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Localization and distribution

of estrogen receptors and

progesterone receptors

in the bovine ovary in

relation to the cell dynamics

Localization and distribution of estrogen receptors and progesterone receptors in the bovine ovary in relation to the cell dynamics

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LIST OF ABBREVIATIONS

А	adenine	LH	luteinising hormone
AEC	amino ethyl carbazole	11	large lutein cells
AS	apoptotic score	mRNA	messenger RNA
С	cytosine	OAF	obliterative atretic follicle
CA	corpus albicans	PBS	phosphate-buffered saline
CAF	cystic atretic follicle	PCR	polymerase chain reaction
СН	corpus hemorrhagicum	PdF	primordial follicle
CL	corpus luteum	PF	primary follicle
cs	capsular stroma	PR	progesterone receptor
DAB	diaminobenzidine	PRID	progesterone releasing intravaginal device
DAPI	4', 6-diamidino-2- phenylindole dihydrochloride	ps	perivascular stroma
DIG	digoxigenin	RNA	ribonucleic acid
DNA	deoxyribonucleic acid	RNase	ribonuclease
DNase	desoxy nuclease	SE	surface epithelium
DS	deep stroma	SEM	standard error of the mean
EDTA	ethylene diamino tetra-acetic acid	SER	score of estrogen receptor
ER	estrogen receptor	SF	secondary follicle
ERα, ESR1	estrogen receptor-alpha	sl	small lutein cells
ER β , ESR2	estrogen receptor-beta	SPR	score of progesterone receptor
ERγ, esr3	estrogen receptor-gamma	SS	superficial stroma
fc	follicle cells	SSPE	sodium chloride sodium phosphate / EDTA
fd	follicle diameter	Т	thymine
FSH	follicle stimulating hormone	ТА	tunica albuginea
G	guanine	te	theca externa
gc	granulosa cells	tf	theca follicularis
HRP	horseradish peroxidase	ti	theca interna
HSS	high sensitive streptavidine	Tris	tris(hydroxymethyl)methylamine
is	internal stroma	TUNEL	terminal deoxynucleotidyl transferase mediated dUTP nick end labeling
kb	kilobase	v/v	volume/volume
kDa	kilodalton	VTF	vital tertiary follicle
KO	knockout		

GENERAL INTRODUCTION

The bovine ovary

Embryology

Harbouring genetic information to be passed down from generation to generation, the ovaries are crucial structures for the survival of the species. The germinal component originates from primordial germ cells that, after migration from the yolk sac, colonize the gonadal primordium. The germ cells together with the surrounding stroma cells form the pool of resting primordial follicles in the ovary (Motta et al., 1997).

The ovary of a bovine foetus of 110 days contains more than 2 million germ cells. However, at birth this number dramatically decreases to ca 100 000 germ cells enclosed in primordial follicles and is progressively further reduced to 2500 in ten years old cattle (Erickson, 1966a, 1966b).

Morphology of the bovine ovary (Fig. 1)

The ovarian cortex and medulla

The bovine ovary consists of a central medulla or zona vasculosa and a peripheral cortex or zona parenchymatosa. The medulla is highly vascularized and consists of connective tissue and a dense network of nerves. The cortex contains both stromal and parenchymatous structures (Schaller, 1992). The ovary is covered by a simple squamous or cuboidal surface epithelium. Beneath the epithelium lies a layer of dense connective tissue named the tunica albuginea and an amount of ovarian stroma, consisting of spindle-shaped cells arranged in whorls (Dellmann and Eurell, 1998). This ovarian stroma can be further subdivided into a superficial part (superficial stroma) and a deep part (deep stroma), both containing ovarian parenchymatous structures.

The ovarian parenchymatous structures

The ovarian parenchymatous structures consist of the follicles in various stages of development, the corpora hemorrhagica, the corpora lutea and the corpora albicantia. The ovarian follicles and corpora hemorrhagica can be found in the superficial stroma, while the corpora lutea and corpora albicantia are mainly embedded in the deep stroma.

Four main types of vital follicles can be distinguished, namely primordial, primary, secondary and tertiary follicles (Nomina Histologica, 1994). A primordial follicle is composed of a primary oocyte enveloped by a single layer of flattened follicular cells (follicle diameter (fd) < 0.04 mm). A primary follicle is composed of a primary oocyte enveloped by a single layer of cuboidal or columnar follicular cells (fd: 0.04-0.08 mm). A secondary follicle consists of a primary oocyte surrounded by a stratified follicular epithelium, a basal membrane, and a developing follicular theca (fd: 0.08-0.13 mm). A tertiary follicle, also called an antral or vesicular follicle, is characterized by the development of a central cavity, the antrum. In the early stages the tertiary follicle has a diameter of 0.13-0.25 mm while in later stages the follicle matures and reaches its maximum size just prior to ovulation (fd: 0.25-20 mm). The latter stage of follicle is termed a preovulatory or Graafian follicle (Gougeon, 1996; Braw-Tal and Yossefi 1997). As the antrum enlarges through the accumulation of liquor folliculi, the primary oocyte is displaced eccentrically, usually in a part of the follicle nearest to the centre of the ovary. The oocyte then lies in a protruding accumulation of granulosa cells, called the cumulus oophorus (Dellmann and Eurell, 1998). The granulosa cells of tertiary follicles are surrounded by a layer of theca cells which differentiates into two layers, viz. an inner vascular theca interna and an outer supportive theca externa. The theca interna cells are spindle-shaped and located in a delicate reticular fiber network. The theca externa layer consists of a thin sheet of loose connective cells arranged concentrically around the theca interna. Blood vessels of the theca externa supply capillaries to the theca interna (Priedkalns et al., 1968; Ham and Cormack, 1979; Braw-Tal and Yossefi, 1997; Dellmann and Eurell, 1998). Graafian follicles are the major source of ovarian estrogens that are produced cyclically during reproductive life. Shortly before ovulation the first meiotic division is finalized, whereby the primary oocyte is transformed into the secondary oocyte and a first polar body. In response to preovulatory gonadotrophin surges the dominant follicle ovulates to release the secondary oocyte for fertilization. After ovulation, the follicular wall collapses and the granulosa layers form large folds protruding into the residual lumen. As a result of the slight hemorrhage, which occurs at the time of ovulation, a blood clot is formed in the centre. The resulting structure is called the corpus hemorrhagicum (Dellmann and Eurell, 1998). This corpus hemorrhagicum further develops into a corpus luteum (or yellow body), which is a transient endocrine gland. The granulosa cells are localized in the centre of the corpus luteum and are referred to as small lutein cells, while theca interna cells are located in the periphery and are called large lutein cells. In the cyclic cow, the corpus luteum is only functional for 17-18 days. In the early luteal stage (days 2-4 of the ovarian cycle) the lutein

cells of the corpus luteum exhibit their greatest ability to proliferate and start secreting progesterone. In the secretory stage (days 5-17) progesterone secretion reaches its maximum levels. A dense network of vessels develops around the corpus luteum and may function in the transport of hormones. Luteal regression occurs from day 18 onwards. Shrinkage of the corpus luteum takes place rapidly after day 19 and is completed 1 to 2 days after oestrus (Dellmann and Eurell, 1998). The corpus albicans is the "white body" and consists of scar tissue formed by regression of the corpus luteum. The colour varies with species and is red or white in the bovine. The corpus albicans is found deeper in the ovary than the corpus luteum and appears very slowly. It is characterized by small lutein cells which enclose a few lipid droplets, and a large amount of connective tissue (Dellmann and Eurell, 1998).

Follicular atresia can be observed at any stage of follicular development. In atretic primordial, primary and secondary follicles the oocyte shrinks and subsequently degenerates, followed by degeneration of the follicle cells (Van Wezel and Rodgers, 1996; Depalo et al., 2003). At the end no scar tissue is left. Atretic changes in tertiary follicles may result in the formation of two different morphological types of atretic follicles, namely cystic and obliterative atretic tertiary follicles. In cystic atretic tertiary follicles the granulosa layer is invaded by vascularized strands of connective tissue, after which the granulosa cells desquamate into the follicular cavity. In more advanced stages of atresia the follicle collapses, the walls are thrown into folds, more connective tissue and blood vessels are found, and rapid resorption of the degenerated granulosa cells takes place. In obliterative atretic follicles, the basement membrane between theca interna cells and granulosa cells thickens and is now referred to as the glassy membrane or membrane of Slavjanski. Subsequently the theca interna cells further increase in size and are arranged in radial layers or cords around the oocyte. Eventually there is a complete breakdown of the obliterative atretic follicles by invading connective tissue cells (Dellmann and Eurell, 1998).

Pathological cystic follicles are different from atretic follicles (Brambell, 1956). A cystic follicle is a large (more than 2.5 cm in diameter), persistent, spherical structure on one of the ovaries. The more common type is a follicular cyst, found when the follicle fails to rupture but continues to grow (Peter, 2004). Less common are luteal cysts where the corpus luteum fails to regress and continues to produce progesterone, blocking further follicle development (Garverick, 1997).



Fig. 1: Diagram of the mammalian ovary illustrating the development and regression of follicles and corpora lutea. (a) Primordial follicle, (b) primary follicle, (c) secondary follicle, (d) tertiary follicle, (e) Graafian follicle, (f) cystic and (g) obliterative atretic follicles, (h) corpus hemorrhagicum, (i) corpus luteum, (j) corpus albicans, (k) rete ovarii, (l) ovarian blood vessels, (m) deep stroma, (n) superficial stroma, (o) tunica albuginea, (p) surface epithelium.

The bovine estrous cycle

Physiological and endocrinological characteristics

Domestic ruminants are polyoestrous animals. In normal heifers the first oestrus starts at the age of 8 to 12 months. Each estrous cycle takes 20 till 21 days, but cycle length may vary between 15 and 25 days (Sirois and Fortune, 1988; Ruckebusch et al., 1991; De Rensis and Peters, 1999). The reproductive cycle of the normal cycling cow is coordinated by hormones produced by the hypothalamus, pituitary gland and ovary. Gonadotrophic releasing hormone, secreted by the hypothalamus, stimulates the adenohypophysis to secrete two gonadotrophic hormones, namely follicle stimulating hormone (FSH) and luteinising hormone (LH). Both gonadotrophins circulate into the blood and regulate the reproductive cycle through a series of feedback mechanisms acting on the hypothalamus and pituitary gland (Peters and Lamming,

1983). In the ovary the regulation of follicular development is a complex process and involves interactions between the gonadotrophins FSH and LH and intraovarian regulators such as steroid hormones, growth factors and cytokines (Asselin et al., 2000).

In cattle, the initial stages of folliculogenesis (from primordial to secondary follicles) occur independent of gonadotrophin stimulation (Crowe et al., 2001; Gong, 2002). From the secondary follicular stage onwards, follicular development is a FSH dependent process as FSH is responsible for survival of the follicles (Gougeon, 1996). A group of secondary follicles are selected to continue growth to tertiary follicles by increasing concentrations of FSH. FSH stimulation causes expression of aromatase in the granulosa cells of the follicles, thus enabling the granulosa cells to convert androgens produced by the theca cells into estrogens (Richards, 1994). The circulating estrogens cause an inhibition of gonadotrophin release. One follicle of a selected group of tertiary follicles, called the dominant follicle, has an aromatase activity much more sensitive to FSH than the other smaller follicles. As a consequence, this dominant follicle is still capable of producing large amounts of estrogens even when the concentration of circulating gonadotrophins decreases (Gougeon, 1996). Concomitantly, increased concentrations of local growth factors and an elaborate perifollicular vasculature contribute to a positive selection of the dominant follicle, thus ensuring its final growth to a Graafian follicle which can ovulate (Coppola et al., 2005).

Progesterone plays also a major role in the selection of the dominant follicle. If circulating progesterone levels remain high after the dominant follicle has attained its maximum size, then all follicles of that particular wave including the dominant follicle undergo atresia (Taylor et al., 1993; Yang and Rajamahendran, 2000a). However, if circulating progesterone levels decrease during the growth phase of the dominant follicle, then this dominant follicle will ovulate while the other follicles of the cohort become atretic. Although the role of steroid hormones is of great importance, the exact mechanism that initiates the selection of the Graafian follicle and the regression of the non-ovulatory dominant follicle is not yet fully elucidated.

When a certain threshold of estrogens in the blood is reached, the negative feedback on gonadotrophin production and secretion is changed into a positive feedback. This results in peak levels of LH in the blood, triggering the ovulation. LH binds to LH receptors, which in secondary follicles are only localized on the theca cells, while in tertiary follicles they are also

present on the granulosa cells (Shima et al., 1987; Yamoto et al., 1992). At ovulation the follicle ruptures and develops into a corpus hemorrhagicum. In response to the binding of pituitary LH to receptors on cells of the ruptured follicular wall, the granulosa cells are converted in lutein cells producing high levels of progesterone (Ruckebusch et al., 1991; Dellmann and Eurell, 1998).

Follicle development in the bovine ovary (Fig. 2)

Primordial follicles undergo initial recruitment to enter the growing pool of primary follicles. Thereafter, the primary follicles continue their growth to the secondary stage and develop further into tertiary follicles (antral stage). The time required between the initial recruitment of a primordial follicle and its growth to the antral stage is about 180 days in cattle (Lussier et al., 1987). In the bovine ovary the growth of antral follicles occurs in a wave-like pattern. Each wave is characterized by the recruitment of a group of three to six follicles and the selection and dominance of a single dominant follicle (D) while the remaining recruited follicles regress (Savio et al., 1988). The number of growth waves during the estrous cycle is determined by the length of the luteal phase. Usually, there are two or three waves of follicular development during each bovine estrous cycle (Celik et al., 2005), although cycles with four waves or even only one wave exist (De Rensis and Peters, 1999). Fig. 2 illustrates the selection of a dominant follicle in a three waves growth pattern. The first wave begins on the day of ovulation. During the next few days, one of the follicles develops to a dominant stage, while the others become subordinate. The dominant follicle of the first wave reaches its largest size on day 5 or 6 of the estrous cycle. This dominant follicle maintains its morphological and functional dominance until around day 7, subsequently becomes atretic and begins to regress between days 8 and 9, to be replaced by a second wave of follicular growth. The second wave emerges at about 9 days and is followed by another wave at 16 days postovulation. The dominant follicle of the third wave becomes the ovulatory follicle (OV, Graafian follicle) (Ginther et al., 1996; De Rensis and Peters, 1999; Mapletoft et al., 2002).



Fig. 2: Schematic presentation of follicle recruitment, selection and dominance in the bovine ovary during the estrous cycle (after McGee and Hsueh, 2000; Aerts and Bols, 2004).

Morphological characteristics

In this study 23 cows were used. The stage of the estrous cycle of each cow was determined at the moment of slaughter on the basis of the morphological appearance of the ovarian follicles and corpora lutea and the presence of uterine cervical mucus (Ireland et al., 1980; Arthur et al., 1983). Additionally, the plasma progesterone concentration was determined for each individual animal and compared with the findings described by Sartori et al. (2004). The ovarian morphological characteristics and the corresponding plasma progesterone levels were required to classify the animals into 5 groups, viz., oestrus, metoestrus, early dioestrus, late dioestrus and procestrus (Fig. 3, Table 1).

Cycle stage	Day	Plasma progesterone concentration ng/ml	Morphologic characteristics
Oestrus	D0	< 0.30	Clear cervical mucus and presence of a mature
			follicle.
			Regressing corpora lutea.
Metoestrus	D1-3	0.20 - 0.40	Collapsed remnants of the ovulated follicle,
			corpus hemorrhagicum.
Early dioestrus	D4-8	0.70 - 2.65	Ovarian epithelium covering the rupture point
			of the ovulated follicle to form the apex of a
			new corpus luteum.
Late dioestrus	D9-16	2.90 - 4.70	Fully developed red or brown corpus luteum
			with vasculature visible around its periphery.
Prooestrus	D17-21	1.50 - 2.20	Brightly orange or yellow corpus luteum.
			Appearance of a mature follicle.

 Table 1: Classification of cows in five different cycle stages with progesterone levels and major ovarian features (after Ireland et al., 1980; Sartori et al., 2004).

Oestrus is the period (16 to 20 hours) of sexual receptiveness. The day of oestrus is considered as 'day 0' of the estrous cycle. About 30 hours after the onset of oestrus, ovulation occurs, preceded by a surge of LH. The estrogen levels decline at the end of oestrus. Metoestrus is the period (2 to 3 days) following oestrus, characterised by initial progesterone secretion. Early dioestrus is the period (about 5 days) of corpus luteum development and increasing progesterone levels. Late dioestrus is the phase (about 8 days) of the active corpus luteum when the progesterone level reaches its maximum. Prooestrus is the period (about 4 days) of regression of the corpus luteum and maturation of a new follicle (Arthur et al., 1983; Dellmann and Eurell, 1998) (Fig. 3).



Fig. 3: Schematic presentation of the hormonal changes that characterize the estrous cycle of the normal cycling cow. (P) Prooestrus, (O) oestrus, (M) metoestrus, (ED) early dioestrus, (LD) late dioestrus, (a) serum progesterone level, (b) serum estrogen level, (c) serum luteinizing hormone level (values from McDonald and Pineda; 1989).

The estrogen receptor (ER) and the progesterone receptor (PR)

The steroid hormone receptor superfamily

The steroid hormone receptor superfamily consists of a large number of proteins, originating from a common ancestral gene and therefore considered a gene superfamily (Tsai and O'Malley, 1994). This family includes receptors for the steroid hormones (estrogen, progesterone, glucocorticoid, mineralocorticoid and androgen), receptors for non-steroids, and several receptors for unknown ligands (orphan receptors). The estrogen receptor and the progesterone receptor are described in more detail hereafter.

The steroid hormone action (Fig. 4)

The concentration of the sex steroid hormone receptors in the various body tissues including the ovary is modulated by circulating and local amounts of sex steroid hormones (Seidman et al., 1997). Therefore sex steroid hormone receptors in the ovary are valuable mirrors of the action and concentration of sex steroid hormones.



Fig. 4: Simplified model of steroid hormone action (after Tsai and O'Malley, 1994).(1) Cell membrane, (2) cytoplasm, (3) nuclear envelope, (4) nucleus.

 \blacktriangleright sex steroid hormone, inactive receptor, \frown activated receptor.

Sex steroid hormones are hydrophobic molecules that pass through the cell membranes and interact with their specific receptors that are localized in the cell nucleus or cytoplasm in an inactive form (King and Greene, 1984; Perrot-Applanat et al., 1985). After binding of the hormone on its receptor, the latter undergoes conformational changes. The activated receptor can bind effectively to a hormone receptor element (HRE), i.e., a particular DNA sequence which is located in the region of hormone responsive genes. The DNA-bound receptor ligand complex is then capable of either activating or repressing target gene transcription (Beekman et al., 1993). The resulting mRNA molecules will be transferred to the cell cytoplasm where they are translated into specific proteins that will induce the cellular response to hormone action.

