

Changes in estrogen receptor- α variant (ER- α 36) expression during mouse ovary development and oocyte meiotic maturation

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Abstract The biological effects of estrogens are largely mediated through estrogen receptors (ERs), which belong to the nuclear receptor gene family of transcription factors. ER- α 36 has been recently identified as a new variant of ER α , but its expression and roles in female reproduction system remain unknown. Immunocytochemistry and confocal microscopy were employed to observe ER- α 36 distribution in mouse ovary during postnatal development and in oocyte during meiotic maturation. ER- α 36 was consistently present in the nuclei of oocytes regardless of follicular growth stage and mouse age until germinal vesicle breakdown (GVBD). Its immunosignal was smeared in granulosa cells. However, the ER- α 36 signal is up-regulated and found in cytoplasm with little or no nuclear staining during corpus luteum development. ER- α 36 was also found in theca cells. We showed by Western blot that ER- α 36 was expressed in mouse oocytes at various maturation stages. When the function of nuclear ER- α 36 was blocked by microinjecting anti-ER- α 36 specific antibody into the germinal vesicle (GV) of mouse oocytes, the first polar body emission occurred earlier in a higher proportion of oocytes compared to the control. These results suggest that ER- α 36 may play critical roles in mouse ovarian folliculogenesis and oocyte development.

Keywords Estrogen receptor (ER) α 36 · Ovary · Oocyte · Follicle · Corpus luteum

Abbreviations

ER Estrogen receptor
GV Germinal vesicle
GVBD Germinal vesicle breakdown

Introduction

Estrogens are essential components of female reproduction, with vital roles in the uterus, ovary, mammary gland and hypothalamic-pituitary axis (Hewitt and Korach 2003). However, the exact roles of estrogens in early ovarian folliculogenesis and oocyte meiotic maturation have not been fully defined. Estrogen receptor (ER) knockout (i.e. ERKO), aromatase null (ArKO), and ER overexpression mouse models developed recently have thrown new light on the actions of estrogens in mammalian reproduction (Britt et al. 2002, 2004; Couse et al. 1999b, 2003; Dupont et al. 2000; Fisher et al. 1998; Tomic et al. 2007). Impairment of follicular development and postnatal sex reversal of the ovaries have been reported for immature ER $\alpha\beta$ knockout mice, suggesting that both ERs are required for early ovarian folliculogenesis and the maintenance of germ and somatic cells in the postnatal ovary (Couse et al. 1999b; Dupont et al. 2000). Moreover, ArKO mouse, which possesses functional ERs but does not make endogenous estrogen, can improve the ovarian phenotype, decreased development of Sertoli cells and induced ovulation in some cases after treatment with exogenous estrogen (Britt et al. 2004). In summary, these data show that the biological effects of estrogens are mediated mainly through ERs.

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Estrogen receptors (ER) are ligand-dependent transcription factor that regulate transcription of estrogen-responsive genes in the cell nucleus and belong to the steroid/thyroid hormone receptor superfamily (Evans 1988; Kong et al. 2003), all of which show an evolutionarily and functionally conserved structure (Laudet et al. 1992). The classical estrogen receptor (ER) exists as two subtypes, α and β (Gustafsson 2000; Katzenellenbogen et al. 2000; Kuiper et al. 1998). On other hand, there are also a number of splice variants for the isoforms, some of which influence the activity of the wild type receptor (Hopp and Fuqua 1998; Jazaeri et al. 1999; Schreihofner et al. 2000; Shupnik et al. 1998; Wang et al. 2006). An alternatively spliced variant of ER- α 66, ER- α 36 has been identified recently (Wang et al. 2005). It lacks both transcriptional activation domains of ER- α 66, retains portions of DNA-binding domain, partial dimerization and ligand-binding domains, and possesses a unique 27 amino acid domain that replaces the last 138 amino acid of ER- α 66 (Wang et al. 2005). Recent reports revealed that ER- α 36 localized to plasma membrane and cytoplasm with little or no nuclear staining in breast cancer patients' tissue and breast cancer cell lines (Lee et al. 2008; Wang et al. 2006). Moreover, ER- α 36 can inhibit the transactivation of both ER- α 66 and ER- β , stimulate MAPK signaling pathway and induce cell growth in breast cancer cell lines (Wang et al. 2006).

