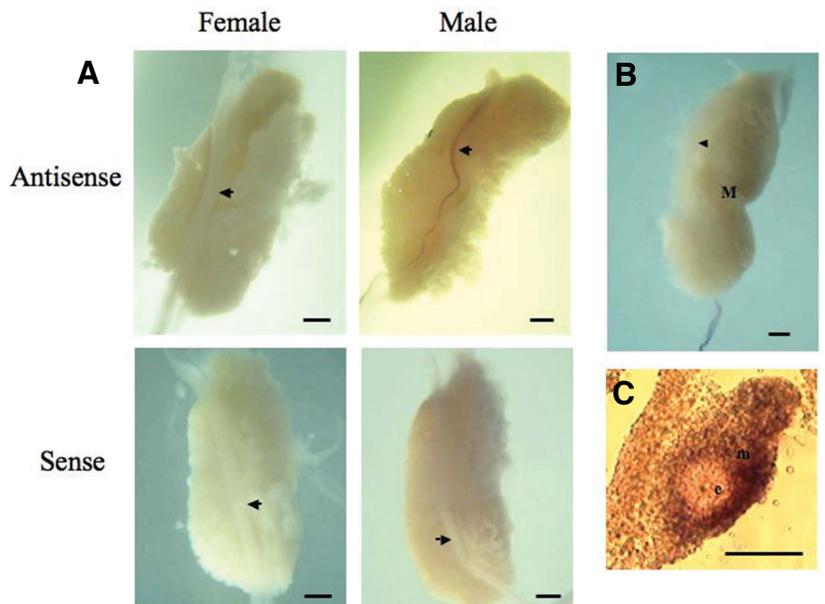
Fig. 4. Expression of *MSX1* in the urogenital region at day 9 of chicken embryo. Whole mount *in situ* hybridization analysis of *MSX1* expression in female and male. **(A)** *MSX1* expression was only detected in the male Müllerian duct, but is not detected in the female Müllerian duct. **(B)** *MSX1* was not detected in testis (arrowhead) and mesonephros (M). **(C)** Cross section of male Müllerian duct shows localization of *MSX1* expression in mesenchymal cells (m); this gene is highly expressed around epithelial cells (e). Scale bars, (A,B) 500 μ m, (C) 0.5 μ m

in the testis (Fig. 6B, C). Although the expression was observed in both epithelial and mesenchymal cells, the expression in epithelial cells was higher than that of mesenchymal cells (Fig. 6D).

Discussion

This is the first report showing that the expression of *MSX1* and *MSX2* genes is involved in the regression of the Müllerian duct of male chicken embryo. Previous studies reported that *MSX1* and *MSX2* are a member of muscle homeobox genes, expressed in a various developing tissue such as limb (Coelho *et al.*, 1991, Coelho *et al.*, 1992), neural tube and crest (Davidson, 1995, Tribulo *et al.*, 2004, Ying *et al.*, 2004), branchial arches (Graham *et al.*, 1996) and regressing Wolffian duct (Yin *et al.*, 2006) where cell death occurs. Our results using RT-PCR assay showed high expression of *MSX1* and *MSX2* in the regressing male duct but not in the developing female duct at day 9 of incubation, indicating a significant role for the simultaneous expression of *MSX1* and *MSX2*.

Fig. 5. Whole mount *in situ* hybridization analysis of *MSX2* expression in the male and female urogenital region at day 9 of incubation. **(A)** *MSX2* expression was only detected in the male Müllerian duct, but was not detected in the female Müllerian duct. **(B)** *MSX2* was not detected in testis (arrowhead) and mesonephros (M). **(C)** Cross section of male Müllerian duct shows localization of *MSX2* expression in mesenchymal cells (m), but not in epithelial cells (e). Scale bars, (A,B) 500 μ m; (C) 0.5 μ m.



Furthermore, whole mount *in situ* hybridization results in this study clearly demonstrated localization of *MSX1* and *MSX2* mRNA in the regressing duct. Namely, both *MSX1* and *MSX2* mRNA were expressed in the mesenchymal cells but not in the epithelial cells, more specifically, *MSX1* mRNA expression in the mesenchymal cells was condensed around the epithelial cell region, whereas *MSX2* mRNA was expressed broadly in the mesenchymal cells. It should be noted that the mesenchymal cells are the sites of *MMP2* expression, which results in degradation of the basement membrane leading to the regression of the Müllerian duct (Roberts *et al.*, 2002, Ha *et al.*, 2004). We propose that *MSX1* and *MSX2* are new mesenchymal factors in the Müllerian duct involved in the duct regression in the male chicken embryo.