The steroid receptors ER and PR are transcription factors composed of six domains (A-F) responsible for specific functions. The N-terminal A/B domain contains an activation function (AF-1) that contributes to transcriptional activity. The C domain is the DNA-binding domain and contains two zinc finger motifs. This domain contributes to receptor dimerization, binding of the liganded receptor to its target genes (= response elements), and finally to the transactivation function of the receptor. Domain D is the hinge region that allows conformational changes during activation of the receptor. This domain contains conserved amino acids which play a role in binding to heat shock proteins, nuclear localization and stabilization of the DNA binding. The E domain is the ligand-binding domain and contains the ligand binding site and the transactivation functions AF-2 and AF-2a (Peters and Khan, 1999). The function of domain F is still unclear, but it is probably involved in modulating transcriptional activation (Pavao and Traish, 2001). Ligand binding may influence the stability of the receptor. In particular, it has been shown that in the absence of ligand, the half-life of ER α is about 4-5 h, whereas estradiol binding accelerates receptor degradation, reducing its half-life to 3-4 h. (Eckert et al., 1984; Nardulli and Katzenellenbogen, 1986).



Fig. 5: Model of the receptor protein with several regions responsible for specific functions. A/B: transcriptional activation domain; C: DNA binding and dimerization domain; D: nuclear localization domain; E: ligand binding, dimerization and transcriptional domain; F: unknown function (Eckert et al., 1984).

Estrogen receptor subtypes

Two estrogen receptor subtypes are known, namely the ER α which was first described by Walter et al. in 1986, and the ER β which was described by Kuiper in 1996. Both receptors are encoded by a different gene, namely the ER α and the ER β gene. In addition to the presence of the two receptor subtypes, the expression of several isoforms from the ER α and ER β genes greatly enlarges the diversity of ER. These isoforms are formed by alternative splicing leading to deletions of certain exons (Walther et al., 1999; Fang et al., 2003). Recently, a third subtype of estrogen receptor, the ER γ , has been detected in the teleost *Micropogonias undulatus* (Hawkins et al., 2000), in mouse (Lorke et al., 2000) and in humans (Heard et al., 2000).

Estrogen receptor a isoforms

Several ER α isoforms have been identified which can be expressed in various species and tissues. In the MCF6 human breast cancer cell line two isoforms of the ER α gene are expressed, namely the ER α 66 isoform, 66 kDa in size, and the ER α 46 isoform, 46 kDa in size (Cullen et al., 2000; Flouriot et al., 2000; Métivier et al., 2004). Additionally, the expression of ER α 46 was reported in osteoblasts (Denger et al, 2001) and in endothelial cells (Li et al., 1998). In chicken two ER α isoforms were identified, the ER α 66 isoform and the ER α 61 isoform, 61 kDa in size. In cockerels and hens, both variants are expressed in the adenohypophysis and basal hypothalamus (Griffin et al., 2001). In rainbow trout two ER α isoforms are found, namely the ER α 1 and ER α 2. Comparison of ER α 1 with ER α 2 reveals a high similarity in the conserved DNA-binding domain (91%) and ligand-binding domain (89%) (Bouma and Nagler, 2001).

Estrogen receptor β isoforms

A large number of ER β isoforms with variable tissue expression has been reported. Several isoforms are found in the human testis. These include the ER β 2 isoform which is also known as ER β cx (Ogawa et al., 1998), the ER β 3, ER β 4 and ER β 5 isoforms (Moore et al., 1998), the ER β 1 δ 5 isoform which lacks exon 5 (Inoue et al., 2000), and the ER β M isoform containing exons 5-8 and exon M (Shoda et al., 2002). Some of these isoforms namely ER β 2 and ER β 2 δ 5 have been associated with human breast cancer (Skrzyczak et al., 2003; Davies et al., 2004). Additionally, 3 ER β isoforms (δ 11, δ 21 and δ 31) were demonstrated in the bovine reproductive organs (Walther et al., 1999).

Progesterone receptor isoforms

All isoforms of PR identified so far are derived from a single gene as a consequence of alternate initiation of transcription by distinct promoters (Kastner et al., 1990b; Taylor et al., 2006).

In humans, macaques, rats, mice and chickens two isoforms of PR have been described, namely PRA, 94 kDa in size, and PRB, 118 kDa in size (Christensen et al., 1991; Ilenchuck and Walters, 1987; Hovland et al., 1998; Wei et al., 1997; Bethea and Widmann, 1998; Pieber et al., 2001; Lessey et al., 1983; Aupperlee et al., 2005; Kastner et al., 1990a; Gonzalez-Moran and Camacho-Arroyo, 2001). PRA is the predominant isoform in humans and rodents (Vegeto et al., 1993; Graham and Clarke, 1997) and is widely expressed in the macaque reproductive system (Bethea and Widmann, 1998), in the human uterus (Mote et al., 1999; Mangal et al., 1997) and in human breast cancers (Graham et al., 1995). A third isoform, called PRC and 60 kDa in size, was detected in the progestin responsive human breast cancer cell line T47D (Wei et al., 1996). This isoform is inactive by itself, but it may inhibit the activity of both PRA and PRB in the presence of progesterone (Wei and Miner, 1994). Recently, two new isoforms PRS and PRM have been identified and partially characterised in human reproductive tissues (Taylor et al., 2006).

Cell dynamics in the bovine ovary

Apoptosis in follicles and corpora lutea of the bovine ovary

Atresia of ovarian follicles is an important process, as it accounts for the loss of 99% of the follicles in the mammalian ovary (Byskov, 1978; Greenwald and Terranova, 1988; Mariana et al., 1991). This follicular atresia is mediated by a highly organized type of cell death called apoptosis or programmed cell death (Hsueh et al., 1994; Billig et al., 1996; Kaipia and Hsueh, 1997; Driancourt, 2001).

During follicular development, endocrine factors such as FSH and LH, and paracrine factors including growth factors, activin and interleukin 1β as well as estrogens activate different intracellular pathways to rescue the follicles from apoptotic demise. In contrast,

tumor necrosis factor α , androgens, and free radicals function as atretogenic factors (Kaipia and Hsueh, 1997). Additionally, specific genes have been identified that encode proteins responsible for the initiation, progression and completion of cell death, particularly a large family of B-cell lymphomal leukemia-2 (Bcl-2)-related proteins (Tilly, 2003). All these diverse signals probably converge on specific intracellular pathways to regulate apoptosis in the ovarian follicles. Because of these multiple signals, it is still not completely understood which are the critical factors that discriminate between follicles destined for elimination by apoptosis and follicles that will continue to develop to reach the final stage of a preovulatory follicle.

During the growth and development of follicles, atresia occurs at every stage of follicular development (Gougeon, 1996; Yang and Rajamahendran, 2000b). In humans, quiescent follicles are the most susceptible to apoptosis (Depalo et al., 2003). In this context, various authors have investigated gonadotrophin control of primordial and primary follicles (Gougeon, 1996; McGee and Hsueh, 2000) and some reports show that in quiescent follicles FSH does not act as a survival factor but that various locally produced growth factors are involved in the cellular control mechanisms (Markstrom et al., 2002). The most common signs of atresia in primordial, primary and secondary bovine follicles are firstly shrinking of the oocyte and degeneration of the follicle cells (Yang and Rajamahendran, 2000b). The follicle will subsequently be resorbed without leaving a trace (Bloom and Fawcett, 1975). In contrast, atresia of tertiary follicles occurs differently and is first reflected by apoptosis in the inner granulosa cell layer and reduction in granulosa cell function, especially the loss of aromatase activity (Amsterdam et al., 2003; Tajima et al., 2002). In a further stage apoptosis leads to almost total destruction of the granulosa cell layer and theca cells (Isobe and Yoshimura, 2000). Studies in bovine Graafian follicles suggest that the local and chronological progress of apoptosis from the granulosa layer towards the embedded oocyte is triggered by a threshold quantity of cells in the theca and granulosa layers and in the cumulus oophorus (Bendell et al., 1988; Zeuner et al., 2003).

Apoptosis has also been implicated as the mechanism underlying structural regression of the corpora lutea. The onset of apoptosis in the bovine corpus luteum is not observed until progesterone production has declined (Juengel et al., 1993). This suggests that progesterone may prevent apoptosis (Okuda et al., 2004). However, it is not known whether the decline in progesterone production and its auto- or paracrine actions within the corpus luteum influence

receptor expression. Therefore, it is of interest to elucidate the role of progesterone and its receptors in the corpus luteum.

Cell proliferation in follicles and corpora lutea of the bovine ovary

During the bovine estrous cycle, the ovary is subjected to extensive tissue remodelling that consists of sequential phases of cell proliferation and cell death in the different ovarian structures (Fortune et al., 1994; Gaytan et al., 1996). The growth of ovarian follicles, ovulation and the formation of the corpus luteum are basic events in the proliferation process. This process is regulated by specific responses to gonadotrophins, steroid hormones and growth factors (Pedersen, 1970; Richards, 1980; Hirshfield, 1991).

In general, follicle growth consists of two distinct consecutive phases. The first phase is characterized by the transformation of follicle cells from a flattened shape to cuboidal cells, and the second phase is characterized by proliferation of the follicle cells. Both phases are regulated by locally produced inhibitory and stimulatory factors (Braw-Tal, 2002). In primordial follicles, the oocyte is surrounded by a single layer of non-dividing follicle cells. Primordial follicles leave this quiescent state and initiate a phase of slow growth. Studies in rat showed that the transition from a primordial follicle to a follicle with a single layer of columnar follicle cells might take several weeks (Hirshfield, 1991). In the secondary and tertiary follicles, the follicle/granulosa cells acquire enhanced responsiveness to FSH and start producing estradiol (Richards, 1975, 1980). Exposure to FSH and estradiol triggers a rapid burst of proliferation that results in the formation of large tertiary follicles (Rao et al., 1978) characterized by the presence of both granulosa cells and a surrounding layer of theca cells (Uilenbroek and Richards, 1979). The LH surge triggers dramatic changes in both follicular structure and function and terminates follicular growth (Rao et al., 1978; Hirshfield, 1991). However, the mechanism by which the growth and differentiation of theca cells in bovine ovarian follicles are controlled is not yet elucidated. Some reports suggest the importance of granulosa-theca interactions (Kotsuji et al., 1990) while others indicate that specific growth factors could promote cell proliferation (Asselin et al., 2000). The follicle ruptures during ovulation and the granulosa cells luteinize to form the corpus luteum (Lobel and Levy, 1968; Meidan and Girsh, 1997; Meidan et al., 1999). The lutein cells undergo intense proliferation during the early stages of luteal development, but during functional luteal regression the

cellular proliferation activity is much lower (Lei et al., 1991). Finally, the lutein cells undergo terminal differentiation and cease to divide (Richards, 1980).

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AIMS OF THE STUDY

During the bovine estrous cycle, the different ovarian structures are subjected to extensive tissue remodelling that consists of sequential phases of cell proliferation and cell death. These processes can be influenced by locally produced estrogens and progesterone. Both hormones exert their functions by binding to specific receptors, the estrogen receptors (ER α and ER β) and the progesterone receptors (PRA and PRB).

Data concerning related changes in the ER and PR expression in the bovine ovary are scarce. This study was undertaken to obtain a better insight into the role of estrogen receptors and progesterone receptors in the different cell types of the bovine ovary and to verify their relation to cell proliferation and apoptotis throughout the estrous cycle.

The specific aims of the present thesis are:

- to describe the morphological localization of estrogen receptors and progesterone receptors in the different cell types of both ovaries in normal cycling cows
- to evaluate whether quantitative differences in the presence of these receptors exist during the different stages of the estrous cycle
- to examine the presence of apoptosis in the bovine ovarian cell types during the different cycle stages using three techniques, i.e., the detection of activated caspase-3, *in situ* DNA end labelling (TUNEL) and a DNA fluorescent staining (DAPI)
- to verify cell proliferation in the bovine ovarian cell types during the various estrous cycle stages by means of immunohistochemical detection of Ki-67 expression
- to study putative differences between the scores in active and inactive ovaries for each parameter, i.e., estrogen and progesterone receptor expression, apoptosis and cell proliferation
- to determine the correlation of each parameter with the plasma progesterone concentration in the various ovarian cell types

CHAPTER 1

SEX STEROID HORMONE RECEPTORS

IN THE BOVINE OVARY

1.1 Localization of estrogen receptors within various bovine ovarian cell types at different stages of the estrous cycle

1.1.1 Cell-specific localization of estrogen receptor beta (ESR2) mRNA within various bovine ovarian cell types using in situ hybridisation

Modified from:

CELL-SPECIFIC LOCALIZATION OF ESTROGEN RECEPTOR BETA (ESR2) mRNA WITHIN VARIOUS BOVINE OVARIAN CELL TYPES USING IN SITU HYBRIDISATION D'Haeseleer M, Van Poucke M, Van den Broeck W Anatomia Histologia Embryologia 2005; <u>34</u>: 265-272

BOS TAURUS ESTROGEN RECEPTOR BETA, PARTIAL SEQUENCE D'Haeseleer M, Van Poucke M, Van den Broeck W NCBI 2005: GenBank Database Accession AY690422

Summary

The localization of estrogen receptor beta (ESR2) mRNA, in this article denominated as (ERβ) mRNA, was examined using *in situ* hybridization in the ovaries of randomly selected cows, irrespective of the cycle stage of the animals. A 602 base pairs fragment of ERB mRNA was cloned, sequenced and digoxigenin (DIG)-labelled. Semi-quantitative evaluation showed that the scores for ER^β mRNA were moderate to high in the follicle cells of both primordial and primary follicles, but lower in granulosa cells of secondary follicles. In vital tertiary follicles, the total ER β mRNA expression was low but varied between the different animals. In both obliterative and cystic atretic follicles, high to moderate ER^β mRNA scores were noticed in the granulosa cells. The stroma cells surrounding primordial and primary follicles and the theca cells of secondary follicles showed moderate ERB mRNA levels, whereas the ERβ mRNA score in theca interna and theca externa cells of vital tertiary follicles was distinctly higher. In the theca cells of atretic follicles the score was even higher. Cells of corpora hemorrhagica and corpora lutea had moderate ERB mRNA scores, while higher scores were seen in cells of corpora albicantia. Cells of the surface epithelium had a moderate score for ER β mRNA, whereas cells of the tunica albuginea and deep stroma showed high ERβ mRNA scores. The present findings have clearly established a cell-specific localization of ER β mRNA in several cell types in the bovine ovary.

Introduction

It has been well established that the gonadal steroid estrogen hormones exert a wide variety of effects on growth, differentiation and functioning of many female and male reproductive organs including the ovary, uterine tube/oviduct, uterus, vagina and mammary gland, and the testis, epididymis and prostate (Clark et al., 1992). To exert their functions, estrogens are retained with high affinity and specificity in target cells by binding to specific intranuclear binding proteins, the estrogen receptors (ER) (Gronemeyer and Laudet, 1995). These receptors belong to a large group of transcription factors, the so-called nuclear receptor superfamily (Tsai and O'Malley, 1994), and are encoded by two different genes resulting in two receptor subtypes, namely estrogen receptor alpha (ER α) and ER β . After ligand binding and dimerization, both receptors function as signal transducers and transcription factors to modulate the expression of specific target genes by binding to target sequences called "response elements" (Enmark and Gustafsson, 1999). In addition to acting as homodimers (ER α /ER α or ER β /ER β), ER α and ER β are also able to form heterodimers (ER α /ER β) (Tremblay et al., 1997).

The cloning and initial characterization of ER β was first described in the rat prostate (Kuiper et al., 1996) and subsequently orthologues have been found in humans (Mosselman et al., 1996), mouse (Tremblay et al., 1997), pig (Kowalski et al., 2002; LaVoie et al., 2002), sheep (Cardenas et al., 2001), monkey (Wu et al., 2000), goldfish (Tchoudakova et al., 1999) and cattle (Rosenfeld et al., 1999; Walther et al., 1999). Within the different species, ER β shows the highest expression in prostate, ovary, lung, bladder, brain and epididymis (Kuiper et al., 1996; Li et al., 1997). In addition, ER β plays an important role in bone maintenance (Turner et al., 1994), in the cardiovascular system (Farhat et al., 1996) and in some breast cancers (Dotzlaw et al., 1997). ER β also mediates activities in the central nervous system (McCarthy and Pfaus, 1996). In the female reproductive tissues, ER β has been found in the ovary, uterus and mammary gland. In the human ovary, the receptor is located in stroma cells of the cortex as well as in the granulosa cells (Byers et al., 1997). Therefore, ER β might play an important role in the regulation of follicular growth and oocyte development (Enmark and Gustafsson, 1999).

The presence of ER β in the bovine ovary has already been demonstrated by *in situ* hybridization (Rosenfeld et al., 1999), but the cell specific localization of ER β remains

unclear. Therefore, the purpose of this study was to localize $ER\beta$ in the different cell types of the bovine ovary.

Materials and methods

Animals and tissue sampling

Three adult cows, without symptoms of any reproductive pathology, were sampled at the moment of slaughter at a local slaughterhouse. Immediately after exsanguination both ovaries were collected. For *in situ* hybridization one part of the ovaries was fixed in a RNase free phosphate buffered saline (PBS) solution containing 4% paraformaldehyde at 4°C for 8 hours, dehydrated in graded methanol concentrations, cleared in xylene and embedded in paraffin wax. Five µm thin sections were cut and mounted on poly-L-lysine-coated (Sigma Diagnostics Inc., St. Louis, MO, USA) SuperFrost[®]Plus-slides (Menzel-Gläser, Braunschweig, Germany) and dried overnight at 37°C. Another part of the ovaries was frozen in liquid nitrogen and used for the generation of a probe for *in situ* hybridization.

Probe design

Total RNA isolation and reverse transcription

Total RNA extraction from bovine ovarian tissue was performed using Trizol reagent (Invitrogen Life Technologies, Merelbeke, Belgium) followed by DNA digestion using DNase 1. Before isolation, 10 pg of rabbit globin mRNA (Invitrogen Life Technologies) was added to each sample as an external standard. RNA isolation efficiency was checked by polymerase chain reaction (PCR) with primers for rabbit globin sense and anti-sense. Reverse transcription was done as described in the protocol from the Reverse Transcription kit (Invitrogen Life Technologies). The bovine glyceraldehyde-3-phosphate dehydrogenase (GAPDH) primers were used as an internal control for the complementary DNA (Yuan *et al.*, 2003).

PCR and verification

In order to amplify a specific fragment of 602 base pairs, two primers (left primer: 5'-CTGCGGTCAATCCATCCTAC-3' and right primer: 5'-TCCGAGAGCCACACTTCAC-3') were designed based on the bovine ER β sequence (Acc No.Y18017) by using the Primer 3 program (Rozen and Skaletsky, 1998). PCR was carried out on a T3 thermocycler (Biometra, Göttingen, Germany) in 10 µl containing 25 ng complementary DNA, reaction buffer, 3 mM MgCl₂ 200 mM of each desoxynucleoside 5'-triphosphate, 200 mM of each primer and 0,5 units Taq Platinum polymerase (Invitrogen Life Technologies). The thermal cycling profile was 10 min at 94°C for denaturation, followed by 40 cycles of 30 sec at 94°C, 30 sec at 58°C, 1 min at 72°C and a final elongation step of 10 min at 72°C. The 602 base pairs PCR product was visualised on a 0.8% agarose gel. cloned in the pCR[®]2.1 Vector (Invitrogen Life Technologies) and sequenced for verification using an ALFexpressTM Sequencing System (Pharmacia, Uppsala, Sweden). In order to generate ER_β-specific RNA probes, the same fragment was subcloned into the pGEM[®]-T easy vector (Promega, USA) according to standard procedures (Sambrook et al., 1989). Subsequently the ligation product was transformed into bacteria, viz., Subcloning Efficiency DH5 α^{TM} Competent Cells (Invitrogen Life Technologies) and recombinant clones were identified by white: blue colour screening chromogenic substrate X-gal (5-bromo-4-chloro-3-indol-β-D-galactoside) using the (Sambrook et al., 1989). After culturing and harvesting the bacteria, plasmid DNA was isolated using the HiSpeed TM Plasmid Purification kit (Oiagen, Westburg, The Netherlands) and the orientation of the insert was determined by restriction digest analysis using the enzymes PstI, SpeI and SalI (Fig. 1).