The expression and function of classical ER α and β in mammalian ovary were reported in several studies. But the distribution of ER- α 36 in mammalian ovary and oocyte, and its role in ovarian folliculogenesis and oocyte meiotic maturation is not known. Therefore, in this study, we examined the expression of ER- α 36 in follicular growth and oocyte maturation in mice.

Materials and methods

Chemicals and antibodies

All chemicals used in this study were purchased from Sigma Chemical Company (St Louis, MO) unless otherwise noted. Rabbit polyclonal anti-ER- α 36 antibody was a gift from Dr Zhao-Yi Wang (Creighton University, California Plaza, USA).

Collection of ovaries and oocytes

Animal care and use were conducted in accordance with the Animal Research Committee Guidelines of the Institute of Zoology, Chinese Academy of Sciences. The mice with color gene type of *aabbcc* were from an inbred strain of Kunming white mice, a native breed widely used in biological research in China. To explore the developmental

expression patterns of ER- α 36 in mouse ovaries, ovaries were removed from postnatal mice at 1, 3, 8, 16, 30, 60 days, respectively, placed in the optimal cutting temperature (OCT) compound (Tissue-Tek), snap frozen in liquid nitrogen, and sectioned at 6 μ m in a leica cryostat microtome (Leica CM1900, Germany) for immunostaining. Oocytes displaying a germinal vesicle (GV) were collected from ovaries of 6-week-old female mice in M2 medium (Sigma, St Louis, MO). Then, the oocytes were washed thoroughly and cultured in M16 medium (Sigma, St Louis, MO) at 37°C in a humidified 5% CO₂ incubator. At different times after culture, only oocytes that underwent GVBD between 1 and 2 h of culture were used for either immunostaining or Western blot analysis.

Immunofluorescent staining and confocal microscopy

Tissue sections from three mice for each time point were randomly selected and immunostained for ER- α 36. Sections were fixed with 4% paraformaldehyde in PBS (pH 7.4) for 5 min, rinsed three times for 5 min each in PBS, permeabilized with 0.5% Triton X-100 for 10 min and then blocked in 1% BSA-supplemented PBS (blocking solution) for 1 h at room temperature. Following the blocking, sections were exposed overnight to anti-ER- α 36 antibody diluted 1:100 with blocking solution. After three washes in PBS containing 0.1% Tween 20 and 0.01% Triton X-100 (washing solution) for 5 min each, the sections were labeled with FITC-conjugated goat anti-rabbit IgG antibody diluted 1:100 with washing solution for 1 h at room temperature. After three washes in washing solution, nuclear status of cells were evaluated by staining with PI (propidium iodide, 10 μ g/ml in PBS) for 5 min. Following extensive washing, sections were mounted on glass slides.

For staining of ER- α 36 in oocytes, oocytes were fixed with 4% paraformaldehyde in PBS (pH 7.4) for at least 30 min at room temperature. After being permeabilized with 0.5% Triton X-100 at room temperature for 20 min, oocytes were processed identically as mentioned above. Each experiment was repeated three times, and at least 30 oocytes were examined each time.

Nonspecific staining was determined by substituting primary antibodies with normal rabbit IgG. Sections and cells were observed under a Confocal Laser-Scanning Microscope (Zeiss LSM 510 META, Germany). The same instrument settings were used for each replicate.

Western blot analysis

For detection of ER- α 36 expression, proteins from 300 oocytes at the appropriate stage of maturation were collected in SDS sample buffer and heated for 4 min at 100°C. After cooling on ice and centrifugation at 10,000g for

3 min, samples were frozen at -20°C until use. The total proteins were separated by SDS-PAGE with a 4% stacking gel and a 10% separating gel at 90 V, 0.5 h and 120 V, 2.5 h, respectively, and then electrophoretically transferred onto polyvinylidene difluoride membrane for 2.5 h, 200 mA, at 4°C . Membranes were blocked in TBST buffer (10 mM Tris, 150 mM NaCl, 0.1% Tween 20, pH 7.4) containing 5% skimmed milk for 2 h at room temperature and then incubated with polyclonal rabbit anti-ER- α 36 antibody, diluted 1:1,000 in TBST containing 0.5% skimmed milk, overnight at 4°C . After three washes of 10 min each in TBST, the membrane was incubated for 1 h at 37°C with HRP-conjugated goat anti-rabbit IgG diluted 1:1,000 in TBST. Then the membrane was processed using the enhanced chemiluminescence (ECL) detection system (Amersham, Piscataway, NJ). The prestained protein molecular weight marker (Fermentas Life Sciences) used for evaluation of molecular weight of ER- α 36 was a mixture of six proteins with apparent molecular weight of 117, 85, 48, 34, 26, and 19 kDa, respectively. Equal protein loading was confirmed by detection of β -actin. For reprob-ing of β -actin, the membrane was washed in stripping buffer (100 mM β -mercaptoethanol, 20% SDS, 62.5 mM Tris, pH 6.7) to strip off bound antibody after ECL detection at 50°C for 30 min. The membrane was reprob-ed with mouse anti- β -actin antibody (Zhongshan Goldenbridge Biotechnology Co.) diluted 1:1,000, using the same procedure as described above. The secondary antibody was HRP-conjugated goat anti-mouse IgG (1:1,000). All experiments were repeated at least three times.