VCP gene was also shown to have sexually dimorphic expression in the Müllerian duct. *VCP* belongs to AAA (*ATPases associated with various cellular activities*) family, and is essential for a wide variety of cellular functions such as ubiquitin/proteasom-dependent protein degradation, membrane fusion, cell cycle regulation, and endoplasmic reticulum-associated degradation (Latterich *et al.*, 1995, Zhang *et al.*, 2000, Dai and Li, 2001, Ye *et al.*, 2003, Wang *et al.*, 2004, Vandermoere *et al.*, 2006). Expression of *VCP* mRNA was female specific as indicated by the RT-PCR results and clearly demonstrated by whole mount *in situ* hybridization in the present study. In particular, *VCP* mRNA was strongly expressed in the posterior part of the duct, and expressed broadly in the epithelial cells and mesenchymal cells in females. It has been reported that RNA interference (RNAi) of *VCP* reduced proliferation and promoted apoptosis of HeLa cells, indicating a crucial role of *VCP* for cell survival (Wojcik *et al.*, 2004, Vandermoere *et al.*, 2006). These results and the present data suggest that *VCP*

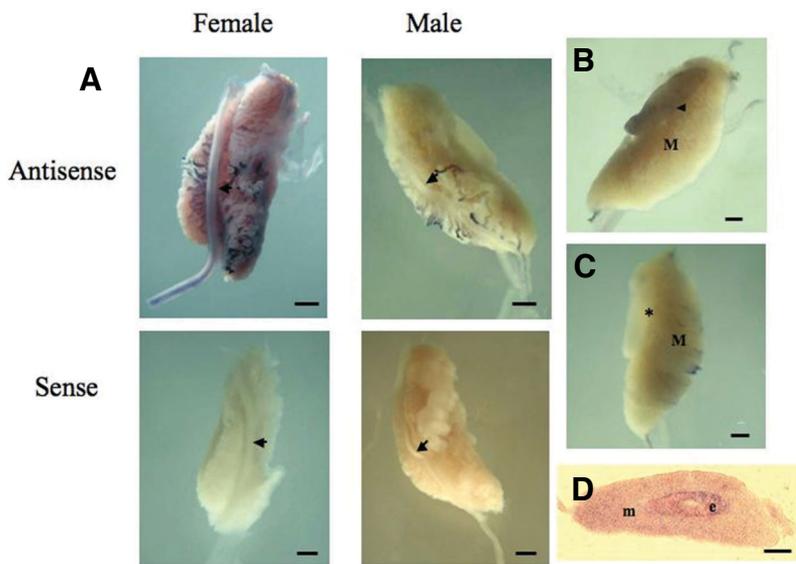


Fig. 6. Whole mount *in situ* hybridization analysis of valosin-containing protein (VCP) expression in the male and female urogenital region at day 9 of incubation. (A) VCP expression was only detected in the female Müllerian duct; it was not detected in the male Müllerian duct. (B) VCP expression was also detected in the left ovary (arrowhead) and mesonephros (M), but not in the testis (*) (C). (D) Cross section of the left female Müllerian duct shows that VCP expression in epithelial cells (e) was higher than that in mesenchymal cells (m). Scale bars, (A,B,C) 500 μ m; (D) 0.5 μ m.

may be involved in cell proliferation and is anti-apoptotic in the Müllerian duct in the female chicken.

PLCH1 is a recently discovered class of isoforms of *phospholipase C* (*PLC*) that is involved in the inositol signaling pathway (Nakahara *et al.*, 2005, Stewart *et al.*, 2007). In females, the expression of *PLCH1* mRNA increased in the Müllerian duct at day 9 of incubation to significantly higher levels than that observed in the males. This suggests that *PLCH1* may play a role in the ducts at about the stage when ducts are committed to either further development or regression. In mammals, *PLCH1* is expressed mainly in neuronal tissue and similar to the other 5 classes of PLC isozymes is involved in protein kinase C activation and calcium metabolism. Since its discovery in 2005, most studies have focused on its role in neurons (Stewart *et al.*, 2007) and this is first report to show that *PLCH1* is expressed in the Müllerian duct during the development. What the functional role may be requires further investigation.

Although left-right asymmetric development of Müllerian ducts is a unique feature of birds, we failed in showing any difference in *MSX1*, *MSX2*, *VCP*, *PLCH1* and *GAL10* mRNA levels between left and right ducts in female chick embryos at least during the period of 7.5 - 9 days of incubation. In this regard, we (Ha *et al.*, 2004) suggested that AMH after binding to AMH receptor in the right duct induces the expression of *MMP2* expression in the mesenchymal cells, which in turn causes the degradation of basement membrane or regressing duct at days 12-18 of incubation in the chicken embryos. Thus, the right duct regression in female starts at day 12 of incubation, we might have missed a crucial day of incubation in this study. Therefore, a possibility that some of the genes are involved in

the regression after day 9 of incubation cannot be excluded. The involvement of *MSX1*, *MSX2* and *VCP* at the later stage of development in the asymmetric formation of Müllerian ducts warrants future study.