Fig. 1: Analysis of DNA by 0.8% agarose gel electrophoresis with ethidium bromide staining; lane 1: 602-bp DNA fragment of bovine ER β following RT-PCR using ER β -specific primers; lane 2: 1 kb λ -marker; lanes 3 and 4: restriction enzyme digest of the cloned DNA fragment using *Spe*I (lane 3) and *Pst*I (lane 4) to ascertain the identity; lane 5: ER β DNA fragment cut out of the plasmid using *EcoR*I restriction enzyme digestion.

Probe generation

Plasmid DNA was linearised with restriction enzymes *Sac*II and *Sac*I to allow *in vitro* runoff synthesis of both sense- and antisense-oriented RNA probes respectively. After phenolchloroform extraction and ethanol precipitation, DIG-labelled RNA probes were generated by transcription using the SP6-T7 DIG RNA Labelling Kit (Roche Molecular Biochemicals, USA) and purified with Mini quick Spin RNA columns (Roche Molecular Biochemicals, USA) (Fig. 2). The yield of the probes was estimated by DIG quantification on DIG control test strips (Roche Diagnostics Corporation, Indianapolis, USA). All experiments were performed according to the manufacturers' instructions unless stated otherwise.



Fig. 2: Analysis of RNA transcripts by 1% agarose gel electrophoresis with ethidium bromide staining; lane 1: λ -marker; lanes 2 and 3: internal controls; lane 4: antisense ER β mRNA; lane 5: sense ER β mRNA.

-602 bp

Histological analysis

In situ hybridization

ER β mRNA was localized in ovarian tissue sections by *in situ* hybridization analysis. Deparaffinized and rehydrated sections were rinsed in double concentrated SSPE (0.1 M NaCl, 10 mM NaH₂PO₄ pH 7.4, 1 mM EDTA) and transferred into Tris-HCl (pH 7.6) for 5 min. Afterwards, slides were treated with 20 µg/ml Proteinase K (Roche Diagnostics, Vilvoorde, Belgium) in 0.05 M Tris-HCl (pH 7.4) at 37°C for 30 min to make the RNA more accessible to hybridization. Sections were fixed in 4% paraformaldehyde in PBS at room temperature for 10 min. The slides were rinsed in PBS and acetylated in 0.2 M HCl for 15 min at room temperature. Prehybridization was performed by incubating the slides with a hybridization buffer (Schwarzacher and Heslop-Harrison, 2000) in a humidified chamber at 42°C for 2 hours.

Hybridization was executed by incubating slides with the hybridization buffer containing the anti-sense probe (200 ng/ml). As negative controls, two sections were incubated either with hybridization buffer alone or with hybridization buffer containing the sense probe. The slides were incubated in a humidified chamber (50 % deionized formamide + 50 % diethyl pyrocarbonate water) at 42°C overnight. Posthybridization was carried out by incubating the slides for 30 min at 37°C in RNase buffer (1 M Tris, pH 8.0, 5 M NaCl, 0.5 M EDTA) containing RNase-A (50 μ g/ml, Sigma, Belgium). Afterwards the slides were rinsed in double concentrated SSPE for 15 min at 42°C, in SSPE for 30 min at 42°C and in SSPE for 15 min at room temperature.

Immunohistochemical detection

Sections were incubated with 1:70 v/v sheep anti-DIG-peroxidase anti-binding fragments (Roche Diagnostics, Vilvoorde, Belgium) for 1 hour at 37°C in a humidified chamber. The sections were subsequently stained by incubation with DAB substrate (DAKO, Prosan, Merelbeke, Belgium) for 10 minutes and counterstained with Mayer's hematoxylin for 5 sec for better visualization of the cellular architecture of ovarian cell types.

Microscopic analysis

In all ovarian sections the different ovarian cell types were analysed by light microscopic inspection. Staining scores were awarded in different follicles, corpora lutea, corpora

albicantia, vascular endothelial cells, rete ovarii, deep and superficial stroma, tunica albuginea and surface epithelium. All follicles were examined and classified into primordial, primary, secondary and tertiary follicles according to the criteria listed in Nomina Histologica (1994). In tertiary follicles the inner layers of granulosa cells facing the antral side were quantified separately from granulosa layers of the basal side because they showed different staining patterns. Atretic tertiary follicles were classified as obliterative or cystic (Dellmann and Eurell, 1998).

For the semi-quantitative evaluation of ER β mRNA the number of ER β mRNA containing cells, identified by a brown staining in the cytoplasm, was counted. For all cell types, at least 100 cells were counted in digital micrographs taken randomly at magnification 600 x (Olympus BX61, Olympus Belgium, Aartselaar, Belgium). If too many cells were present in one micrograph, 4 grids of 2500 μ m² were placed randomly onto the micrograph and cells were counted in these areas.

Staining intensity scores ranged from 0 (no staining), 1 (weak staining), 2 (moderate staining) to 3 (strong staining). ER β mRNA presence was evaluated using a score (SER β mRNA) which was calculated according to the formula:

SER β mRNA = 0 n₀ + 1 n₁+ 2 n₂ + 3 n₃

with n_0 , n_1 , n_2 and n_3 are the percentage of cells exhibiting a staining intensity 0, 1, 2 and 3, respectively (Boos et al., 1996).

Results

Production of ER β mRNA probes for in situ hybridization

After sequence comparison of bovine ER α (Acc. No. 538775) with ER β (Acc. No. Y18017), an ER β -specific region of 602 base pairs was selected from exon 1 (5'-UTR) to exon 4. The amplified fragment (Acc. No. AY 690422) showed 99% sequence identity with the sequence on which the primers were based because of 4 different nucleotides at position 19, 26, 57 and 202 (Table 1). With this fragment as a template, both sense- and antisense-oriented RNA probes were synthesized with a yield of 200 ng/ml.

	ERβ sequence			
	Acc. No. Y18017		Acc. No. AY690422	
	Nucleotides	Amino acid	Nucleotides	Amino acid
Codon 7 (nucleotides 19-21)	GAT	Asp	AAT	Asn
Codon 9 (nucleotides 25-27)	TCA	Ser	TGA	STOP
Codon 19 (nucleotides 55-57)	AGT	Ser	AGC	Ser
Codon 68 (nucleotides 202-204)	CTG	Leu	TTG	Leu

Table 1: Sequence comparison of bovine ERβ Acc. No. Y18017 with Acc. No. AY 690422.

Localization of ER β mRNA in the bovine ovary

The *in situ* hybridization technique confirmed the presence of a specific and intense signal in the different ovarian tissues when using the DIG-labelled RNA anti-sense probe (Fig. 3a) while no signal was visible using the DIG- labelled RNA sense probe (Fig. 3b).

The presence of ER β mRNA, identified by brown cytoplasmic staining, was found in all animals in most follicular stages, in the corpora lutea and corpora albicantia, in the superficial and deep stroma, in vascular endothelial cells, in the rete ovarii, in the tunica albuginea and in the surface epithelium.



Fig. 3: Micrograph of a tertiary follicle following *in situ* hybridization using the ER β -specific antisense probe (**Fig. 3a**) and sense probe (negative control) (**Fig. 3b**). Bar = 100 μ m.

Cell-specific variation of ER β mRNA expression

In situ localization results indicated that the intensity of specific staining and the proportion of positive cells varied between the different cell types (Fig. 4 and 5). The scores for ER β mRNA were moderate to high in the follicle cells of both primordial and primary follicles, but clearly higher than in follicle cells of secondary follicles. In vital tertiary follicles, the granulosa cells of both the cumulus oophorus and the granulosa layers showed a higher expression of ER β mRNA compared with that in follicle cells of secondary follicles. However, this expression in vital tertiary follicles was only seen in one of the 3 cows, whereas no expression of ER β mRNA could be detected in this cell type of the other animals. In both obliterative and cystic atretic follicles, high to moderate ER β mRNA scores were noticed in the granulosa cells.

A moderate SER β mRNA was observed in superficial stroma cells surrounding the primordial and primary follicles. In stroma cells around secondary follicles (theca follicularis cells) this score was similar, whereas the score in theca interna and externa cells of vital tertiary follicles was manifestly higher. A similar high score was found in the theca interna and theca externa layers of cystic atretic follicles. In both theca layers of obliterative follicles SER β mRNA were higher than in any other follicular cell type. In obliterative follicles, the membrane of Slavjansky was also strongly positive.

A corpus hemorrhagicum was only present in one animal, with moderate scores for ER β mRNA in the granulosa cells and higher scores in both theca layers. Corpora lutea could only be investigated in two animals, and within these structures, moderate ER β mRNA scores were present in both large and small lutein cells, vascular endothelial cells, perivascular stroma cells and peripheral stroma cells (comparable with theca externa cells of tertiary follicles). High ER β mRNA levels were observed in corpora albicantia with similar high scores in internal and peripheral stroma cells.

Finally, the cells of the surface epithelium had a moderate SER β mRNA, whereas cells of the tunica albuginea, deep stroma, rete ovarii and vascular endothelial cells showed high ER β mRNA levels.



Fig. 4: Mean scores (\pm SEM) for ER β mRNA (SER β mRNA) in different groups of bovine ovarian cells; II = secondary follicle, vital III = vital tertiary follicle, obliterative III = obliterative attractic tertiary follicle, cystic III = cystic attractic tertiary follicle, CH = corpus hemorrhagicum, CL = corpus albicans; (*n*=1, 2, 3).



Fig. 5: Micrographs of different bovine ovarian cell types following *in situ* hybridization using the ER β -specific antisense probe; **Fig. 5a and 5b**: primordial (1), primary (2) and secondary (3) follicles with moderate to low scores for ER β mRNA (SER β mRNA) in the follicle / granulosa cells, and moderate to high SER β mRNA in stroma cells (4); **Fig. 5c**: wall of a vital tertiary follicle with low ER β mRNA scores in the granulosa cells (5) and high SER β mRNA in theca interna (6) and externa cells (7); **Fig. 5d**: wall of a cystic atretic

tertiary follicle with moderate to high SER β mRNA in apoptotic granulosa cells (5) and in theca interna (6) and externa cells (7); **Fig. 5e**: overview of an obliterative attretic tertiary follicle with high SER β mRNA in the collapsed theca layers (8) and the thickened basal membrane (9); **Fig. 5f**: overview of a corpus luteum with moderate to high SER β mRNA in the lutein cells (10); **Fig. 5g**: overview of a corpus albicans with moderate to high SER β mRNA in internal stroma cells (11); **Fig. 5h**: cells of the surface epithelium (12) with moderate SER β mRNA and tunica albuginea cells (13) with low SER β mRNA; **Fig. 5i**: cells of the deep stroma (14) with high SER β mRNA. Bar = 50 µm in Fig. 5a, 5b, 5c, 5d, 5h and 5i, 100 µm in Fig. 5e and 5g, and 500 µm in Fig. 5f.

Discussion

The primary aim of this study was to localize ER β mRNA in different cell types of the bovine ovary by *in situ* hybridization using DIG labelled bovine ER β mRNA probes. In comparison with the early published bovine ER β sequence different nucleotides were observed in the coding sequence. A G to A transition at position 19, results in an aspartic acid (Asp) to asparagin (Asn) substitution at amino acid 7. The difference between Asn and Asp is the neutral character of Asn (NH₂-group) and the acid character of Asp (OH-group). This missense mutation might have an impact on the protein structure, folding and related cell response to hormone actions. A C to G transversion at position 26 results in the incorporation of a STOP-codon at amino acid 9. Although, this nonsense mutation alters the translation, it doesn't influence our *in situ* hybridization experiments, because we are interested in the binding of the ovarian tissue mRNA with the mRNA probes. A T to C and a C to T transition at pos 57 and 202 respectively are silent mutations, resulting in the incorporation of the amino acids serine (Ser) and leucine (Leu) at pos 19 and 68.

Since this was a single-pass sequence, the question remains whether these 4 polymorphisms are time single nucleotides polymorphisms (SNPs) or PCR sequencing errors. If they are SNPs it should be worthwhile to investigate the nonsense and the missense mutations.

Localization of ER β mRNA and protein in ovarian cells of various species has already been reported as reviewed by Rosenfeld and others (2001). The present study revealed that the expression level of ER β mRNA varied between the different follicular types. Indeed, ER β mRNA levels were high in the follicle cells of primordial and primary follicles and were

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manifestly lower in the granulosa cells of secondary and tertiary follicles. Similar findings are described in the ewe, namely a high specific expression of ER β in the follicle/granulosa cell layer of small and developing follicles that declines with increasing follicle size (Jansen et al., 2001). In contrast, human (Pelletier and El-Alfy, 2000) and rat (Pelletier et al., 2000) granulosa cells of pre-antral and antral follicles showed a high expression of ERB. Rosenfeld and co-workers (Rosenfeld et al., 1999) found high ERB mRNA levels in granulosa cells of tertiary follicles of cattle, but they did not compare this expression with that in other follicular structures. Despite the differences in follicle-specific expression between different species, these and our findings support the hypothesis that estrogens are involved in folliculogenesis through interaction with ER β and not with ER α as the latter receptor was rarely if ever found in bovine ovarian follicles (Van den Broeck et al., 2002). Furthermore, the moderate to high expression of ER β mRNA in primordial and primary ovarian follicles of cattle indicates that $ER\beta$ is mainly involved in the earlier stages of development. Again, this is in contrast with studies in ERaß double knock-out (KO) mice, in which early follicular growth and development occurred normally but without leading to mature Graafian follicles, suggesting that ERα and ERβ may not be necessary for the earlier stages of folliculogenesis (Dupont et al., 2000).

Remarkably, a moderate to high ER β mRNA expression was present in the granulosa cell layer of obliterative and cystic atretic tertiary follicles. This atresia is due to apoptosis of the granulosa cells (Amsterdam et al., 1999) which in turn can be inhibited by estrogens and promoted by androgens (Billig et al., 1993). Studies in ER α KO mice have demonstrated that this inhibition is not conferred through ER α (Schomberg et al., 1999). In our study, high levels of ER β mRNA were present in the atretic follicles, so apparently ER β could not prevent the atresia. This can be due to the fact that other pro- and anti-apoptotic factors are present and that the final outcome is the result of a complex crosstalk among multiple intracellular pathways in which the protective capacity of estrogens is overruled. An alternative possibility is that the ER β demonstrated here is characterized by a lower affinity for estrogens or for estrogen response elements, so that no apoptotic inhibiting effect is exerted. This also means that there should be another yet unidentified estrogen receptor that can prevent apoptosis. A third estrogen receptor, ER γ , has been identified in fish (Hawkins et al., 2000) and there is some evidence that at least one additional estrogen receptor is present in the mouse ovary and uterus (Ghosh et al., 1999). Similarly, it might be possible that the bovine ovary also contain a third ER type that prevents follicular atresia. This hypothesis, although highly speculative, should be kept in mind and may form the basis for further experimental research.

Moderate ER β mRNA levels were noted in granulosa cells of a corpus hemorrhagicum present in one animal, and in small and large lutein cells of corpora lutea present in two animals. These observations are suggestive of a role of estrogens in the formation and maintenance of a corpus luteum. The specific role of ER β and ER α in this process in cattle is unclear, as moderate levels of ER α are also present in corpora lutea cells (Van den Broeck et al., 2002). Studies in mice have elucidated that in both ER α KO and ER β KO animals steroidogenically functional corpora lutea are formed (Krege et al., 1998; Rosenfeld et al., 2000). In contrast, ER $\alpha\beta$ double KO mice do not undergo luteinization and corpus luteum formation (Couse et al., 1999). These results indicate that estrogen is needed for corpus luteum formation and maintenance acting through ER α or ER β , and luteinization cannot take place when both receptors are lacking. A similar mechanism can occur in cattle, but remains to be proven.

Cells of both the tunica albuginea and deep stroma showed high ER β mRNA expression levels, which has never been described before. The exact role of ER β in these cell types remains to be clarified, but a putative function is that upon estrogen binding, these cells alter the stromal composition in order to allow the dynamical processes of follicle development, ovulation and corpus luteum formation. Further studies including cell culture experiments are needed to confirm this hypothesis. In stroma cells immediately surrounding primordial and primary follicles ER β mRNA expression was lower than in cells of the tunica albuginea and deep stroma, and in follicle cells of primordial and primary follicles. However, the theca layers of secondary, vital tertiary and atretic tertiary follicles showed a gradually increased ERβ mRNA expression which was always higher than in the granulosa cells of the corresponding follicle. These findings might indicate that for primordial and primary follicles, estrogens act via ER^β that are located within the follicle cells, whereas the indirect influence via the surrounding stroma cells is lower. In contrast, ER^β levels in secondary and tertiary follicles are decreased in the granulosa cells and increased in the theca cells, which is suggestive of an indirect effect of estrogens on the developing follicles via the theca cells. These stroma-epithelial interactions have already been described in earlier studies for progesterone (Hild-Petito et al., 1988), and theca cells may produce factors that promote follicle cell growth or differentiation (Hsueh et al., 1984). Similar mechanisms may exist in the bovine ovary for estrogens, but further studies are needed to identify and reveal these processes.

In the present study, the localization of ER β mRNA in the ovary was examined in randomly selected cows irrespective of the cycle stage of these animals, and therefore without taking into account the effects of circulating ovarian hormones and gonadotrophins. It has been demonstrated that hormonal superovulatory treatment in cows does not affect the topographical distribution and staining intensity of receptors for ER α and progesterone in the endometrium (Dall'Aglio et al., 1999). Nevertheless, in the ovary estrogens downregulate granulosa cell expression of ER β protein (Sharma et al., 1999) and that treatment of rodents with LH results in a marked decrease of ER β in the ovary (Byers et al., 1997). Consequently, cycle-dependent differences in ER β mRNA expression in the ovary may be expected, and at this moment an elaborate experiment is performed in which the presence of ER β mRNA is studied in detail in a higher number of cows in each of the five different cycle stages, i.e., prooestrus, oestrus, metoestrus, early dioestrus and late dioestrus.

In summary, the present findings have clearly established a cell-specific localization of $ER\beta$ mRNA in several cell types in the bovine ovary. Some of these findings are in agreement with earlier studies in various mammalian species, while others are contradictory even to findings in cattle. Further comprehensive studies are needed and planned to confirm and explain the findings and hypotheses proposed here.