Antibody microinjection

Microinjection was performed using a Nikon Diaphot ECLIPSE TE 300 (Nikon UK Ltd, Kingston upon Thames Surrey, UK) inverted microscope equipped with Narishige MM0-202N hydraulic three-dimensional micromanipulators (Narishige Inc., Sea Cliff, NY, USA). A microinjection volume of 7 pl of anti-ER α 36 antibody was injected into the nucleus of a fully-grown GV oocyte in all experiment. The oocytes were kept in M2 medium supplemented with 0.2 mM IBMX to prevent GV breakdown during the injection period. After microinjection, oocytes were washed twice in M16 and then cultured. Control oocytes were microinjected with the same amount of rabbit IgG. Each experiment was repeated three times, and at least 30 oocytes were examined each time.

Statistical analysis

All percentages from three repeated experiments were expressed as mean \pm SEM and the number of oocytes observed was labeled in brackets as (n). Data were

analyzed by χ^2 test. Differences at $p < 0.05$ were considered to be significant.

Results

ER- α 36 protein expression in developing mouse ovary

The distribution of ER- α 36 in mouse ovaries collected from mice at different ages was shown in Fig. 1. Strong ER- α 36 immunoreactivity was detected in surface epithelium cells and oocytes at all post-natal age groups examined. In the ovaries of 1-day post-natal (P1) mouse, ER- α 36 immunore-activity was evident in both oocytes and somatic cells. However, oocyte clusters showed more intense immunosig-nal compared to other clusters (Fig. 1 P1). When morpho-logically distinct follicles can be identified at P3, strong ER- α 36 staining was primarily associated with granu-losa cells and oocytes, whereas interstitial cells exhibited low staining. In oocytes, ER- α 36 immunosignal was more intense in nucleus than in cytoplasm, while ER- α 36 immu-nostaining was not clearly delineated to a specific subcellu-lar compartment in granulosa cells. However, the nuclear staining was somewhat more stronger than cytoplasm com-partment (Fig. 1 P3). With further development of the folli-cles by P8, a distinct theca cell layer is formed in secondary follicles. Beside oocytes and granulosa cells, these long-stretched fiber-like theca cells showed evident ER- α 36 immunostaining (Fig. 1 P8). The patterns of ER- α 36 expression had no obvious changes in subsequent develop-ment before ovulation (Fig. 1 P16 and P30). After adminis-tering the ovulatory stimulus, granulosa cells of mature follicle were luteinized and transformed into corpus luteum cells, which presented stronger ER- α 36 immunostaining than that of granulosa cells. In corpus luteum cells, ER- α 36 localized to cytoplasm with little or no nuclear staining (Fig. 1 P60). No ER- α 36-specific immunoreactivity could be detected in sections incubated with normal rabbit IgG, which is a substitute for the primary antibody (Fig. 1 P1nc).

Expression of ER- α 36 in mouse oocytes during meiotic maturation

The ER- α 36 distribution during oocyte meiotic maturation was examined by immunofluorescence. As shown in Fig. 2, in oocytes at germinal vesicle (GV) stage, ER- α 36 was mainly localized to the nucleus. From germinal vesicle breakdown (GVBD) to metaphase II stage during mouse oocyte maturation, there was no concentrated signal of ER- α 36 immunoactivity in oocytes. ER- α 36 could distribute homogenously in the cytoplasm, since Western blot analy-sis showed that ER- α 36, which has an approximate molecu-lar weight of 36 kDa, was expressed in mouse oocytes at all