In summary, we propose a possible molecular mechanism underlying the regression of Müllerian ducts in the chicken as follows: In males, upregulation of *MSX1* and *MSX2* expression in mesenchymal cells of the duct and low expression of *VCP* around day 9 of incubation may stimulate *MMP2* expression, which provokes degradation of the basement membrane of the Müllerian ducts, resulting to the duct regression. In contrast, in females, high expression of *VCP* mRNA and low expression of *MSX1* and *MSX2* may suppress *MMP2* expression, hence, inhibiting cell death but promoting estrogen-induced cell proliferation or oviductal growth.

Materials and Methods

Embryos

A total of 120 fertilized chicken eggs (White Leghorn) were obtained from a commercial supplier and incubated at 37.5°C under humid conditions. Müllerian ducts were collected as a pooled sample on days 7.5, 8, 9 of incubation for ACP-based RT-PCR analysis. For RT-PCR assay, Müllerian ducts were collected individually at days 7.5, 8 and 9 of incubation. The collected tissues were immediately immersed in RNAlater (Qiagen, Hilden, Germany) and kept at 4°C until use. All procedures for the use and the care of animals were conducted after approval by the Institutional Animal Care and Use Committee of Nagoya University.

ACP-based RT-PCR analysis

Total RNA was isolated from both left and right Müllerian ducts of male embryo and only left Müllerian duct of female embryo on days 7.5, 8, 9 of incubation using Macherey-Nagel Nucleospin RNA II kit (Macherey-Nagel, Germany). Reverse transcription was performed for 90 min at 42°C in a final volume of 20 μ l containing 3 μ g of total RNA, 4 μ l of 5X RT buffer, 5 μ l of dNTP (each 2 mM), 2 μ l of 10 μ M cDNA synthesis primer dT-ACP1 (Gene-Fishing™DEG kits, Seegene, Korea), 0.5 μ l of RNase inhibitor (40 U/ μ l, Invitrogen, Carlsbad, CA, USA) and 1 μ l of Superscript II a reverse transcriptase (200 U/ μ l, Invitrogen, Carlsbad, CA, USA). The reaction was stopped by incubation for 2 mins at 94°C. First-strand cDNAs were diluted by the addition of 80 μ l of diethyl pyrocarbonate (DEPC) treated water.

The PCR reactions were carried out in a final volume of 20 μ l containing the 3 μ l of cDNA, 10 μ M of dT ACP-2 primer, 5 mM of arbitrary ACP primer and 2XSeeAMP™ Master Mix (Gene-Fishing™DEG kits, Seegene, Korea). After incubation at 94°C for 3 min, 50°C for 3 min and 72°C for 1 min, 40 cycles were performed. Each cycle consisted of denaturation at 94°C for 40 sec, annealing at 65°C for 40 sec, and extension at 72°C for 40 sec. A final extension step of 5 min at 72°C was performed to complete the PCR. The 3 μ l of PCR products were run on a 2% agarose gel in 1XTAE buffer, and stained with ethidium bromide.

Sequence of differentially expressed genes

The gel band of the differentially expressed gene was extracted and cloned into a pGEM® T Easy vector (Promega, Madison, WI, USA) and used to transform competent JM109 *E. coli* cells (Takara bio Inc., Shiga, Japan). The colonies were grown for 16 hr at 37°C on LB agar plates

containing ampicillin, X-gal and IPTG for colony selection. The plasmids were extracted and sequenced using an automated DNA sequencer (Applied Biosystems, Foster City, CA, USA). Sequences were analyzed by searching for similarities using the BLAST of NCBI GenBank.