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1.1.2 Localization of estrogen receptors within various bovine ovarian cell types at different stages of the estrous cycle

Modified from:

LOCALIZATION OF ESTROGEN RECEPTORS WITHIN VARIOUS BOVINE OVARIAN CELL TYPES AT DIFFERENT STAGES OF THE ESTROUS CYCLE D'Haeseleer M, Simoens P, Van den Broeck W Anatomia Histologia Embryologia 2006; in press

Summary

In the present study ER α and ER β mRNA were localized in various ovarian cell types of 23 cows at different cycle stages. ER α was detected by immunohistochemistry and the localization of ER β mRNA was examined using *in situ* hybridization. The immunostaining of ER α was low in the ovarian follicles, tunica albuginea and surface epithelium, but high in cells of the deep stroma and superficial stroma, which indicates a functional role of ER α in the cells surrounding the follicles. In contrast, ER β mRNA scores were low to moderate in primordial and primary follicles and increased with the development of the follicle. ER β mRNA scores were higher in cystic follicles than in obliterative follicles. In corpora lutea and corpora albicantia the scores for ER β mRNA were moderate. Furthermore, in corpora lutea ER β mRNA levels showed cyclic variations and were low during early dioestrus. The correlation between the plasma progesterone levels and the score for ER β was low and negative in all ovarian cell types. This study demonstrates the predominant role of ER β mRNA and ER α in most cell types suggests possible interactions between both ER subtypes.

Introduction

Estrogens regulate follicular growth and ovarian function by binding to estrogen receptors (ERs) (Couse et al., 1999). Binding of the ER-ligand complex to target DNA at estrogen response elements (ERE) results in the transcription of various genes. In many species two subtypes of the estrogen receptor have been demonstrated, viz. estrogen receptor α (ER α) and estrogen receptor β (ER β). A third type of estrogen receptor ER γ has been detected in the teleost *Micropogonias undulatus* (Hawkins et al., 2000), in mouse (Lorke et al., 2000) and in humans (Heard et al., 2000). ER α and ER β may form both homodimers and heterodimers resulting in receptor molecules with different functional activities (Pettersson et al., 1997; Enmark and Gustafsson, 1999). Experiments in mice lacking genes for either or both ER types have demonstrated the biological significance of each subtype in male and female reproduction (Lubahn et al., 1993; Krege et al., 1998; Couse et al., 1999).

The tissue distribution of ER α and ER β has been described for a wide variety of species in different organs. The presence of ER α has been shown in humans (Coppens et al., 1993), cattle (Dall'Aglio et al., 1999; Van den Broeck et al., 2002), dog (Vermeirsch, 2001) and rat (Pelletier et al., 2000). ER α mRNA has been localized in various species including humans (Revelli et al., 1996), cattle (Berisha et al., 2002), pig (Slomczynska et al., 2001), dog (Hatoya et al., 2003), rat (Kuiper et al., 1996) and mouse (Tremblay et al., 1997), and is expressed predominantly in the pituitary gland, ovary, uterus, testis, epididymis and adrenal gland (Kuiper et al., 1997).

ER β has been found in many vertebrate species and is expressed predominantly in the ovary and the prostate (Enmark et al., 1997). ER β mRNA has been described in different ovarian structures of various species including humans (Enmark et al., 1997), cattle (Rosenfeld et al., 1999), pig (Slomczynska et al., 2001), sheep (Jansen et al., 2001), dog (Hatoya et al., 2003) and rat (Kuiper et al., 1996; Wang et al., 2000).

Although the localization of ER α and ER β has already been described in the bovine ovary, the localization patterns of the different ovarian tissues are not well known (Rosenfeld et al., 1999; Manikkam et al., 2001b). Therefore, in a preliminary study, a probe for *in situ* hybridization was generated to localize ER β mRNA in the bovine ovary and an immunohistochemical method was developed to investigate the presence of ER α (D'Haeseleer

et al., 2005; Van den Broeck et al., 2002). The aim of the present study was to determine the morphological localization of ER α and ER β in the different cell types of the bovine ovary. Furthermore, putative differences between the receptor score in active and inactive ovaries were examined. Finally, possible cyclic alterations of receptor localization in the different cell types, and their correlation with the plasma progesterone level were determined.

Materials and methods

Animals and samples

Twenty-three animals were used, namely 7 cows that were experimentally synchronized and 16 slaughterhouse cows. For oestrus synchronization 7 cows were treated by means of PRID[®] insertion (Ceva Sante Animal, Brussels, Belgium) for 12 days. Blood samples for LH surge and plasma progesterone determination were taken every 4 hours in heparinized tubes from 36 hrs until 60 hrs after PRID[®] removal (Verberckmoes et al., 2004; Henry et al., 1987). Six of the seven cows were slaughtered 72 hrs after PRID[®] removal, i.e., at the day of oestrus (D0). One cow was slaughtered 15 days after PRID[®] removal (D13 of the estrous cycle) when a functional corpus luteum was present. From 16 other cows, without symptoms of any reproductive pathology, the genital tracts and blood samples were collected at the moment of slaughter. Both ovaries were classified using morphologic parameters as described by Ireland et al. (Ireland et al., 1980). The ovary showing a mature follicle or a young corpus luteum was classified as the active ovary. Subsequently, each ovary was bisected and sampled.

For the immunohistochemical determination of ER α , one part of the ovarian samples was fixed for 6 h in 3.5% phosphate-buffered formaldehyde (pH 6.7) at room temperature and dehydrated in increasing concentrations of ethanol. Subsequently, all tissue samples were embedded in paraffin, and 5 μ m thin sections were cut. The sections were mounted on 3-aminopropyl-triethoxysilane-coated slides (Sigma, S. Louis, MO, USA) and dried overnight at 37°C.

For the detection of ER β mRNA the other part of the ovarian samples was fixed in a Rnase-free phosphate buffered saline (PBS) solution containing 4% paraformaldehyde at 4°C
for 8 hours and dehydrated in increasing methanol concentrations. Afterwards the tissue samples were embedded in RNase free paraffin, and 5 µm thin sections were cut. The sections were mounted on poly-L-lysine-coated (Sigma Diagnostics Inc., S. Louis, MO, USA) SuperFrost[®]Plus-slides (Menzel-Gläser, Braunschweig, Germany) and dried overnight at 37°C.

The stage of the estrous cycle was determined for each cow on the basis of the morphological appearance of ovarian follicles and corpora lutea and the presence of uterine cervical mucus. Besides, the plasma progesterone concentration of each individual animal was determined and compared with the criteria described by Sartori et al. (2004). The cows were classified in 5 groups: 6 animals were in oestrus (synchronized cows), 2 animals in metoestrus (slaughterhouse cows), 5 animals in early dioestrus (4 slaughterhouse cows and one synchronized cow), 5 animals in late dioestrus (slaughterhouse cows) and 5 animals in prooestrus (slaughterhouse cows) (Fig. 1).



Fig. 1: Plasma progesterone concentrations of 23 cows in various stages of the estrous cycle.

Immunohistochemical detection of ERa

For the detection of ER α , the assay described by Van den Broeck et al. (2002) was used. The primary antibody was a monoclonal mouse-anti-human antibody clone 1D5 (DAKO, Prosan, Merelbeke, Belgium) which is ER α -specific. The antigen-antibody complex was visualized using biotinylated rabbit anti-mouse antibodies (DAKO) and diaminobenzidine was used as chromogen substrate (DAKO). For each cell type, the presence of ER α was assessed by counting the cells presenting a brown nuclear staining.

In every staining procedure, positive and negative controls were included. Bovine uterine tissues known to be positive for ER α from a previous study served as positive controls. As negative controls tissue sections incubated with dilution buffer instead of the primary antibody, secondary antibody, and/or peroxidase-conjugated streptavidin (BioGenex, San Ramon, CA, USA) were used.

In situ hybridization of ER\$ mRNA

Probe multiplication

The ligated complementary DNA fragment containing the 602 base pairs fragment of ERβ (D'Haeseleer et al., 2005) was purified and the plasmid DNA was linearized using restriction enzymes *Sac*II and *Sac*I to allow *in vitro* run-off synthesis of sense- and antisense-oriented RNA probes, respectively. After phenol-chloroform extraction and ethanol precipitation, digoxigenin (DIG)-labelled sense and antisense riboprobes were generated by transcription using the SP6-T7 DIG RNA Labelling Kit (Roche Molecular Biochemicals, Indianapolis, USA) and purified with Mini quick Spin RNA columns (Roche Molecular Biochemicals, Indianapolis, USA). The yield of the probes was estimated by DIG quantification on DIG control test strips (Roche Diagnostics Corporation, Indianapolis, USA).

In situ hybridization

In situ hybridization was carried out in all ovarian tissue sections as described previously (D'Haeseleer et al., 2005). The presence of ER β mRNA was assessed by counting the cells presenting a brown cytoplasmic staining. Two controls were performed to confirm that the antisense riboprobe hybridized specifically to ER β mRNA, viz., hybridization with the

labelled sense probe and hybridization with hybridization buffer only. None of these treatments showed in any detectable ribopobe binding.

Microscopic analysis

In each ovary the different follicle types, corpora hemorrhagica, corpora lutea, corpora albicantia, deep stroma, superficial stroma, tunica albuginea and surface epithelium were microscopically evaluated. All follicles were examined and classified by histological stage into primordial, primary, secondary and tertiary follicles according to the criteria listed in the Nomina Histologica (1994). Tertiary follicles containing less than 1 % apoptotic cells were classified as vital tertiary follicles, whereas follicles containing more than 1% apoptotic cells were regarded as atretic (D'Haeseleer et al., 2006a). The atretic follicles were further classified as obliterative or cystic according to the criteria described by Dellman and Eurell (1998).

At least 100 cells of every cell type were counted on digital micrographs taken randomly at magnification 600 x with an Olympus DP 50 digital camera mounted on an Olympus BX61 light microscope (Olympus Belgium N.V., Aartselaar, Belgium). If too many cells were present in one micrograph, cells were counted in 4 grids of 2500 μ m² that were projected randomly over the micrograph. Staining intensity scores ranged from 0 (no staining), 1 (weak staining), 2 (moderate staining) to 3 (strong staining). The presence of ER α was evaluated using a score SER α which was calculated according to the formula modified after Boos et al. (1996): SER α = 0 n₀ + 1 n₁ + 2 n₂ + 3 n₃ with n₀, n₁, n₂ and n₃ are the percentage of cells exhibiting a staining intensity 0, 1, 2 and 3, respectively. A similar formula (SER β mRNA) was used for calculating the score for ER β mRNA.

Statistical analysis

Throughout the study, results were represented as mean values and variation was expressed as standard error of the mean (SEM). Possible differences in the ER score between active and inactive ovaries, between the various cell types and between the 5 stages of the estrous cycle were analysed by means of the Kruskall-Wallis test. The Tukey's test was applied to determine significant (P < 0.05) or highly significant (P < 0.01) differences between particular cycle stages. The Spearman rank correlation test was used for determination of correlations between the plasma progesterone levels and the scores for ER α and ER β mRNA. Statistical analyses were performed with procedures available in SPSS 12.0 (SPSS Inc., Chicago, Illinois, USA).

Results

Localization of ER α and ER β mRNA in the bovine ovary

In all cows immunoreactivity of ER α was completely absent in primordial and primary follicles, but was observed in low amounts in secondary follicles, vital and atretic tertiary follicles, corpora lutea and corpora albicantia. Furthermore, immunostaining of ER α was detected in cells of the deep and superficial stroma, tunica albuginea and surface epithelium (Fig. 2a).

 $ER\beta$ mRNA was demonstrated in all different ovarian cell types, viz., the various vital and atretic follicles, corpora hemorrhagica, corpora lutea, corpora albicantia, deep stroma, superficial stroma, tunica albuginea and surface epithelium (Fig. 2b).



Fig. 2: Mean scores for estrogen receptors in ovarian tissues of 23 cows (pooled data of active and inactive ovaries). Fig. 2a: Mean scores for estrogen receptor α in the different bovine ovarian tissues. Fig. 2b: Mean scores for estrogen receptor β mRNA in the different bovine ovarian tissues.

PdF: primordial follicle, PF: primary follicle, SF: secondary follicle, VTF: vital tertiary follicle, OAF: obliterative atretic follicle, CAF: cystic atretic follicle, CH: corpus hemorrhagicum, CL: corpus luteum, CA: corpus albicans, DS deep stroma, SS: superficial stroma, TA: tunica albuginea, SE: surface epithelium, fc: follicle cells, tf: theca follicularis, gc: granulosa cells, ti: theca interna, te: theca externa, ll: large lutein cells, sl: small lutein cells, ps: perivascular stroma, is: internal stroma, cs: capsular stroma. Between the various cell types of a particular ovarian structure, scores are significantly different when indicated by different superscripts.

Cell-specific variation of ER α and ER β mRNA

The score for ER α varied from the score for ER β mRNA, and for each receptor subtype the score varied between the different ovarian cell types.

In general the immunoreactivity of ER α was low in all ovarian structures (Fig. 2a). It was never observed in the follicle cells of primordial, primary and secondary follicles, apart from a very weak immunostaining in the theca layers of the secondary follicles (P < 0.01). In vital tertiary follicles the SER α was low in the granulosa cells and slightly higher in the theca externa cells (P < 0.01), whereas in atretic tertiary follicles the score for ER α was low and only observed in the theca externa cells. ER α was not detected in cells of the corpora hemorrhagica. In corpora lutea scores for both large and small lutein cells were low, whereas in the stroma cells the score was still lower. In corpora albicantia the ER α immunostaining was stronger in the capsular stroma than in the internal stroma. ER α scores were high in cells of the deep stroma and the superficial stroma, while low values were found in cells of the tunica albuginea (P < 0.01). Finally, the score for ER α was very low in cells of the surface epithelium.

The scores for ER β mRNA in the different ovarian cell types are shown in fig. 2b. These scores were moderate in the follicle/granulosa cells of primordial, primary, secondary and vital tertiary follicles, and much higher in the granulosa cells of atretic follicles. In obliterative atretic follicles a nearly similar SER β mRNA was observed in the granulosa and theca layers, whereas in cystic atretic follicles the score was much higher in the theca externa than in the granulosa layers (P < 0.01). The SER β mRNA in the theca cells was generally higher than in the follicle/granulosa cells, but this difference was only significant for secondary follicles (P < 0.01) and for vital tertiary follicles (P < 0.05). The granulosa cells of the corpora hemorrhagica, the lutein cells of the corpora lutea and the internal stroma cells of the corpora albicantia showed a similar moderate score for ER β mRNA. Furthermore, the SER β mRNA was high in cells of the deep stroma and superficial stroma, and slightly lower in the tunica albuginea and the surface epithelium.

Ovary-specific variation

Comparison between active and inactive ovaries revealed no significant differences for SER α , despite the distinct differences in the deep and superficial stroma (Fig. 3a).

The score for ER β mRNA was generally higher in inactive than in active ovaries. Significant differences were observed in the granulosa (P < 0.05) and theca interna layer (P < 0.05) of vital tertiary follicles and in the internal stroma of the corpora albicantia (P < 0.05) (Fig. 3b).



Fig. 3: Mean scores for estrogen receptors in the different cell types of the active and inactive ovaries of 23 cows. Fig. 3a: Mean scores for ER α in the various cell types of the bovine active and inactive ovary. Fig. 3b: Mean scores for ER β mRNA in the various cell types of the bovine active and inactive ovary. Abbreviations: See fig. 2. For a particular ovarian structure significant differences between the active and inactive ovary were observed for ER β mRNA and are indicated by different superscripts.

Cycle-specific variation

In all ovarian parenchymatous structures the immunoreactivity of ER α was very low in each of the cycle stages (Fig. 4a). In contrast, statistically significant cycle-dependent differences were seen in the deep and superficial stroma (Fig. 5a-c). In these cell types, the levels were highest in oestrus, decreased in metoestrus and early dioestrus, increased in late dioestrus and decreased again during procestrus (P < 0.01).

High albeit statistically non-significant variations in the scores for ER β mRNA during the bovine cycle stages were observed in the different cell types (Fig. 4b). In both the primordial and primary follicles similar patterns were seen, with the highest scores in early dioestrus and the lowest score during metoestrus. In secondary follicles the follicle cells showed a gradually increasing ER β mRNA level from oestrus to prooestrus, whereas in the theca layers a nearly similar level was observed in all cycle stages, apart from higher scores present during metoestrus. In the cell layers of vital tertiary follicles the SER β mRNA was moderate to high during oestrus, increased in metoestrus, decreased during early dioestrus, and increased again during late dioestrus and prooestrus. A similar cyclic pattern was seen in the superficial stroma cells surrounding the follicles.

The staining pattern of ER β mRNA in obliterative atretic follicles differed from that in cystic atretic follicles. Obliterative follicles showed high levels in oestrus and early dioestrus, while in cystic follicles the highest level was seen during metoestrus and prooestrus. The corpora hemorrhagica, which were only observed during oestrus, metoestrus and early dioestrus, showed moderate SER β mRNA. In both the large and small lutein cells of corpora lutea the score for ER β mRNA was moderate in oestrus, increased during metoestrus, decreased during early dioestrus and increased again during late dioestrus and prooestrus. In the stroma cells of corpora lutea the score was generally lower than in the lutein cells in all cycle stages, except during prooestrus.

In corpora albicantia nearly similar levels for ER β mRNA were seen during all cycle stages. In cells of the deep stroma and tunica albuginea a similar pattern for ER β mRNA was present, although the SER β mRNA in the latter cell type was lower. In cells of the surface epithelium moderate scores were detected with no cycle dependent variations (Fig. 5).

In general, the correlations between the plasma progesterone levels and the SER of all ovarian cell types were low and negative (Table 1). These correlations were significant for ER α in the theca cells of vital tertiary follicles (p < 0.01), in cells of the corpora albicantia (p < 0.05), in stroma cells (p < 0.01) and in the cells of the tunica albuginea (p < 0.01). Significant correlations were found for ER β mRNA in the corpora hemorrhagica.

		Plasma Progesterone Level	
		ERα	ERβ
Secondary Follicle	follicle cells	/	+ 0,47
	theca follicularis	- 0,27	- 0,08
Vital Tertiary Follicle	granulosa cells	- 0,26	+0,10
	theca interna	/	- 0,36
	theca externa	- 0,55**	- 0,23
Obliterative Atretic Follicle	granulosa cells	/	- 0,33
	theca interna	/	- 0,27
	theca externa	- 0,26	- 0,24
Cystic Atretic Follicle	granulosa cells	/	- 0,13
	theca interna	/	- 0,10
	theca externa	- 0,37	+ 0,15
Corpus Hemorrhagicum	granulosa cells	/	/
	theca interna	/	$+0,87^{**}$
	theca externa	/	$+0,87^{**}$
Corpus Luteum	large lutein cells	+0,29	+ 0,20
	small lutein cells	+0,29	+ 0,23
	perivascular stroma cells	+0,29	- 0,05
	capsular stroma cells	+0,29	- 0,25
Corpus Albicans	internal stroma cells	- 0,44*	+0,04
	capsular stroma cells	- 0,45*	- 0,35
Deep Stroma		- 0,55**	+ 0,24
Superficial Stroma		- 0,66**	- 0,21
Tunica Albuginea		- 0,63**	+ 0,24
Surface Epithelium		- 0,26	+0,03

Table 1: Correlations between the plasma progesterone levels and the progesterone receptor scores in the different ovarian cell types. "/" not detected; * P < 0.05; ** P < 0.01





Fig. 4: Mean scores for estrogen receptors in various bovine ovarian cell types at different stages of the estrous cycle. Fig. 4a: Mean scores for ER α in the various cell types of the bovine ovary at different cycle stages. Fig. 4b: Mean scores for ER β mRNA in the various cell types of the bovine ovary at different cycle stages. Abbreviations: See fig. 2. Significant differences between the cycle stages were observed in the deep and superficial stroma for ER α and are indicated by different superscripts.