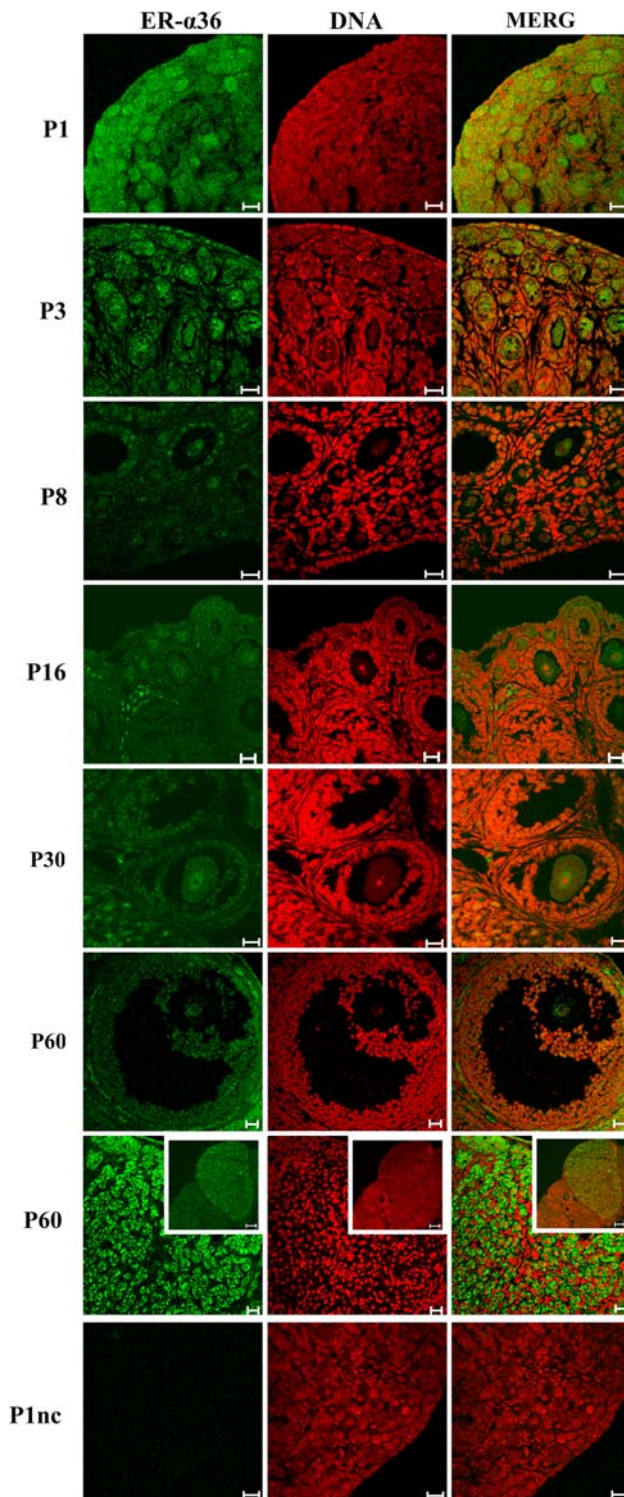


Fig. 1 Immunofluorescent localization of ER- α 36 in mouse ovary during the postnatal development. *Green* ER- α 36, *red* chromatin, *yellow* overlapping of green and red. Sections were from ovaries collected at 1-day-old (P1), 3-days-old (P3), 8-days-old (P8), 16-days-old (P16), 30-days-old (P30) and 60-days-old (P60) post birth. Sections of 1-day-old (P1nc) postnatal mice ovaries stained when the primary antibody was substituted with normal rabbit IgG were used as control. *Bar* 20 μ m. The upper inserts in P60 showed ER- α 36 immunosignal was more intense in corpus luteum cells than in granulosa cell. *Bar* 100 μ m

stages during meiosis with no evident changes (Fig. 2b). To investigate the potential role of ER- α 36 in regulating meiotic maturation of mouse oocyte, we depleted the protein function by microinjecting anti-ER- α 36 specific antibody into the nucleus. After 9.5 h of culture, 32.3% (61/195) of anti-ER- α 36 antibody injected oocytes displayed first polar body (PB1), which was significantly higher than 17.0% (18/106) in the control group ($p < 0.05$) (Fig. 3).