Semi-quantitative RT-PCR

Total RNA was isolated from Müllerian ducts of male embryo and left Müllerian duct of female embryo at 7.5, 8 and 9 days of incubation using Trizol™ Reagent (Invitrogen, Carlsbad, CA, USA). 0.4 µg of total RNA was reverse transcribed in 20 µl reverse transcription buffer (50 mM Tris-HCl, 75 mM KCl, 3 mM MgCl₂, pH 8.3) containing 10 mM dithiothreitol (DTT), 0.5 mM dNTPs, 0.5 µg oligo (dT) 12-18 and 100 units Superscript II a reverse transcriptase (200U/µl, Invitrogen, Carlsbad, CA). Reverse transcription was carried out at 42°C for 50 min, and incubated at 70°C for 15 min to inactivate the reverse transcriptase. First-strand cDNA was diluted by the addition of 30 µl of diethyl pyrocarbonate (DEPC) treated water. Two microlitres of reverse transcribed cDNA was used for each PCR amplification. PCR reactions were carried out in 25 µl PCR buffer (20 mM Tris-HCl, 100 mM KCl, 2 mM MgCl₂, pH 8.0) containing 0.2 mM dNTPs, 0.5 µM each gene specific primers (Table 1) and 0.25 unit of Taq DNA polymerase (Takara bio Inc., Shiga, Japan). All PCR reactions were denatured for 2 min at 94°C. PCR condition for chicken *MSX2*, *GAL10* and *VCP* consisted of 30 sec of denaturation at 94°C, 30sec of annealing of 54°C, and 30 sec of extension at 72°C for 24 cycles. PCR condition for chicken *MSX1* consisted of 30 sec of denaturation at 94°C, 30sec of annealing of 50°C, and 30 sec of extension at 72°C for 32 cycles. PCR condition for chicken *PLCH1* and *S17* ribosomal protein consisted of 30 sec of denaturation at 94°C, 30sec of annealing of 60°C, and 30 sec of extension at 72°C each for 25 and 24 cycles. PCR reactions were completed with an additional extension at 72°C for 5 min. *S17* was used for standardization. The optimal number of cycles for each gene was determined by comparing band intensity after 17, 22, 27, 32 and 37 cycles of PCR. The 24 cycles for *MSX2*, *GAL10*, *PLCH1*, *VCP* and *S17*, and 32 cycles for *MSX1* were selected for the optimal cycles, respectively. All PCR reactions were carried out on a Perkin-Elmer 9700 thermocycler (Applied Biosystems, Foster City, CA, USA). PCR products were run on a 0.8% agarose gel in 1XTAE buffer, and bands were visualized by ethidium bromide staining. The bands of the electrophoretic gels were analyzed using NIH image program (developed at the U.S. National Institutes of Health and available on the Internet at <http://rsb.info.nih.gov/nih-image/>) and the level of *MSX1*, *MSX2*, *GAL10*, *VCP* and *PLCH1* were estimated after normalization to the levels of ribosomal protein *S17* mRNA in each sample.

Whole mount in situ hybridization

Urogenital systems were isolated from embryos on day 9 of incubation. Tissues were fixed overnight at 4°C in 4% paraformaldehyde in phosphate buffered saline (PBS), then dehydrated using a graded metha-

nol series on ice. Sense and antisense riboprobes, specific to each gene, were transcribed *in vitro* from linearized plasmid containing the *MSX1*, *MSX2* and *VCP* cDNA fragment (TABLE 1) using digoxigenin-labeled UTP (Roche, Germany) and SP6 or T7 RNA polymerase (Roche, Germany). Tissues were hybridized in a solution containing 500ng/ml of riboprobe overnight at 55°C. After hybridization, staining was conducted by incubation in nitroblue tetrazolium/5-bromo-4-chloro-3-indolyl phosphate (NBT/BCIP). Staining was only observed in tissues hybridized with antisense probe. For sectioning, tissues were embedded in paraffin and 8 µm sections cut on a microtome. Sections were observed after counterstaining with 0.02% eosin.

Statistical analysis

Data were presented as means ± SEM. The data were analyzed by a two-way analysis of variance (ANOVA) and a Tukey-Kramer honest significant difference (HSD) test ($\alpha=0.05$) using JMP version 5.1.2 (SAS Institute, Cary, NC, USA).

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TABLE 1

PRIMER SEQUENCES USED IN RT-PCR

Gene	Location	Sequence (5'→3')	Accession number
<i>MSX1</i>	790-806	GCTGCATTCCGGCATCTC	NM_205488
	1105-1121	AGAAATGTCCTACTGGG	Coelho CND et al. 1992
<i>MSX2</i>	829-846	CCGTTGGACTCTATGCTA	NM_204559
	1087-1105	ACACTCTGAGGATCAGACT	
<i>VCP</i>	2031-2047	GACATCATTGACCCAGCT	NM001044664
	2431-2448	GGCTCTGCTGTAGAGTCT	
<i>PLCH1</i>	6500-6519	GCTAAAGTGTGTAGTCTGT	Xm422832
	6667-6683	ACTTACAGCAAGGCAA	
<i>GAL10</i>	218-235	CCCACCTTCCACATCTCT	NM00100160
	583-602	CTTCATATGCTCCAAGAGCT	
<i>S17</i>	56-75	GCAACGACTTCCACACCAAC	X07257
	387-406	CAACATAACGAGCGGCTCAG	

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