Fig. 5: Micrographs of different bovine ovarian cell types at various estrous stages showing immunostaining of ER α (5a-c) and ER β mRNA (5d-k). Fig. 5a: Outer region of the ovarian cortex during oestrus with low scores for ER α (SER α) in the surface epithelium (1), higher scores in the superficial stroma (3), and no expression in the tunica albuginea (2) nor in the primary follicles (4) (Bar = $100 \mu m$). Fig. 5b: Secondary follicle during procestrus showing no expression for ER α in the granulosa cells (5), low SER α in the theca cells (6) and high SER α in cells of the superficial stroma (7) (Bar = 100 μ m). Fig. 5c: Overview of a corpus albicans during oestrus with low SER α in the internal stroma cells (8) and higher SER α in the capsular stroma cells (9) (Bar = 50 μ m). Fig. 5d: Detail of the outer region of the bovine ovarian cortex during early dioestrus with moderate scores for ER β mRNA (SER β mRNA) in cells of the surface epithelium (10), higher scores in the tunica albuginea (11) and moderate scores in the superficial stroma (12) (Bar = 100 μ m). Fig. 5e: Primary (13) and secondary (14) follicles at oestrus with moderate SER β mRNA in the follicle cells and high scores in the superficial stroma (15) (Bar = 50 μ m). Fig. 5f: Vital tertiary follicle at procestrus with high SER β mRNA in the granulosa cells (18) and cells of the cumulus opphorus (17). The oocyte (16) is surrounded by the zona pellucida (Bar = 100 μ m). Fig. 5g: Vital tertiary follicle at early dioestrus with low to moderate SER β mRNA in the granulosa layer (19), higher scores in the theca interna (20) and still higher scores in the theca externa (21) (Bar = $100 \ \mu m$). Fig. 5h: Cystic attentic tertiary follicle at metoestrus with low to moderate SER β mRNA in the granulosa layer (22) and higher scores in the theca interna (23) and theca externa layers (24) (Bar = $100 \mu m$). Fig. 5i: Overview of an obliterative attetic tertiary follicle at oestrus with low SERB mRNA in cells of the collapsed theca interna layers (25), and higher scores in cells that are located in the thickened basal lamina (26) and in the theca externa layers (27) (Bar = $200 \ \mu m$). Fig. 5j: High magnification of a corpus luteum at early dioestrus with low to moderate SER β mRNA in the lutein cells (28) (Bar = 200 μ m). Fig. 5k: Overview of a corpus albicans at oestrus with low to moderate SER β mRNA in the internal stroma cells (29) and high SER β mRNA in cells of the deep stroma (30) (Bar = $100 \mu m$).

Discussion

In this study the presence of ER α and ER β mRNA was demonstrated in the different bovine ovarian cell types at different stages of the estrous cycle.

The results show that both types of ER have different tissue localization patterns, suggesting that each receptor type has a specific function.

In the follicle cells of primordial, primary and secondary follicles moderate scores for ER β mRNA were observed, whereas ER α was not detected. This indicates that estrogens, which play an important role in follicular growth (Hulshof et al., 1995) and development (Manikkam et al., 2001a), act upon the early follicular stages via ER β . These findings are in agreement with previous studies in the mouse (Couse et al., 1997) and dog (Hatoya et al., 2003).

In our study, the score for ER β mRNA was lower in the granulosa cells of vital tertiary follicles than in the follicle cells of secondary follicles. Moreover, the score for ER β mRNA in vital tertiary follicles was high in procestrus, decreased during cestrus, to increase again during metoestrus. These observations may reflect a down-regulation of ER β mRNA by pituitary gonadotrophins during oestrus as earlier described in the rat ovary (Byers et al., 1997). In contrast, low levels of ER α were localized in the granulosa and theca externa layers of vital tertiary follicles. This is in accordance with studies of mouse (Orimo et al., 1995), rat (Byers et al., 1997), rabbit (Iwai et al., 1991) and baboon (Billiar et al., 1992). According to Couse et al. (1997), the presence and interaction of both receptors within vital tertiary follicles are required for the late stage of follicular growth. Because ER α and ER β mRNA were expressed in the same cells, both ER α and ER β homodimers and/or ER α /ER β heterodimers could be formed (Pettersson et al., 1997; Pepe et al., 2002). These homo- and heterodimers can also have affinities for other ligands that have estrogen-like activity (Sun et al., 1999). ERβ mRNA is localized in vital tertiary follicles with moderate levels in the granulosa cells, higher levels in the theca interna and still higher levels in the theca externa. This observation is consistent with reports of ER^β mRNA or protein in the pig (Slomczynska et al., 2001) and monkey (Pelletier et al., 1999) and it indicates that $ER\beta$ has functions that are specific in the different cell layers of tertiary follicles.

The SER β mRNA in the granulosa cells of both obliterative and cystic atretic follicles was nearly two times higher than in vital tertiary follicles. This suggests that the level of ER β mRNA in the granulosa layers of tertiary follicles depends upon the physiological status of the follicle. Studies of ER α KO and ER β KO mice have suggested that the presence of either ER α or ER β is sufficient to mediate the protective effects of estrogens against atresia of tertiary follicles (Emmen et al., 2005). Our results suggest that in cattle ER β is not sufficient in conferring the protective capacity of estrogens. Another explanation, which is rather speculative, is that the score for ER β mRNA increases when follicular atresia occurred. It is possible that other factors or another ER, maybe the recently detected ER γ , contributed to the protective capacity of estrogens in bovine tertiary follicles.

In our study both ER α and ER β mRNA were demonstrated in the corpora lutea. This observation is in accordance with studies of mouse (Rosenfeld et al., 2001), rat (Wang et al., 2000) and pig (Slomczynska et al., 2001). The presence of ER α and ER β mRNA in the corpus

luteum is suggestive of their involvement in the process of maintenance and luteinization of the corpus luteum (Rosenfeld et al., 2001).

In all cell types of the corpus luteum ER β mRNA had a similar localization pattern with increasing levels after early dioestrus, while immunostaining of ER α was exclusively observed during late dioestrus. These findings are suggestive of an up-regulation of both types of ER by progesterone. It is of interest that the enhanced secretion of progesterone in the corpus luteum only leads to up-regulation of ERs, whereas the progesterone receptors remain unaltered (D'Haeseleer et al., 2006c). The simultaneous presence of ER α and ER β mRNA in the corpus luteum during dioestrus may lead to heterodimerisation and possible involvement of ER α/β heterodimers in corpus luteum regression.

High scores for ER β mRNA and relatively high amounts of ER α positive cells were found in cells of the deep stroma and superficial stroma. Generally, an up-regulation of ER α and ER β mRNA was obvious around the time of oestrus. This may account for specific compositional changes of the stromal cells surrounding the follicular structures during oestrus, which has also been described in a study of estrogen receptors in the bovine oviduct (Ulbrich et al., 2003). Furthermore, the high scores for ER α in the superficial stroma cells which are in direct contact with the follicles and in the deep stroma cells which surround the corpora albicantia are suggestive of possible stromal-epithelial interactions as reviewed by Cunha et al. (2004).

For all ovarian structures the score for ER β mRNA was higher in inactive than in active ovaries, except during the early follicular stages where the highest levels for ER β mRNA were seen in the active ovaries. This implicates a cell specific involvement of ER β mRNA in the development of follicles in the active ovary.

Although the biological significance of ER α and ER β has not yet been fully established, the different tissue localization patterns and the cyclic alterations in the bovine ovarian tissues suggest a specific cellular function of each receptor type. This is in agreement with the study of Kuiper et al. (1997) who proved by analysis of mRNA distribution that ER β expression does not exactly match with the distribution of ER α , although colocalization of both types of ER can occur. Further studies are needed to elucidate the precise role of the different types of estrogen receptors, including ER γ , in the different organs and in particular in the ovary.

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1.2 Cell-specific localization of progesterone receptors in the bovine ovary at different stages of the estrous cycle

Modified from:

CELL-SPECIFIC LOCALIZATION OF PROGESTERONE RECEPTORS IN THE BOVINE OVARY AT DIFFERENT STAGES OF THE ESTROUS CYCLE D'Haeseleer M, Simoens P, Van den Broeck W Animal Reproduction Science 2006; in press

Summary

This immunohistochemical study describes the localization of progesterone receptors (PR) in the bovine ovary of 23 cows at different stages of the estrous cycle. In primordial, primary and secondary follicles the score for PR in the follicle cells increased progressively with the maturation of the follicle. In vital tertiary follicles and cystic atretic follicles a moderate score for PR was found, while in obliterative atretic follicles the score was much lower. Scores were high in corpora hemorrhagica, low in corpora lutea and still lower in corpora albicantia. Low PR scores were also found in the tunica albuginea and surface epithelium. Cyclic variations of PR immunoreactivity were manifest in most ovarian tissues. Follicular scores for PR were high in oestrus and decreased during the following stages, whereas scores in corpora lutea cells varied according to a characteristic pattern with high levels during oestrus and metoestrus.

The variations in the scores for PR in the different ovarian cell types suggest a cell-specific and cycle-dependent influence of progesterone. A negative correlation was found between the PR scores and the plasma progesterone concentration.

Introduction

Progesterone, synthesized and secreted by the corpus luteum, is one of the major regulators of the reproductive cycle in mammals. It exerts its effects on the growth and differentiation of ovarian structures and renders the endometrium receptive to the implantation of the embryo (Pinter et al., 1996; Dellmann and Eurell, 1998). Progesterone acts on its target tissues after binding to a specific intracellular progesterone receptor (PR). Two functionally different receptor isoforms (PRA and PRB) have been identified (Vegeto et al., 1993; Graham and Clarcke, 1997; Mangal et al., 1997). An additional progesterone binding protein PRC, detected in the human breast cancer cell line T47D, may inhibit the activity of both PRA and PRB in the presence of progesterone (Wei and Minner, 1994).

The presence of PR has been demonstrated in the ovary of a number of species, viz. in man (Iwai et al., 1990), primates (Hild-Petito et al., 1988), cattle (Van den Broeck et al., 2002), pig (Slomczynska et al., 2000), rabbit (Iwai et al., 1991a), dog (Vermeirsch et al., 2001) and mouse (Lydon et al., 1995).

Although the distribution of PR has already been studied in the bovine ovary, it is not well known whether the localization of PR varies during the estrous cycle stages. Therefore, the aim of the present study was to determine the morphological localization of PR (PRA and PRB) in various cell types of the bovine ovary at the different stages of the estrous cycle. Putative differences between the scores of receptor immunoreactivity in active and inactive ovaries were also examined. Furthermore, the correlation between the plasma progesterone level and the receptor immunopresence in the different ovarian cell types was determined.

Materials and methods

Animals and samples

Samples of both ovaries and blood were obtained from 23 cows at the moment of slaughter in a local slaughterhouse. All animals had been used in our previous study on the localization of the estrogen receptors in the bovine ovary (D'Haeseleer et al., 2006b). The cows were classified in 5 groups: 6 animals were in oestrus (synchronized cows), 2 animals in metoestrus (slaughterhouse cows), 5 animals in early dioestrus (4 slaughterhouse cows and one synchronized cow), 5 animals in late dioestrus (slaughterhouse cows) and 5 animals in procestrus (slaughterhouse cows) (Fig. 1).



Fig. 1: Plasma progesterone concentrations of 23 cows in various stages of the estrous cycle.

For immunohistochemical determination of PR the ovarian tissue samples were fixed for 6 hours in 3.5% phosphate-buffered formaldehyde (pH 6.7) at room temperature and dehydrated in increasing concentrations of ethanol. Subsequently, all tissue samples were embedded in paraffin and 5 µm thin sections were cut. The sections were mounted on 3-aminopropyl-triethoxysilane-coated slides (APES, Sigma, St. Louis, MO, USA) and dried overnight at 37° C.

Immunohistochemical detection of PR

The presence of PR was determined immunohistochemically as described by Van den Broeck et al. (2002) by means of a monoclonal mouse-anti-human-PR antibody (clone 10A9, ImmunoTech, Marseille, France) which binds to both PRA and PRB. Rabbit anti-mouse antibodies conjugated to biotin (DAKO, Prosan, Merelbeke, Belgium) were used as secondary antibodies and peroxidase-conjugated streptavidine as enzyme label. Reaction of DAB chromogen substrate (Liquid DAB+, DAKO, Prosan, Merelbeke, Belgium) with peroxidase resulted in a brown nuclear staining. Positive and negative controls were included in every staining procedure. Positive controls were bovine ovarian tissues known to be positive for PR, whereas similar tissue sections incubated with dilution buffer instead of the primary antibody, biotinylated secondary antibody, and/or peroxidase-conjugated streptavidine served as negative controls.

Microscopic analysis

In each ovary the different follicle types, corpora hemorrhagica, corpora lutea, corpora albicantia, deep stroma, superficial stroma, tunica albuginea and surface epithelium were evaluated. All follicles were examined and classified by histological stage into primordial, primary, secondary and tertiary follicles according to the criteria listed in the Nomina Histologica (1994). Tertiary follicles containing less than 1 % apoptotic cells were classified as vital tertiary follicles, whereas follicles containing more than 1% apoptotic cells were regarded as atretic (D'Haeseleer et al., 2006a). The atretic follicles were further classified as obliterative or cystic according to the criteria described by Dellmann and Eurell (1998). For each cell type the presence of PR was assessed by counting the cells presenting a brown nuclear staining.

At least 100 cells of every cell type were counted on digital micrographs taken randomly at magnification 600 x with an Olympus DP 50 digital camera mounted on an Olympus BX61 light microscope (Olympus Belgium N.V., Aartselaar, Belgium). If too many cells were present in one micrograph, cells were counted in 4 grids of 2500 μ m² that were projected randomly over the micrograph. Staining intensity scores ranged from 0 (no staining), 1 (weak staining), 2 (moderate staining) to 3 (strong staining). The immunopresence of PR was evaluated using a score (SPR) calculated according to the formula modified after Boos et al. (1996): SPR= 0 n₀ + 1 n₁+ 2 n₂ + 3 n₃ with n₀, n₁, n₂ and n₃ are the percentage of cells exhibiting a staining intensity 0, 1, 2 and 3, respectively.

Statistical analysis

Throughout the study, results were represented as mean values and variation was expressed as standard error of the mean (SEM). Possible differences in PR score between active and inactive ovaries, between the various cell types and between the 5 stages of the estrous cycle were analysed by means of the Kruskall-Wallis test. The Tukey's test was applied to determine significant (P < 0.05) or highly significant (P < 0.01) differences between particular cycle stages.

The Spearman rank correlation test was used for determination of correlations between the plasma progesterone levels and the scores for PR. Statistical analyses were performed with procedures available in SPSS 12.0.

Results

Localization of PR in the bovine ovary (Fig. 2a)

Immunoreactivity of PR was detected in the primordial, primary, secondary and tertiary vital follicles, in cystic and obliterative atretic follicles, in corpora hemorrhagica, corpora lutea and corpora albicantia, in cells of the deep and superficial stroma, and in the tunica albuginea and surface epithelium.



Fig. 2: Mean scores for progesterone receptors in ovarian tissues of 23 cows. **Fig. 2a:** Mean scores for the progesterone receptors in the different bovine ovarian cell types (pooled data of the active and inactive ovaries). Between the various cell types of a particular ovarian structure, scores are significantly different when indicated by different superscripts. **Fig. 2b:** Mean scores for the progesterone receptors in the various cell types of the active and inactive bovine ovary. No significant differences of the scores for progesterone receptors were observed between the active and inactive ovary.

PdF: primordial follicle, PF: primary follicle, SF: secondary follicle, VTF: vital tertiary follicle, OAF: obliterative atretic follicle, CAF: cystic atretic follicle, CH: corpus hemorrhagicum, CL: corpus luteum, CA: corpus albicans, DS deep stroma, SS: superficial stroma, TA: tunica albuginea, SE: surface epithelium, fc: follicle cells, tf: theca follicularis, gc: granulosa cells, ti: theca interna, te: theca externa, ll: large lutein cells, sl: small lutein cells, ps: perivascular stroma, is: internal stroma, cs: capsular stroma.

Cell-specific variation of the PR expression (Fig. 2a)

Many differences were observed in the scores for PR of the different ovarian cell types. The PR immunopresence in the follicle cells of primordial, primary and secondary follicles was high and increased from primordial to secondary follicles (P < 0.05), whereas the presence of PR in granulosa cells of vital tertiary follicles was clearly lower (P < 0.05). The score for PR in the granulosa cells of cystic atretic tertiary follicles was similar to that of the granulosa cells of vital tertiary follicles, whereas the score in the granulosa cells of obliterative atretic follicles was much lower (P < 0.01). Granulosa cells of corpora hemorrhagica showed a very high PR immunoreactivity. In the subsequent stage of corpora lutea the score for PR decreased in both the large and small lutein cells (P < 0.01) and reached the lowest level in internal stroma cells of corpora albicantia. In general, the score for PR in theca cells was lower than in the follicle/granulosa cells of the corresponding structures, and this difference was significant in secondary follicles (P < 0.01) and corpora hemorrhagica (P< 0.05). Furthermore, theca externa cells always showed a lower score compared to theca interna cells. The immunostaining for PR was moderate in cells of the deep stroma, whereas superficial stroma cells had a lower staining intensity, nearly similar to that of the theca cells in secondary and vital tertiary follicles. The surface epithelium showed a low score for PR, while in the tunica albuginea the score was lower than in any other ovarian structure (P <0.01).

Ovary-specific variation of the PR expression (Fig. 2b)

Comparison between active and inactive ovaries revealed no significant differences, although the PR immunoreactivity in the majority of the ovarian cell types was slightly higher in the active than in the inactive ovaries.

Cycle-specific variation of the PR expression (Fig. 2c, Fig. 3)

High variations in the PR immunoreactivity, concomitant with the bovine cycle stages, were observed in the different ovarian cell types. In primordial, primary and secondary follicles the score for PR was highest in oestrus, decreased during metoestrus, reached the lowest levels in early dioestrus, and increased again to moderate scores during the subsequent period of dioestrus and procestrus. These differences between the various cycle stages were

significant in the follicle cells (P < 0.05) and theca cells (P < 0.01) of secondary follicles. A similar staining pattern was seen in the granulosa and theca layers of vital tertiary follicles, although the PR immunoreactivity in theca interna and theca externa cells was slightly lower in procestrus than in late dioestrus.

In obliterative atretic tertiary follicles the PR immunostaining in granulosa and theca cell was low in oestrus and metoestrus, and minimal in the other stages. In cystic atretic tertiary follicles the granulosa and theca cells showed high levels during oestrus and the score for PR decreased during the next stages in granulosa cells, whereas in the theca layers this score decreased in metoestrus and early dioestrus to increase again during late dioestrus and prooestrus (P < 0.05).

The corpora hemorrhagica, which were only observed during oestrus, metoestrus and early dioestrus, showed a higher PR immunostaining than any other ovarian cell type.