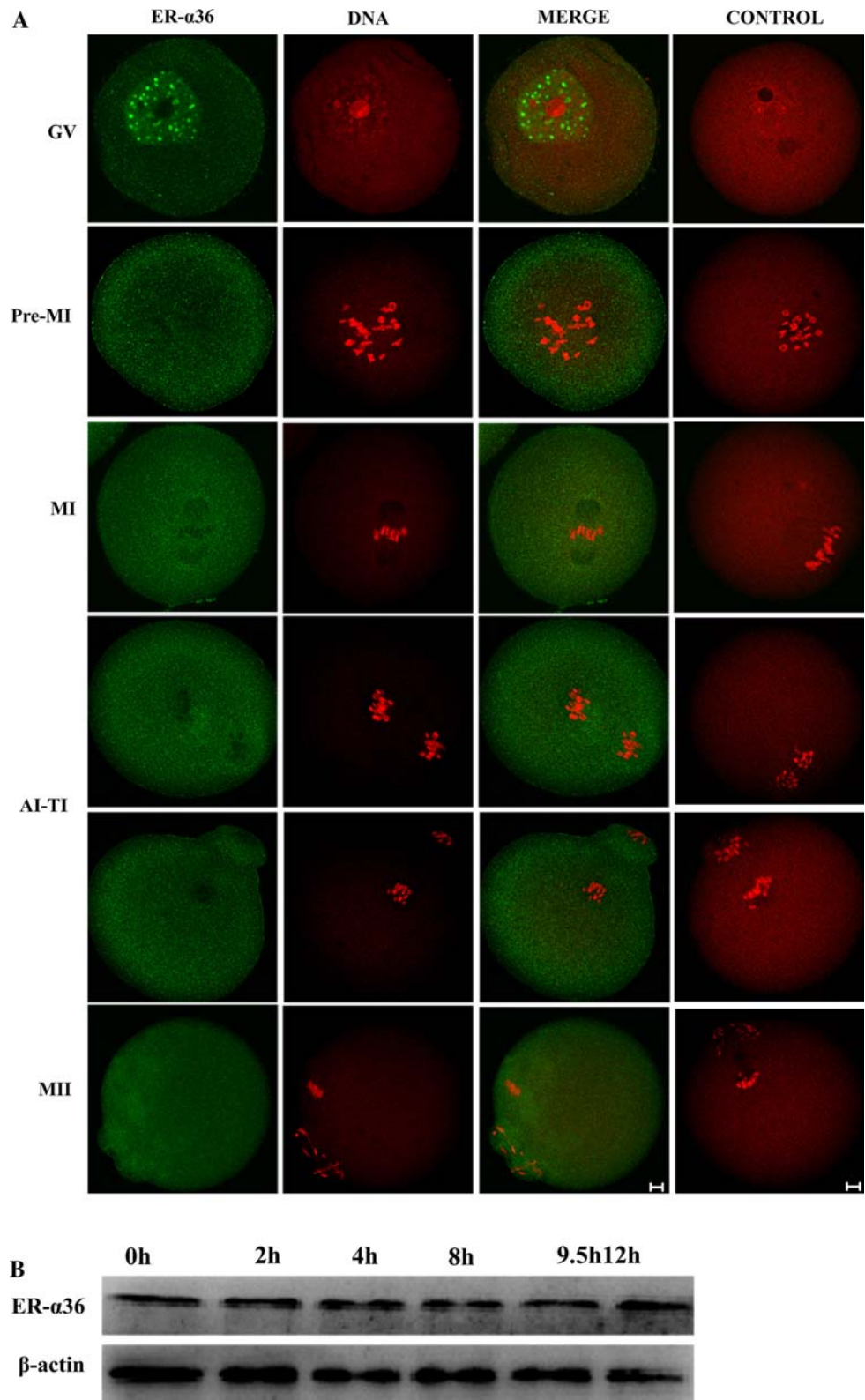
Discussion

This study for the first time revealed the ER- α 36 distribution in postnatal mouse ovary development and in oocyte at different maturation stages. It is found that ER- α 36 was constitutively present in the nuclei of oocytes in spite of follicular growth stage and mouse age until GVBD occurrence. Its distribution changes in granulosa cells, theca cells, and corpus luteum cells were also observed. These data may lay ground for the future research on the critical roles of ER- α 36 in mouse follicle/oocyte growth and maturation.

As described above, ER- α 36 is devoid of both the AF-1 and AF-2 transactivation domains of ER- α 66 but retains the DNA-binding domain, partial dimerization and ligand-binding domains, which implies that ER- α 36 may effectively compete with ER- α 66 for the DNA-binding elements (EREs) in estrogen-responsive genes and suppress the transactivation of ER- α 66. It has been demonstrated that ER- α 36 inhibits estrogen-dependent and estrogen-independent transactivation activity of ER- α 66 and ER β in vitro (Wang et al. 2006). However, the correlation between ER- α 36 and ER- α 66 or ER β in vivo as well as the roles of ER- α 36 in early ovarian folliculogenesis and oocyte meiotic maturation is not known. In the present study, our data showed that ER- α 36 immunosignal is evident in theca and granulosa cells. Several lines of evidence show that ER α is highly expressed in the interstitial/theca compartment, whereas ER β expression is limited to granulosa cells of growing follicles in the mouse ovary (Couse et al. 1997; Tremblay et al. 1997). Our results imply the possibility of the actions between ER- α 36 and ER α or ER β within the same cell type. Additionally, examination of the ovaries of ER α β KO mice indicated a phenotype distinct from those observed in ovaries from ER α KO or ER β KO mice (Couse et al. 1999a, b; Dupont et al. 2000). Therefore, it is reasonable to speculate that a delicate balance of ERs interactions may ultimately determine the nature of estrogen signaling, which is critical for the differentiation of ovarian somatic cells.

The mechanism by which granulosa cells start to differentiate and undergo luteinization has not been well defined. Evidence has accumulated to indicate that ER α is involved

Fig. 2 Localization and expression of ER- α 36 during mouse oocyte maturation. **a.** Immunofluorescent localization of ER- α 36 in mouse oocytes. *Green* ER- α 36, *red* chromatin, *yellow* overlapping of green and red. *GV* oocytes at germinal vesicle, *Pro-MI* oocytes at first prometaphase, *MI* oocytes at first metaphase, *AI-TI* oocytes at first anaphase and telophase, *MII* oocytes at second metaphase. Pictures stained when normal rabbit IgG substituted for primary antibody were used as control. *Bar* 10 μ m. **b** ER- α 36 immunoblotting of cellular extracts from 300 oocytes cultured for 0, 2, 4, 8, 9.5, and 12 h, respectively. A control blot shows equivalent levels of protein loading between various samples



in this process. Female mice null for ER α (ER α KO) are anovulatory, infertile, and possess hyperemic ovaries devoid of corpus luteum in the presence of elevated LH levels (Couse et al. 1999a, 2003). Although immature ER α KO

females do successfully ovulate when treated with exogenous gonadotropins if administered before onset of the overt hypergonadotropic phenotypes in the ovary, but the number of produced corpus luteum is lesser than controls

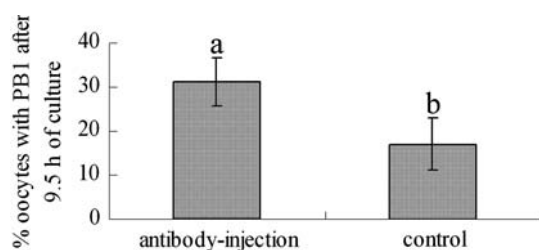


Fig. 3 Microinjection of anti-ER- α 36 antibody into the nucleus accelerates the emission of first polar body in mouse oocytes. Data are presented as mean percentage of PB1 \pm SEM of three independent experiments after oocytes were cultured for 9.5 h. PB1 oocytes with first polar body. Different superscripts denote statistical difference at a $p < 0.05$ level of significance