In both the large and small lutein cell types of corpora lutea the score for PR was moderate at oestrus, increased during metoestrus and decreased during the subsequent cycle stages.

Perivascular stroma cells showed a similar staining pattern, whereas in the capsular stroma the cyclic differences were minimal. In corpora albicantia the score for PR was very low and little variation was seen between the different cycle stages.

In both deep and superficial stroma, the PR immunostaining was high during oestrus and metoestrus, very low during early dioestrus and higher again in the subsequent stage of dioestrus and in procestrus. The scores for PR were low in the surface epithelium and in the tunica albuginea, with minimal cyclic variations in the latter structure. In general, the correlation between the plasma progesterone levels and the PR score was negative for all ovarian cell types (Table 1). These correlations were highly significant (P < 0.01) in the obliterative follicles, the surface epithelium and the internal stroma cells of the corpora albicantia.



Fig. 2c: Mean scores for progesterone receptors in various bovine ovarian cell types during the different estrous cycle stages. Abbreviations: See fig. 2. Between the various cell types of a particular ovarian structure, PR scores are significantly different (* P < 0.05; ** P < 0.01) when indicated by different superscripts.

		Plasma Progesterone Level
Secondary Follicle	follicle cells	- 0,34
	theca follicularis	- 0,52*
Vital Tertiary Follicle	granulosa cells	- 0,13
	theca interna	- 0,06
	theca externa	- 0,31
Obliterative Atretic Follicle	granulosa cells	- 0,71**
	theca interna	- 0,55**
	theca externa	- 0,56**
Cystic Atretic Follicle	granulosa cells	- 0,24
	theca interna	- 0,39
	theca externa	- 0,46*
Corpus Hemorrhagicum	granulosa cells	- 0,44
	theca interna	+ 0,04
	theca externa	- 0,12
Corpus Luteum	large lutein cells	- 0,40
	small lutein cells	- 0,38
	perivascular stroma cells	- 0,23
	capsular stroma cells	- 0,16
Corpus Albicans	internal stroma cells	- 0,62**
	capsular stroma cells	- 0,24
Deep Stroma		- 0,09
Superficial Stroma		- 0,51*
Tunica Albuginea		- 0,25
Surface Epithelium		- 0,56 **

Table 1: Correlations between the plasma progesterone levels and the progesterone receptor scores in the different ovarian cell types. * P < 0.05; ** P < 0.01

Fig. 3: Micrographs of different bovine ovarian cell types showing PR expression (arrows). **Fig. 3a:** Primary (1) and early secondary (2) follicles at late dioestrus show immunoreaction for PR in the follicle cells (Bar = 50 μ m). **Fig. 3b:** Secondary follicle at early dioestrus with low scores for PR (SPR) in the granulosa cells (3) and in the theca cells (4) (Bar = 50 μ m). **Fig. 3c:** Overview of a vital tertiary follicle at procestrus with high SPR in cells of the cumulus oophorus (5), granulosa layer (6), theca interna (7) and theca externa (8) (Bar = 200 μ m). **Fig. 3d:** Cystic attetic tertiary follicle at oestrus with high SPR in apoptotic granulosa cells (9) and in theca interna cells (10), and moderate scores in cells of the theca externa (11) (Bar = 50 μ m). **Fig. 3e:** Overview of an obliterative attetic tertiary follicle at oestrus with low SPR in cells of the collapsed theca interna layer (12) and in cells that are located in the thickened basal lamina (13) (Bar = 200 μ m). **Fig. 3f:** High magnification of corpus luteum lutein cells (14) during procestrus with low SPR in the internal stroma cells (15) (Bar = 50 μ m). **Fig. 3h:** Cells of the deep stroma (16) at late dioestrus showing high SPR (Bar = 50 μ m). **Fig. 3i:** Outer region of the ovarian cortex during oestrus showing high immunoreactivity for PR in cells of the surface epithelium (17), low scores in the tunica albuginea cells (18) and high scores in the superficial stroma (19) (Bar = 50 μ m).



Discussion

The present study confirms previous data on the localization of PR in the bovine ovarian structures (Van den Broeck et al., 2002) and additionally shows that the various ovarian cell types exhibit different patterns of PR immunoreactivity during the estrous cycle. In the follicle cells of primordial, primary and secondary follicles the scores for PR were high and increased from primordial to secondary follicles. These data are in accordance with findings in primates (Hild-Pepito et al., 1988) and dogs (Vermeirsch et al., 2001), and they indicate that progesterone may regulate follicular growth during the early stages of follicular development.

Studies in the monkey ovary (Hild-Petito et al., 1988) showed that PR was exclusively located in the theca layers of secondary and tertiary follicles. In contrast, in our study PR immunoreactivity was observed in the different cell types of secondary and tertiary follicles, viz. the follicle/granulosa cells, theca interna and theca externa cells. Moreover, the scores for PR in the theca layers of secondary and tertiary follicles were similar to the scores of the surrounding superficial stroma cells. These data indicate that progesterone may regulate follicular development in the bovine ovary not only via granulosa cells, but also through theca and stroma cells. The possibility of such a paracrine action of ovarian steroids on follicles mediated through stromal steroid receptors has also been suggested by Revelli et al. (1996).

In all bovine follicular structures examined, the follicle/granulosa cells showed the highest PR immunostaining during oestrus, when ovulation occurs. These results are concomitant with earlier observations in the dog (Vermeirsch et al., 2001) and with a study on PR mRNA in the bovine ovary (Cassar et al., 2002). The crucial role of PR in the ovulatory process has been demonstrated in PR-deficient mice, since such mice develop large follicles but fail to ovulate (Lydon et al., 1995). All these findings emphasize the important role of progesterone and its receptor in the ovulation process. The expression of PR in granulosa cells in tertiary follicles is induced by the LH surge (Hild-Petito et al., 1988; Jo et al., 2002). The induction of PR mRNA has also been observed in macaque granulosa cells during periovulatory stages (Chandrasekher et al., 1994) and in porcine granulosa cells cultured *in vitro* after LH stimulation (Iwai et al., 1991b).

The finding that the PR expression in the bovine ovarian follicles presented a cyclic pattern that was similar in all follicle types, indicates that the role of PR in follicle development is specific in each stage of the ovarian cycle.

The score for PR in the bovine ovarian tissues was negatively correlated with the plasma progesterone concentrations. This finding indicates that progesterone may downregulate the expression of PR in the bovine ovary. A similar negative feed back process has also been described for the distribution of PR in the bovine endometrium (Boos et al., 1996).

The progesterone receptors are reported to mediate the protective effects of progesterone against apoptosis in the granulosa cells of bovine preovulatory follicles (Quirk et al., 2004). In our study vital tertiary and cystic atretic follicles showed nearly similar PR levels. These observations indicate that PR is sufficient in conferring the protective effects of progesterone in vital tertiary follicles. If this PR expression decreases, the protective influence of progesterone is lost. As a consequence, apoptosis can be induced in the vital tertiary follicle so that the latter degenerates into an obliterative atretic follicle. This explains the low expression level of PR in obliterative atretic follicles. However, in cystic atretic follicles the PR score is as high as in vital tertiary follicles. Apparently, in some cases progesterone cannot protect vital tertiary follicles to become atretic. A possible explanation is that in those cases the effect of pro-apoptotic factors surpasses the protective effect of progesterone. Alternatively, the presence of receptor-binding proteins in cells of cystic atretic follicles may result in a lower affinity of PR for progesterone than in vital tertiary follicles, so that apoptosis is induced despite the relatively high concentration of PR. Further studies are needed to confirm these statements.

The presence of PR in corpora lutea reflects the role of progesterone in corpus luteum activity (Revelli et al., 1996; Rueda et al., 2000). Progesterone regulates the proliferation and development of luteinized granulosa and theca cells in an autocrine and paracrine way (Sasano and Suzuki, 1997). The presence of PR in all lutein cells of the bovine corpora lutea suggests the influence of progesterone in the luteinization process (Revelli et al., 1996; Duffy et al., 1997). In the present study, however, the PR immunostaining in the corpus luteum was lower than in most other ovarian structures, which can be due to a negative effect of the locally produced high levels of progesterone to the PR production.
In contrast to all other ovarian cells, the lutein cells of the corpora lutea showed PR immunostaining not only in the nuclei, but in the cytoplasm as well. It is unlikely that this cytoplasmic staining is due to an artefact, because it was absent in all other ovarian cell types and in the negative controls. Therefore the idea is favoured that this cytoplasmic staining reflects the cycle-dependent localization of PR in the cytoplasm of this specific ovarian cell type. Whether this cytoplasmic PR reflects unbound and consequently inactive receptors in cells that produce high concentrations of progesterone remains to be elucidated.

A low but manifest PR immunoreactivity was observed in cells of the tunica albuginea and the surface epithelium. This corresponds with a study on ovine ovaries in which it has been suggested that cells of the ovarian surface epithelium are enzymatically involved in the ovulatory process by the influence of progesterone and its receptors (Murdoch, 1998). Further investigations in cattle are necessary to verify the role of PR and progesterone in the ovarian surface epithelium.

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CHAPTER 2

CELL DEATH IN THE CYCLIC BOVINE OVARY

Modified from:

CELL-SPECIFIC LOCALIZATION OF APOPTOSIS IN THE BOVINE OVARY AT DIFFERENT STAGES OF THE ESTROUS CYCLE D'Haeseleer M, Cocquyt G, Van Cruchten S, Simoens P, Van den Broeck W Theriogenology 2006; <u>65</u>: 757-772

Summary

Apoptosis was localized in all ovarian cell types of 23 cows in various stages of the estrous cycle, using the detection of active caspase-3, *in situ* end labelling (TUNEL) and DNA fluorescent staining (DAPI). Very few apoptotic cells were found in primordial, primary, secondary and vital tertiary follicles. In contrast, apoptosis in attetic tertiary follicles was much more frequent, and high apoptotic scores were recorded when using the TUNEL technique and lower scores with the caspase-3 assay. Cystic attetic follicles showed in general a higher apoptotic score than obliterative attetic follicles, with intermediate to high scores in granulosa cells and lower scores in theca cells. In corpora lutea, large and small lutein cells had intermediate to high scores using the caspase-3 assay, and intermediate to low scores using the TUNEL assay. Irrespective of the detection method, the scores were higher in lutein cells than in the capsular stroma cells. In all ovarian structures examined, variations in apoptotic scores were seen in the different cycle stages, suggesting a cycle-dependent influence on apoptosis, although correlations with plasma progesterone concentrations were low.

Introduction

Apoptosis in the mammalian ovary is involved in the mechanism of both follicular atresia and luteal regression. Follicular atresia is the phenomenon through which the ovary eliminates those follicles that are not selected for ovulation (Tilly et al., 1991; Yang and Rajamahendran, 2000a), while degeneration of corpora lutea is essential for maintaining the estrous or menstrual cycle (Amsterdam et al. 2003). In general, apoptotic cells are identified by cellular shrinking, ruffling of the plasma membrane, chromatin margination and condensation of the cytoplasm and nucleus.

Subsequently, the cells become fragmented and break up into apoptotic bodies (Van Cruchten et al., 2003). Ovarian apoptosis has already been documented abundantly in a number of species, viz. in human (Depalo et al., 2003), baboon (Kugu et al., 1998), mouse (Fenwick and Hurst, 2002), rat (Kaipia and Hsueh, 1997), sheep (Jolly et al., 1997), goat (Sharma, 2000), pig (Berardinelli et al., 2004) and cattle (Manikkam and Rajamahendran, 1997; Porter et al., 2001). Despite data concerning the frequency and cyclic changes of apoptosis in bovine ovarian follicles (Austin et al., 2001; Van Wezel et al., 1999; Zeuner et al., 2003) and corpora lutea (Zheng et al., 1994; Juengel et al., 1993), data concerning apoptosis in all different ovarian cell types are still incomplete. Therefore, in this study the presence of apoptosis in the bovine ovarian cell types was examined using three techniques, i.e., the detection of activated caspase-3, *in situ* DNA end labelling (TUNEL) and a DNA fluorescent staining (DAPI). Furthermore, putative differences between the apoptotic scores in active and inactive ovaries were studied. Finally, possible cyclic changes of apoptosis in the different cell types and their correlation with the plasma progesterone level were determined.

Materials and methods

Animals and samples

Samples of both ovaries and blood were obtained from 23 cows at the moment of slaughter at a local slaughterhouse. All animals had been used in our previous studies on the localization of progesterone receptor and estrogen receptors (D'Haeseleer et al., 2006b; 2006 c).

The ovarian tissue samples were fixed for 6 hours in 3.5 % phosphate-buffered formaldehyde (pH 6.7) at room temperature. Afterwards, the samples were embedded in paraffin and from each sample three sections (5 μ m thin) were cut, mounted on 3-aminopropyl-triethoxysilane-coated slides (APES, Sigma, St. Louis, MO, USA) and dried overnight at 37°C. These sections were used for TUNEL, caspase-3 assay and DAPI staining.

The stage of the estrous cycle of each animal was earlier determined (D'Haeseleer et al., 2006b). The cows were classified in 5 groups: 6 animals in oestrus (synchronized cows), 2 animals in metoestrus (slaughterhouse cows), 5 animals in early dioestrus (4 slaughterhouse cows and one synchronized cow), 5 animals in late dioestrus (slaughterhouse cows) and 5 animals in procestrus (slaughterhouse cows).

Analysis of DNA cleavage by TUNEL

DNA breaks occurring during the apoptotic process were detected by means of the TACS TM TdT kit (R&D systems Europe Ltd, Abingdon, UK) a TUNEL (TdT-mediated dUTP Nick-End Labelling) assay as described by Van Cruchten (2003). DAB chromogen substrate (Liquid DAB ⁺, DAKO, glostrup, Denmark) was administered for 5 min at room temperature, resulting in a brown staining. Finally, sections were counterstained with Mayer's haematoxylin for 10 sec and mounted. In every staining procedure positive and negative controls were included.

Immunohistochemical detection of active caspase-3

The caspase-3 technique is an immunohistochemical assay for the detection of the enzyme caspase-3 which can be activated during the apoptotic process and which in turn eventually activates endonucleases that cause the characteristic morphology of apoptotic cells (Van Cruchten et al., 2003). For the detection of activated caspase-3 a polyclonal rabbit-anti-mouse caspase-3 antibody AF835 and the Cell and Tissue Staining Kit HRP-AEC System (R&D Systems Europe Ltd, Abingdon, UK) were used. The protocol described by Van Cruchten (2003) was followed with slight modification. After incubation of the sections with

Streptavidin-HRP detection solution the slides were rinsed in PBS and 50 μ l 3-amino-9ethylcarbazole (AEC) chromogenic substrate solution was administered for 6 min at room temperature. The reaction of AEC with peroxidase resulted into a red coloured precipitate. Sections were counterstained with Mayer's haematoxylin for 10 sec and mounted with an aqueous mounting medium. In every staining procedure positive and negative controls were included.

DAPI-staining

Nuclear DNA fragmentation was assessed using a cell permeable fluorescent DNA binding probe, viz. 4', 6-Diamidino-2-Phenylindole (DAPI, Sigma-Aldrich Inc., St. Louis, USA). Therefore, the ovarian tissue sections were rehydrated, rinsed in PBS and incubated with DAPI (5µg/ml PBS) for 12 min at room temperature in a humidified chamber. After incubation, sections were rinsed in PBS, mounted in a specific mounting medium for fluorescence microscopy (Sigma-Aldrich Inc.) and stored at 4°C until observation using an Olympus BX61 fluorescence microscope (Olympus Belgium N.V., Aartselaar, Belgium).

Microscopic analysis

Apoptosis was examined by using all of the above-mentioned assays applied in serial tissue sections of every ovary. In each ovary, the different follicles, corpora hemorrhagica, corpora lutea, corpora albicantia, stroma cells, tunica albuginea and surface epithelium were evaluated. Follicles were classified as primordial, primary, secondary and tertiary follicles according to the criteria listed in the Nomina Histologica (1994). Tertiary follicles containing less than 1 % apoptotic cells were regarded as vital tertiary follicles, whereas follicles containing more than 1% apoptotic cells were classified as atretic. The latter were further classified as obliterative or cystic according to Dellmann and Eurell (1998).

For each cell type the presence of apoptosis demonstrated by the TUNEL assay and caspase-3 assay was determined by counting the cells presenting a brown nuclear staining or brown apoptotic bodies, and a red cytoplasmic staining, respectively. Using the DAPI technique strongly fluorescent apoptotic nuclei or fragmented nuclear bodies were counted. At least 100 cells of each cell type were counted on digital micrographs taken randomly at magnification 600 x with an Olympus DP 50 digital camera mounted on an Olympus BX61

light microscope (Olympus Belgium N.V., Aartselaar, Belgium). If too many cells were present in one micrograph, cells were counted in 4 grids of 2500 μ m² that were projected randomly over the micrograph.

For the TUNEL and caspase-3 assay staining intensity scores ranged from 0 (no staining), 1 (weak staining), 2 (moderate staining) to 3 (strong staining). Apoptosis was evaluated using an apoptotic score (AS) calculated according to the formula:

$$AS = 0 n_0 + 1 n_1 + 2 n_2 + 3 n_3$$

with n_0 , n_1 , n_2 and n_3 are the percentage of cells exhibiting a staining intensity 0, 1, 2 and 3, respectively (Boos et al., 1996). For the DAPI staining procedure, the apoptotic score indicated the mean percentage of apoptotic cells for each cell type.

Statistical analysis

Throughout the study, results were represented as means and variation was expressed as standard error of the mean (SEM). Possible differences in the AS between active and inactive ovaries, cystic and obliterative atretic follicles, and between the various cell types were analysed using the Kruskall-Wallis test. This test was also used to compare the apoptotic scores in the 5 stages of the estrous cycle. The Tukey's test was used to determine between which particular cycle stages significant differences were present (P < 0.05, P < 0.01).

The Spearman rank correlation test was used for determination of correlations between the plasma progesterone levels and the apoptotic score (Table 1). Statistical analyses were performed using SPSS 12.0 (SPSS Inc., Chicago, Illinois, USA).

		TUNEL	caspase-3	DAPI
Secondary Follicle	follicle cells	-	-	- 0,20
	theca follicularis	-	-	-
Vital Tertiary Follicle	granulosa cells	- 0,12	+ 0,01	- 0,29
	theca interna	- 0,43*	+ 0,01	+ 0,23
	theca externa	- 0,26	-	+ 0,03
Obliterative Atretic Follicle	granulosa cells	+ 0,20	+ 0,01	+ 0,13
	theca interna	+ 0,06	+ 0,01	- 0,14
	theca externa	- 0,03	-	- 0,15
Cystic Atretic Follicle	granulosa cells	- 0,14	+ 0,34	- 0,18
	theca interna	- 0,56	- 0,10	+ 0,22
	theca externa	+ 0,42	- 0,03	+ 0,20
Corpus Hemorrhagicum	granulosa cells	+ 0,80	+ 0,95*	+ 0,16
	theca interna	+ 0,36	-	-
	theca externa	+ 0,00	-	-
Corpus Luteum	large lutein cells	- 0,39	+ 0,20	- 0,23
	small lutein cells	- 0,58**	+ 0,18	- 0,04
	perivascular stroma cells	- 0,20	- 0,24	+ 0,06
	capsular stroma cells	+ 0,13	- 0,32	+ 0,04
Corpus Albicans	internal stroma	+ 0,01	- 0,21	+ 0,03
	capsular stroma cells	+ 0,09	- 0,25	+ 0,16
Deep Stroma		- 0,31	+ 0,02	+ 0,03
Superficial Stroma		- 0,32	-	+ 0,15
Tunica Albuginea		- 0,13	- 0,35	- 0,26
Surface Epithelium		- 0,42*	_	- 0,52*

Table 1: Correlations between the plasma progesterone levels and apoptotic scores detected by various techniques (TUNEL, caspase-3 and DAPI) in the different ovarian cell types. "-" no correlation; * P < 0.05; ** P < 0.01.

Results

Localisation of apoptosis in the bovine ovary

In all animals TUNEL and caspase-3 positive cells were completely absent in primordial, primary and secondary follicles (Figs. 1a, 1b, 1c). DAPI-staining showed a small amount of apoptotic cells in the secondary follicles. In vital tertiary follicles low levels of apoptotic cells were seen for the three assays. In contrast, apoptosis was obvious in atretic tertiary follicles. Apoptotic cells were also noticed in corpora hemorrhagica, corpora lutea and corpora albicantia. The deep stroma, superficial stroma, surface epithelium and tunica albuginea also contained apoptotic cells.



Fig. 1: Mean scores of apoptosis using the TUNEL (Fig. 1a), caspase-3 (Fig. 1b) and DAPI (Fig. 1c) assay in the different groups of bovine ovarian cell types. Within the different cell groups, apoptotic scores are significantly different (P < 0.05) when indicated by different superscripts.

PdF: primordial follicle, PF: primary follicle, SF: secondary follicle, VTF: vital tertiary follicle, OAF: obliterative atretic follicle, CAF: cystic atretic follicle, CH: corpus hemorrhagicum, CL: corpus luteum, CA: corpus albicans, DS deep stroma, SS: superficial stroma, TA: tunica albuginea, SE: surface epithelium, fc: follicle cells, tf: theca follicularis, gc: granulosa cells, ti: theca interna, te: theca externa, ll: large lutein cells, sl: small lutein cells, ps: perivascular stroma, is: internal stroma, cs: capsular stroma.

Cell-specific variation

Many differences were observed in the staining scores of the different ovarian cell types (Figs. 1a, 1b, 1c). In vital tertiary follicles low apoptotic levels were found in granulosa and theca cells. In contrast, apoptosis was clearly present in cystic and obliterative atretic tertiary follicles with a high apoptotic score in the granulosa cells, an intermediate score in the theca interna and a low score in the theca externa cells. These differences were statistically significant (P < 0.01). Comparison between cystic and obliterative atretic follicles revealed that cystic atretic follicles showed in general higher apoptotic scores for all cell types, irrespective of the staining procedure.

Many variations in apoptotic levels were found in the corpora hemorrhagica, corpora lutea and corpora albicantia. In corpora hemorrhagica the TUNEL assay revealed high to intermediate apoptotic scores with significant differences between the granulosa and theca cells (P < 0.05), whereas the caspase-3 assay and DAPI staining revealed a lower apoptotic score in granulosa cells and no apoptotic score in theca cells. In contrast, the large and small lutein cells of corpora lutea showed high scores using the caspase-3 assay, and lower scores using the TUNEL assay. Apoptosis in capsular stroma cells of corpora lutea was only demonstrated by TUNEL assay and DAPI staining. Furthermore, all assays revealed a significantly higher apoptotic score in lutein cells than in stroma cells (P < 0.05). In corpora albicantia, low to intermediate scores were seen irrespective of the staining protocol. Comparing corpora hemorrhagica, corpora lutea and corpora albicantia with each other, higher scores were observed in corpora lutea, apart from the very high value found in the granulosa cells of the corpora hemorrhagica when using the TUNEL assay. Deep stroma cells showed a low apoptotic score, comparable with capsular stroma cells of corpora lutea and theca externa cells of atretic tertiary follicles. In the superficial stroma and tunica albuginea, the apoptotic score was significantly lower than in the deep stroma for all assays (P < 0.01). Intermediate apoptotic scores were seen in the surface epithelium following TUNEL assay.

Comparison between the different follicular stages revealed a significant increase in apoptotic score for follicle/granulosa cells from secondary to vital tertiary to atretic tertiary follicles (P < 0.05).

Ovary-specific variation

Comparison between the active and inactive ovaries revealed no significant differences (Figs. 2a, 2b, 2c), although, the DAPI-staining showed more apoptosis in the active ovary than in the inactive ovary for almost all cell types.



Fig. 2: Mean scores of apoptosis in the different cell types of the active and inactive bovine ovary using the TUNEL (Fig. 2a), caspase-3 (Fig. 2b) and DAPI (Fig. 2c) assay. Abbreviations: See fig. 1.

Cycle-specific variation

Variations in apoptotic score, concomitant with the bovine cycle stages, were seen in the different ovarian structures and cell types (Figs. 3a, 3b, 3c).

In primordial, primary and secondary follicles no detectable expression for apoptosis was observed in any of the cycle stages. However, DAPI clearly showed the presence of a few apoptotic cells in secondary follicles during metoestrus and early dioestrus.

In vital tertiary follicles, the caspase-3 assay showed a similar expression for all cycle stages. Using the TUNEL assay a higher apoptotic score was seen in oestrus than in the other cycle stages, while the DAPI staining showed intermediate scores in oestrus, higher scores in early dioestrus and intermediate to low scores in late dioestrus and prooestrus. In all different cell types a similar cyclic pattern was observed. In attretic tertiary follicles high variations in apoptotic levels were found between the different cycle stages. The apoptotic scores of all cell types of obliterative attretic follicles showed a similar cyclic pattern with values that were intermediate to high during oestrus and metoestrus, intermediate during dioestrus, and high again during prooestrus. In granulosa cells of cystic attretic follicles the apoptotic scores were high in oestrus, low in metoestrus, high in early dioestrus and intermediate during the subsequent stages of dioestrus and prooestrus. A similar cyclic variation was observed in the theca interna cells. Using the caspase-3 assay, clear cycle-dependent variations were only seen in cystic attretic follicles, with an intermediate score during oestrus, a low score during metoestrus, intermediate score again during dioestrus, and the highest score during prooestrus.

Corpora hemorrhagica were only present in oestrus and early dioestrus. The granulosa cells of the corpora hemorrhagica always showed a higher apoptotic score in early dioestrus than in oestrus.







Fig. 3: Mean apoptotic scores measured by the TUNEL (Fig. 3a), caspase-3 (Fig. 3b) and DAPI (Fig. 3c) assay in various bovine ovarian cell types during the different oestrous cycle stages. Abbreviations: See fig. 1. Significant differences between the cycle stages observed in the cystic follicles using the caspase-3 assay are indicated by different superscripts.

In both the large and small lutein cells of corpora lutea, the highest apoptotic scores following the TUNEL assay were seen in oestrus and metoestrus. These scores decreased in early dioestrus and the subsequent period of dioestrus, and increased again in prooestrus. A nearly similar pattern was observed in the stroma cells of corpora lutea. The caspase-3 assay showed a different pattern, with the lowest apoptotic score in oestrus, higher scores in metoestrus and both dioestrus stages, and the highest score in prooestrus. However, in stroma cells the scores were extremely low in all stages. Using the DAPI staining, little variation was seen in lutein cells, with scores that were relatively high in oestrus, lower in metoestrus and both dioestrus stages, and high again in prooestrus. In stroma cells of corpora lutea, the scores were low during oestrus, high during metoestrus, low again during both dioestrus stages, and minimal during prooestrus.

In corpora albicantia, the apoptotic scores obtained by the TUNEL technique were intermediate in oestrus, decreased in metoestrus to increase again in the next stages. According to the caspase-3 assay and the DAPI staining the scores were low to intermediate, with small differences between the various stages.

The apoptotic scores of deep stroma cells were low at oestrus, high in metoestrus, and low again during the following stages. Using the TUNEL assay, apoptosis in superficial stroma cells and tunica albuginea cells was clearly demonstrable during oestrus, and in surface epithelial cells also at oestrus and during metoestrus (Fig. 4).

In addition to the cycle-specific variation, the correlations between plasma progesterone levels and apoptotic scores were examined (Table 1). Positive correlations were found for the granulosa cells of corpora hemorrhagica after using the TUNEL technique (0.80) and caspase-3 assay (0.95, P < 0.05). Remarkably, the correlation with TUNEL scores in lutein cells of corpora lutea was negative (-0.39 for large lutein cells and -0.58 (P < 0.01) for small lutein cells) whereas the correlation was slightly positive following the caspase-3 assay (0.20 for large and 0.18 for small lutein cells). In the surface epithelium, a significant and faintly negative correlation was seen between the plasma progesterone levels and the apoptotic scores determined by the TUNEL assay (-0.24, P < 0.05) and DAPI staining (-0.52, P < 0.05).











Fig. 4: Apoptosis detection (arrows) using TUNEL assay (**Fig. 4a-d**), immunohistochemical detection of caspase-3 (**Fig. 4e-f**) and DAPI staining (**Fig. 4g-h**) in the different bovine ovarian cell types. **Fig. 4a**: Cystic tertiary follicle at early dioestrus with numerous TUNEL positive cells in the granulosa layer (1), the cells of the cumulus oophorus (2) and the theca interna layer (3). Bar = 50 μ m. **Fig. 4b**: Overview of an obliterative attretic tertiary follicle at oestrus showing TUNEL positive cells in the collapsed theca layers (4). Bar = 100 μ m. **Fig. 4c**: TUNEL positive cells in the surface epithelium (5) and tunica albuginea (6) during early dioestrus; superficial stroma (7) and primary follicles (8) are negative. Bar = 50 μ m. **Fig. 4d**: TUNEL positivity in lutein cells of the corpus luteum (9) at oestrus. Bar = 50 μ m. **Fig. 4e**: Cystic tertiary follicle at oestrus with numerous caspase-3 positive cells in the granulosa layer (10) and lower numbers in the theca layers (11). Bar = 50 μ m. **Fig. 4f**: Overview of a corpus luteum during prooestrus with high numbers of caspase-3 positive lutein cells (12). Bar = 500 μ m. **Fig. 4g**: Cystic attetic follicle during prooestrus with high numbers of apoptotic cells in the granulosa (13), moderate levels in the theca interna (14) and low levels in the theca externa (15). DAPI staining, bar = 50 μ m. **Fig. 4h**: Overview of an obliterative attetic tertiary follicle during prooestrus with apoptotic cells (arrows) in the collapsed theca layers (16). DAPI staining, bar = 100 μ m.

Discussion

Literature reports concerning cell death and the use of techniques for apoptosis detection in the bovine ovary contain many apparent contradictions (Van Wezel et al., 1999; Hagemann et al., 1999). Therefore, in cell death research it is very important to detect and characterize apoptosis by multiple methods. In the present study, apoptosis in various cell types of the bovine ovary at the different stages of the estrous cycle was demonstrated by using three techniques, viz., the detection of active caspase-3, *in situ* end labelling (TUNEL) and DNA fluorescence staining (DAPI).

Our results show that in general more significant differences in apoptotic scores were noticed when using TUNEL-assay than caspase-3 assay. This discrepancy can be explained by the fact that the used assays detect different aspects in the apoptotic process. The caspase-3 assay detects activated caspase-3, which is present at the onset of the apoptotic process, whereas the TUNEL-assay and DAPI-staining demonstrate DNA fragmentation and chromatin condensation, respectively, which are only visible at later stages of the apoptotic process. Furthermore, the number of TUNEL positive cells was in general higher than the number of caspase-3 positive cells. This can be due to the fact that DNA fragmentation remains present and detectable for a longer period of time than activated caspase-3. However, it should be noticed that the number of TUNEL positive cells could also have been

overestimated, due to the fact that it is not always clear whether a cluster of apoptotic bodies originated from a single cell or from different cells. Furthermore, DNA fragmentation is not specific for apoptosis, and oncotic cells can be misinterpreted as apoptotic cells (Van Cruchten et al., 2003). In corpora lutea, however, the number of caspase-3 positive cells was two times higher than the number of TUNEL positive cells. This finding might indicate that many caspase-3 positive cells disappear, probably by phagocytosis, before DNA fragmentation occurs.

The present study demonstrates that apoptotic cells were extremely scarce in primordial, primary and secondary follicles and very abundant in tertiary follicles in all cows at the different cycle stages. Although it has been described that atresia may begin at any stage of development of the follicle (Bloom and Fawcett, 1975), our findings would suggest that atresia of tertiary follicles is of major importance for the reduction of the thousands of follicles that will never ovulate. The very low frequency of atresia in the earlier follicular stages has also been demonstrated in the human ovary in a recent immunohistochemical study using TUNEL and active caspase-3 detection (Glamoclija et al., 2005). A possible explanation for this observation could be that apoptosis in early stage follicles is caused by a mechanism in which caspase-3 is not involved and which is not detectable by the TUNEL assay. A more plausible reason might be the fact that for each cow in our study only a single snapshot evaluation of ovarian structures was made. As follicular atresia starts already before birth and continues for many years, the chance to detect atretic primordial and primary follicles is clearly reduced when only a few histological sections are examined.

The finding of low apoptotic levels in secondary follicles is in line with a recent study on apoptosis in human ovarian follicles, which demonstrates that, once the primary follicle develops to a secondary follicle, the follicle cells of secondary follicles showed Bcl-2 expression and consequently were protected against apoptosis (Depalo et al., 2003). Another study in rat ovaries demonstrated that apoptosis in the granulosa cells of early developing follicles is prevented by progesterone (Peluso, 2003). However, in the present study no correlations were found between apoptotic scores in secondary follicles and the plasma progesterone concentration, indicating that progesterone acts via paracrine rather than via endocrine route. In order to elucidate the putative influence of progesterone on secondary

follicles, the localisation of progesterone-specific receptors in ovarian cells needs to be investigated.

High numbers of atretic tertiary follicles were observed in all bovine ovaries at the different cycle stages. This corresponds with the physiologic process of the bovine estrous cycle in which only one tertiary follicle is selected for ovulation, while the remaining tertiary follicles regress. In our study, the number of apoptotic cells in the different layers of atretic tertiary follicles showed significant differences. These findings correspond with studies of ovarian apoptosis in rats (Vickers et al., 2000), pigs (Tilly et al., 1992) and chicken (Tilly et al., 1991). Cystic atretic follicles show higher apoptotic levels compared to obliterative follicles, which can be due to their different morphology. Disruption of the intercellular contacts in cystic atretic follicles may stimulate apoptosis (Zeuner et al., 2003). In contrast, in obliterative follicles the granulosa and theca layers both collapse, while the cell-to-cell contacts remain intact and the intercellular integrity inhibits the induction of apoptosis. Therefore, apoptosis in obliterative tertiary follicles would be the result of morphological changes during atresia, rather than the cause of follicular atresia. However, this hypothesis is highly speculative and should be confirmed.

In many ovaries, cystic follicles with high apoptotic scores were located nearby cystic follicles with low scores, suggesting that morphologically identical follicles showed a high variability in apoptosis. In an earlier study we also found a variable expression of estrogen receptor beta mRNA levels in neighbouring bovine follicles that were morphologically identical (D'Haeseleer et al., 2005). These observations suggest that ovarian follicles are mainly influenced by local factors produced by adjacent follicles or other ovarian cells, rather than by systemic endocrine factors.

Earlier studies indicated that high progesterone levels trigger apoptosis in non-ovulatory tertiary follicles during the bovine estrous cycle (Taylor et al., 1993; Yang and Rajamahendran, 2000b). A similar observation was made in granulosa cells of obliterative follicles in which the TUNEL assay demonstrated high apoptotic levels in early dioestrus, when plasma progesterone concentrations are increasing. However, the apoptotic scores in cystic follicles were always lower in early dioestrus, than in the other stages. Furthermore, no significant correlations were found between apoptosis in tertiary follicles and plasma

progesterone concentration. This suggests that the influence of progesterone on apoptosis in tertiary follicles is complex and that other factors may also be involved.

In lutein cells, the apoptotic score detected by the caspase-3 assay increased from low levels at oestrus to maximal levels at dioestrus / prooestrus, whereas the opposite was seen using the TUNEL assay. This is in accordance with the positive and negative correlations observed between the plasma progesterone concentrations and the caspase-3 and TUNEL scores, respectively. This observation indicates that progesterone induces apoptosis in corpora lutea by caspase-3 activation, whereas the DNA fragmentation occurs in a later stage, explaining the negative correlation. This is in contrast with the results obtained from an *in vitro* study on progesterone treated cultured bovine lutein cells, in which it was demonstrated that progesterone prevents apoptosis of lutein cells (Rueda et al., 2000).

In general, apoptotic scores were low in cells of the deep stroma, superficial stroma, tunica albuginea and surface epithelium with relative high levels during metoestrus. This indicates that ovulation might have an influence on the regression of stroma and epithelial cells but data to elucidate this process are still lacking.

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PROLIFERATION PATTERNS IN THE BOVINE OVARY AT DIFFERENT STAGES OF THE ESTROUS CYCLE

Modified from:

PROLIFERATION PATTERNS IN THE BOVINE OVARY AT DIFFERENT STAGES OF THE ESTROUS CYCLE D'Haeseleer M, Simoens P, Van den Broeck W In preparation

Summary

Proliferation was examined in the ovaries of 23 cows at different stages of the estrous cycle using immunohistochemical detection of Ki-67. Ki-67 was not observed in primordial follicles. In primary and secondary follicles the score for Ki-67 increased progressively with the maturation of the follicle. In vital tertiary and cystic atretic follicles high to moderate scores for Ki-67 were found, whereas in obliterative atretic follicles the score was much lower. Scores were high in all cell types of corpora hemorrhagica and in the lutein cells of the corpora lutea, moderate in the stroma cells of the corpora lutea and low in the cells of the corpora albicantia. Low Ki-67 scores were also found in the deep and superficial stroma.

In vital tertiary and cystic atretic follicles the scores for Ki-67 were high at oestrus, decreased during metoestrus, increased during dioestrus to decrease again during procestrus. In corpora lutea, corpora albicantia and deep stroma a different cyclic pattern was observed with high scores during metoestrus and lower scores during dioestrus and procestrus. These variations suggest a cell-specific and cycle-dependent influence on cell proliferation, although correlations with the plasma progesterone concentrations were low.

Introduction

During the bovine estrous cycle, the ovary is subjected to extensive tissue remodelling that consists of sequential phases of cell proliferation and cell death in the different ovarian structures such as the follicles, the corpus hemorrhagic, the corpus luteum, the corpus albicans, the stroma and the surface epithelium. Cell proliferation is morphological and physiological event that is controlled by a combination of circulating steroid hormones and locally produced growth factors (Quirk et al., 2004).

Cell proliferation in the ovary has been studied in a number of species such as human (Funayama et al., 1996), the baboon (Wandji et al., 1997), the marmoset (Young et al., 2000), the horse (Al-zi'abi et al., 2003), the ewe (Campbell et al., 2004; Jablonka-Shariff et al., 1993; Jolly et al., 1997), the porcine (May et al., 1992), the dog (Spanel-Borowski a, b et al., 1984), the rat (Gaytan et al., 1996; Richards et al., 1986; Uilenbroek et al., 1979; Oktay et al., 1995) and the mouse (Britt et al., 2004). Many papers have been published on cell proliferation in bovine ovarian follicles (Tajima et al., 2002; Kotsuyi et al., 1990; Isobe et al., 2000; van Wezel, 1999) and in corpora lutea (Boos, 1998; Zheng et al., 1994).