(Couse et al. 1999a; Rosenfeld et al. 2000). Furthermore, in ER α KO ovaries, there is an almost total lack of detectable progesterone receptor expression after PMSG stimulation (Couse et al. 2005). Our results demonstrated that ER- α 36 expression rapidly increased and its prominent staining was observed in cytoplasm with little or no nucleus staining following ovulation in luteinizing granulosa cells that were undergoing terminal differentiation to form the corpus luteum. During this process, ER- α 36 immunoreactivity displayed a shift from nuclear to cytoplasmic distribution. What might be the actual cause of these changes in ER- α 36 expression is currently unknown. Several groups have shown that expression and subcellular localization of ER α are related to follicular developmental stage and/or hormonal environment in vitro (Lenie and Smitz 2008; Sharma et al. 1999). There is accumulating evidence showing that ERs continuously shuttle between targets located within various cellular components, that is, membrane, cytoplasm, nucleus, etc., and their steady state localization is a consequence of a fine balance between operational strengths of ‘nuclear localization signal’ (NLS) and ‘nuclear export signal’ (NES) (Kumar et al. 2006; Leclercq et al. 2006). This process is regulated by ligand and/or protein-induced ER conformational changes (Leclercq et al. 2006). Another group has found that a pure antiestrogen RU 58668 profoundly modified the subcellular distribution of ER in a specific time-, dose- and protein synthesis-dependent manner. Furthermore, this effect was reversed by RU 58668 withdrawal, together with estradiol treatment (Devin-Leclercq et al. 1998). When granulosa cells proceed to luteinization and shift their main products from estrogen to progesterone, significant changes in the expression of many genes occur. Therefore, we could not exclude the possibility that the NLS of ER- α 36 is sequestered by some protein(s), which results in ER- α 36 staying in cytoplasm, the site of protein synthesis or that NES of ER- α 36 is activated by some protein(s), which leads to ER- α 36 translocation from nucleus to cytoplasm. Report has revealed that prolactin-mediated pathway is highly important in the induction

of ER α in rodent corpus luteum (Frasor and Gibori 2003; Telleria et al. 1998). A similar intense cytoplasmic ER α staining has been reported in luteinized cells of ovine and mouse (Lenie and Smitz 2008; Zieba et al. 2000). It is now accepted that ERs exert their roles in cells by a number of distinct mechanisms. The classical mechanism of ER action involves estrogen binding to receptors in the nucleus, after which the receptors dimerize and bind to specific response elements known as estrogen response elements (EREs) located in the promoters of target genes to activate or suppress the transcriptional activity of target gene (Bjornstrom and Sjoberg 2005). On other hand, the nongenomic actions of membrane-associated ERs result mainly from rapid activation of cellular signaling systems that eventually act on target transcription factors (Bjornstrom and Sjoberg 2005). Activation of the MAPK signaling pathway by ER- α 36 has been established (Wang et al. 2006). Based upon those findings we could speculate that ER- α 36 plays an important role in ovulation and luteinization through nongenomic effects, rapidly responding to LH surge. However, all these above speculations need support by solid evidence from ER- α 36 knockout mice.

Previous study reported that in response to estrogen, an ER α /p300 complex is recruited to an AP-1 domain located in the proximal BRCA-1 promoter and activates BRCA-1 transcription (Jeffy et al. 2005), whereas BRCA-1 could be a candidate controlling spindle checkpoint component in mitotic and meiotic cells (Joukov et al. 2006; Pan et al. 2008). During GVBD, mammalian oocytes undergo a dramatic change in nuclear organization in which chromatin becomes progressively condensed and forms heterochromatin/chromosomes (De La Fuente 2006). When chromatin is diffused, transcription and translation can take place, but when chromatin becomes condensed, transcription is practically inactive (Curtis et al. 1995; Tomek et al. 2002). In the present study, ER- α 36 was mainly concentrated to GV of oocytes regardless of follicular growth stage and mouse age until GVBD occurrence. After GVBD, the signal of ER- α 36 was distributed evenly in cytoplasm. Moreover, we blocked the function of nuclear ER- α 36 by microinjecting anti-ER- α 36 specific antibody into GV of mouse oocytes, and found that the depletion of ER- α 36 induced a significantly higher percentage of oocytes with first polar body earlier compared to control. Thus, we speculate that ER- α 36 may be involved in regulating transcription or translation of some protein related to cell cycle checkpoint proteins to control cytoplasmic maturation in mammalian oocyte maturation. Of course, we can not exclude the possibility that the new element of antibody–antigen complex may stimulate polar body emission after injection of ER- α 36 antibody into the cell. To elucidate and confirm how ER- α 36 regulates the progress of meiotic cell cycle, additional biochemical and molecular biological studies are needed.

In summary, our results indicated that ER- α 36 was mainly present in oocytes, granulosa cells, and theca cells in mouse ovary at different developmental stages. Notably, its immunoreactivity increased during luteinization. When mouse oocytes were cultured *in vitro*, the signal of ER- α 36 was converted from concentrated GV distribution to homogenous cytoplasmic distribution after GVBD. Moreover, ER- α 36 depletion from the GV could cause earlier emission of the first polar body. These data imply that ER- α 36 may play critical roles in female mammalian reproduction, which needs further evidence *in vivo*.

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