However, the cyclic proliferation patterns in the different bovine ovarian cell types are not well known. Therefore, the aim of the present study was to determine proliferation in the different cell types of the bovine ovary at the different stages of the estrous cycle by using immunohistochemical detection of Ki-67. Ki-67 is a nuclear antigen expressed by proliferating cells and is occurring in all active parts of the cellular cycle, except in the resting phase (G0) (Scholzen et al., 2000). Furthermore, putative differences between the scores of Ki-67 in active and inactive ovaries were also examined. Finally, the correlation between the plasma progesterone level and proliferation in the different ovarian cell types was determined.

Materials and methods

Animals and samples

Samples of both ovaries and blood were obtained from 23 cows at the moment of slaughter at a local slaughterhouse. All animals had been used in our previous study on apoptosis in the bovine ovary (D'Haeseleer et al., 2005).

For immunohistochemical determination of PR the ovarian tissue samples were fixed for 6 h in 3.5% phosphate-buffered formaldehyde (pH 6.7) at room temperature and dehydrated in increasing concentrations of ethanol. Subsequently, all tissue samples were embedded in paraffin and 5 μ m thin sections were cut. The sections were mounted on 3-aminopropyl-triethoxysilane-coated slides (APES, Sigma, St. Louis, MO, USA) and dried overnight at 37°C.

The stage of the estrous cycle of each animal was earlier determined (D'Haeseleer et al., 2006b). The cows were classified in 5 groups: 6 animals in oestrus (synchronized cows), 2 animals in metoestrus (slaughterhouse cows), 5 animals in early dioestrus (4 slaughterhouse cows and one synchronized cow), 5 animals in late dioestrus (slaughterhouse cows) and 5 animals in procestrus (slaughterhouse cows).

Immunohistochemical detection of Ki-67

For the detection of Ki-67, the assay described by Van Cruchten et al. (2004) was used. The primary antibody was a monoclonal mouse-anti-human NCL-Ki-67-MM1 antibody (Novocastra Laboratories Ltd., Newcastle upon Tyne, UK) which cross-reacts with the bovine Ki-67. Rabbit anti-mouse antibodies conjugated to biotin (DAKO A/S, Glostrup, Denmark) were used as secondary antibodies and peroxidase-conjugated streptavidine (DAKO A/S, Glostrup, Denmark) as enzyme label. DAB (Liquid DAB+, DAKO, Prosan, Merelbeke, Belgium) was used as chromogen substrate, resulting in a brown nuclear precipitate. Mayer's haematoxylin was applied as a nuclear counterstain.

Positive and negative controls were included in every staining procedure. Canine uterine tissues known to be positive for Ki-67 served as positive controls. As negative controls tissue

sections incubated with dilution buffer instead of the primary antibody, secondary antibody, and/or peroxidase-conjugated streptavidine were used.

Microscopic analysis

In each ovary the different follicle types, corpora hemorrhagica, corpora lutea, corpora albicantia, deep stroma, superficial stroma, tunica albuginea and surface epithelium were microscopically evaluated. All follicles were examined and classified by histological stage into primordial, primary, secondary and tertiary follicles according to the criteria listed in the Nomina Histologica (1994). Tertiary follicles containing less than 1 % apoptotic cells were classified as vital tertiary follicles, whereas follicles containing more than 1% apoptotic cells were regarded as atretic. The latter were further classified as obliterative or cystic according to the criteria described by Dellmann and Eurell (1998). For each cell type, the presence of Ki-67 was assessed by counting the cells presenting a brown nuclear staining.

At least 100 cells of every cell type were counted on digital micrographs taken randomly at magnification 600 x with an Olympus DP 50 digital camera mounted on an Olympus BX61 light microscope (Olympus Belgium N.V., Aartselaar, Belgium). If too many cells were present in one micrograph, cells were counted in 4 grids of 2500 μ m² that were projected randomly over the micrograph. Staining intensity scores ranged from 0 (no staining), 1 (weak staining), 2 (moderate staining) to 3 (strong staining). The presence of Ki-67 was evaluated using a score S(Ki-67) which was calculated according to the formula:

 $S(Ki-67) = 0 n_0 + 1 n_1 + 2 n_2 + 3 n_3$ with n_0 , n_1 , n_2 and n_3 are the percentage of cells exhibiting a staining intensity 0, 1, 2 and 3, respectively (Boos et al., 1996).

Statistical analysis

Throughout the study, results were represented as mean values and variation was expressed as standard error of the mean (SEM). Possible differences between active and inactive ovaries, between the various cell types and between the 5 stages of the estrous cycle were analysed by means of the Kruskall-Wallis test. The Tukey's test was used to determine significant (P < 0,05) or highly significant (P < 0,01) differences between particular cycle stages. The Spearman rank correlation test was used for determination of correlations between the plasma progesterone levels and the scores for Ki-67 (Table 1). Statistical analyses were performed with procedures available in SPSS 12.0 (SPSS Inc. Headquarters, Chicago, Illinois, USA).

Results

Localization of Ki-67 in the bovine ovary (Fig. 1a)

Ki-67 was not detected in primordial follicles, tunica albuginea and surface epithelium but was observed in low amounts in primary and secondary follicles, obliterative atretic follicles, corpora albicantia, deep and superficial stroma. The expression of Ki-67 was high in vital tertiary follicles, cystic atretic follicles, corpora hemorrhagica and corpora lutea (Fig. 1a).

Cell-specific variation of the Ki-67 expression (Fig. 1a)

Many differences were observed in the staining scores of Ki-67 between the different ovarian cell types. The Ki-67 score in the follicle cells of primary follicles was low but increased in secondary follicles (p < 0.05) to reach high numbers in vital tertiary follicles (p < 0.01). The score in the granulosa cells of obliterative atretic follicles was very low, while in the granulosa cells of cystic atretic tertiary follicles a very high score was observed, similar to that of the granulosa cells of vital tertiary follicles (p < 0.01). Granulosa cells of corpora hemorrhagica showed a very high Ki-67 score and this score slowly decreased in both the large and small lutein cell of corpora lutea and reached the lowest level in the internal stroma cells of corpora albicantia.

The Ki-67 score in theca cells was always lower than that in the follicle / granulosa cells of the corresponding structures, and this difference was significant in secondary follicles (P < 0.01), vital tertiary follicles (P < 0.05) and cystic atretic tertiary follicles (p < 0.01). Furthermore, theca externa cells always showed a lower Ki-67 score compared to theca interna cells.

Cells of the deep stroma and superficial stroma cells showed a lower Ki-67 expression, similar to that of the theca cells in secondary follicles.
Ovary-specific variation of the Ki-67 expression

Comparison between active and inactive ovaries revealed that the Ki-67 expression was higher in the ovarian structures of the active than in the inactive ovaries, except in the cystic atretic tertiary follicles and in the corpora hemorrhagica where higher scores in the inactive ovaries were found. Significant differences were observed in the large lutein cells (P < 0.05) and capsular stroma cells (P < 0.05) of corpora lutea (Fig. 1b).



Cycle-specific variation of the Ki-67 expression

High variations in the Ki-67 expression, concomitant with the bovine cycle stages, were observed in the different ovarian cell types. However these differences were never significant (Fig. 1c).

In primordial follicles Ki-67 expression was not detected in any of the cycle stages. In the follicle cells of primary follicles the Ki-67 expression was low during oestrus, decreased during metoestrus and early dioestrus and was not longer detectable at late dioestrus and prooestrus.

In secondary follicles, the follicle cells showed low levels during oestrus which slightly increased from metoestrus to late dioestrus and further decreased during procestrus. In the theca layers of latter follicles the expression was low and slightly changed during the subsequent stages. In the granulosa layer of vital tertiary follicles the Ki-67 expression was high in oestrus, slightly decreased during metoestrus, increased again during dioestrus to decrease again in procestrus. In the theca layers lower but similar cyclic patterns were observed. In cystic atretic tertiary follicles the Ki-67 expression pattern was nearly similar to that of vital tertiary follicles. However, in obliterative atretic tertiary follicles, the Ki-67 expression was very low and slightly differed between the cyclic stages.

The corpora hemorrhagica, which were only visible during oestrus, metoestrus and early dioestrus showed moderate to high Ki-67 expression with the highest expression during early dioestrus in the granulosa layer and during metoestrus in the theca layers.

In both the large and small lutein cells of corpora lutea, Ki-67 expression was moderate during oestrus, increased during metoestrus and early dioestrus and decreased during the other subsequent stages. A similar pattern with lower Ki-67 expression was observed in the stroma cells of the corpora lutea, corpora albicantia and in the deep stroma cells. In the superficial stroma, the Ki-67 expression was moderate during oestrus and low in the subsequent cycle stages whereas in the tunica albuginea and surface epithelium no Ki-67 expression was detected.

		Plasma Progesterone Level
Secondary Follicle	follicle cells	- 0,35
	theca follicularis	- 0,47*
Vital Tertiary Follicle	granulosa cells	+ 0,13
	theca interna	+ 0,30
	theca externa	+0,28
Obliterative Atretic Follicle	granulosa cells	- 0,40
	theca interna	- 0,31
	theca externa	- 0,44*
Cystic Atretic Follicle	granulosa cells	- 0,11
	theca interna	- 0,12
	theca externa	+ 0,19
Corpus Hemorrhagicum	granulosa cells	- 0,07
	theca interna	+0,18
	theca externa	+0,25
Corpus Luteum	large lutein cells	- 0,09
	small lutein cells	- 0,39
	perivascular stroma cells	- 0,27
	capsular stroma cells	- 0,53*
Corpus Albicans	internal stroma cells	- 0,37
	capsular stroma cells	- 0,66**
Deep Stroma		- 0,10
Superficial Stroma		- 0,72**
Tunica Albuginea		/
Surface Epithelium		/

Table 1: Correlations between the plasma progesterone levels and the Ki-67 scores in the differentovarian cell types. "/" not detected; * P < 0.05; ** P < 0.01



Discussion

In the present study, proliferation in various bovine ovarian cell types during the different stages of the estrous cycle was examined using immunohistochemical detection of Ki-67. Ki-67 was not observed in the primordial follicles, while increasing numbers of Ki-67 were found in primary, secondary and tertiary follicles. These findings correspond with earlier data on bovine (Kotsuji et al., 1990) and with observations in man (Funayama et al., 2005) and rat (Oktay et al., 1995) and implicates that the primordial follicles are quiescent. The transformation of primordial follicles from quiescence to growth is important in early folliculogenesis (Funayama et al., 1996). It is of evidence that the significant increase in the Ki-67 score of follicle/granulosa cells in primary, secondary and tertiary follicles reflects the high number of cells in growing follicles.

Theca cells show a significant increase in secondary and tertiary follicles. These findings are in accordance with a study on canine ovary and may indicate that thecal cell proliferation is necessary for the development of vital follicles (Spanel-Borowski et al., 1974a).

Two proliferation peaks were seen in the vital tertiary follicles and in the cystic atretic follicles during the estrous cycle. The first peak was observed during oestrus. Spannel-Borowski (1974b) who found a similar observation in dog ovarian follicles ascribed this peak to the synergetic action of both follicle stimulating hormone (FSH) and estrogens. It cannot be excluded that the proliferative activity is higher during procestrus because activation by follicle stimulating hormone (FSH) starts at procestrus.

The second proliferation peak, which is seen more or less in all ovarian structures except obliterative atretic follicles, develops during metoestrus and early dioestrus to reach a high level during late dioestrus. This proliferation activity may be ascribed to the increasing plasma progesterone levels. It is most probable that rising levels of progesterone stimulated the above-mentioned cell types sensitive for both estrogens and progesterone. The capacity of progesterone to stimulate cell proliferation in cooperation with estrogens is distinctly shown by studies in mouse (Bresciani et al., 1968) and dog (Spanel-Borowski et al., 1984a).

In our study the proliferative activity in cystic atretic follicles was nearly similar to that in vital tertiary follicles. A similar observation was made in a study on human ovaries (Funayama et al., 2006). These observations may suggest that in atretic tertiary follicles the

effect of other factors, namely factors involved in cell turnover rather than cell proliferation may be involved.

The corpus luteum undergoes extensive proliferation starting from metoestrus to late dioestrus while this proliferating level decreased during procestrus. Similar results were obtained by studies in man (Funayama et al., 1996) and rat (Gaytan et al., 1996). These data implicate that cells in the corpus luteum proliferate during their luteal phase, when the cells actively produce and secrete hormones (Juengel et al., 1993). Furthermore, the present study demonstrates that cell proliferation in the luteal phase of the corpus luteum is associated with the circulating plasma progesterone levels.

Moreover, the low proliferation rate in the corpus luteum during procestrus continued in the cells of the corpora albicantia. This finding corresponds with earlier data on cell proliferation in bovine luteal tissues of different reproduction states by Boos (1998), who found that these proliferation was high in young corpora lutea and markedly reduced in the corpora lutea regression stages. Cell proliferation in the bovine corpus luteum was therefore limited to one cycle, whereas, morphological involutions takes several cycles (Zheng et al., 1994).

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GENERAL DISCUSSION

This study was undertaken to obtain a better insight into the role of estrogen receptors and progesterone receptors in the different cell types of the bovine ovary, and to verify their relation to cell proliferation and apoptosis throughout the estrous cycle of normal cycling cows.

The first aim of this thesis was to examine the morphological localization of the estrogen receptors (ER α and ER β) and the progesterone receptor in the different cell types of both ovaries of 23 cows. For the detection of ER α and PR, monoclonal antibodies were used, based on previous studies (Van den Broeck et al., 2002a, 2002b) demonstrating that these antibodies crossreact with bovine tissue. In contrast, for ER β neither monoclonal nor polyclonal antibodies useful for bovine studies are readily available and therefore, in a preliminary study, a probe for *in situ* hybridisation to localize ER β mRNA was generated.

The second aim of the present study was to examine whether ER and PR were differentially localized throughout the estrous cycle. Therefore, quantitative differences in the immunostaining for ER and PR were analysed during five cycle stages of the bovine estrous cycle, namely procestrus, cerly metoestrus, late metoestrus and dioestrus. Furthermore, a putative relation between the presence of the steroid receptors and the cellular dynamics was studied. To achieve this, apoptosis and cell proliferation were quantified in all different ovarian cell types during the five mentioned cycle stages. Apoptosis was demonstrated by the use of three techniques which detect different aspects in the apoptotic process, whereas the TUNEL-assay and the DAPI staining demonstrate DNA fragmentation and chromatin condensation, respectively, which are only visible at later stages of the apoptotic process. The proliferative activity in the bovine ovarian cell types was assessed by immunohistochemical detection of the Ki-67 proliferation associated antigen.

In primordial follicles ER α was completely absent, while moderate levels of ER β mRNA and PR were detected. These findings are in accordance with other studies in cattle and indicate that both estrogen and progesterone play a role in the early process of folliculogenesis (Hulshof et al., 1995; Van den Broeck et al., 2002b). Alternatively, the presence of ER and PR in the follicle cells of primordial follicles may be necessary to keep the primordial follicle in the resting state. However, the exact influence of the steroid hormones during initial recruitment remains to be elucidated.

No apoptosis was detected in primordial follicles. This finding however is in contrast with earlier observations in human, rat and mouse ovaries (McGee and Hsueh, 2000; Depalo et al., 2003), showing that the number of primordial follicles decreases over time due to apoptosis of some of the follicles. This was confirmed by radioactive decay models of rats and mice, which proposed a constant loss of a fixed number of primordial follicles from the resting pool over time (Edwards et al., 1977). Furthermore, studies in humans showed that during lifetime, the continuing recruitment of primordial follicles leads to a gradual decrease of the number of these follicles. Additionally, an accelerated loss of follicles was seen during initial recruitment in humans (McGee and Hsueh, 2000; Gougeon et al., 1994) while a decreasing loss of follicles with increasing ages was seen in mouse ovaries (Jones and Krohn, 1961). The continuous initial recruitment and the loss of primordial follicles through apoptosis leads to depletion of the original follicle pool. In this way, the diminishing follicle reserve leads to the onset of menopause (Richardson et al., 1987; Faddy et al., 1992; Faddy and Gosden, 1996). In the present study, apoptosis was not detected in primary and secondary follicles. This is in discordance with many reports in mammals, which show that atresia affects the early stages of follicular development (Gougeon, 1996; Yang and Rajamahendran, 2000; Depalo et al., 2003). At the beginning of reproductive life, the number of primordial follicles in a normal cycling cow is ca 100 000 (Erickson, 1966). Only some 1000 of these follicles will develop to the preovulatory stage when considering 180 cycles in a reproductive lifespan of 10 years in cattle with in each cycle an average of 6 developing secondary follicles (Richards, 1994). Earlier studies showed that only 2500 follicles were found in the ovaries of 10-year old cattle (Käppeli, 1908). This implicates that more than 96 % of the early follicular stages, i.e. primordial, primary and secondary follicles, become atretic. Despite our results, it is very unlikely that atresia of primordial, primary and secondary follicles does not occur in cattle and that depletion of the pool of primordial follicles is only caused by the continuous initial recruitment during lifetime. A plausible explanation might be that the number of atretic primordial, primary and secondary follicles is underestimated in our study by the fact that only a single snapshot evaluation of ovarian structures was made for each cow. As follicular atresia starts already before birth and continues for many years, the chance to detect these atretic follicles is clearly reduced when only three histological sections are examined. Elaborate studies in which the complete ovaries are analysed by means of stereology might help to answer this question.

Increasing scores of Ki-67 were seen in primary and secondary follicles. In addition, the follicle/granulosa cells of primary and secondary follicles showed moderate scores of ER β mRNA and PR whereas ER α was completely absent. The increasing scores for ER β mRNA and PR in the follicle/granulosa cells in the early follicle stages suggest a pivotal role of ER β and PR in follicular growth and development.

Although little is known about the estrogen action in the ovary, the constant presence of ER β mRNA in the primordial, primary, secondary and tertiary follicles suggests that estrogens may modulate the proliferation and differentiation of the follicular cells. In vital tertiary follicles, ER β mRNA levels were higher than in primordial, primary and secondary follicles. These findings may reflect the role of estrogens in the recruitment and dominance of tertiary follicles. Scores for ER β mRNA were different in the various cell layers of the tertiary follicles. ER β mRNA levels were significantly higher in the theca externa layer than in the theca interna layer and significantly higher in the theca interna layer than in the granulosa layer. This observation is consistent with reports in other species and indicates that ER β mRNA has cell-specific functions (Pelletier et al., 1999; Pelletier et al., 2000; Slomczynska et al. 2001). Furthermore, it has been suggested that the locally produced estrogens by the granulosa cells itself (Rosenfeld et al., 1999). Additionally, in these tertiary follicles low levels of ER α were found. According to Couse et al. (1997) the presence and interaction of both receptors within vital tertiary follicles are required for the late stage of follicular growth.

In vital tertiary follicles the PR levels increased towards oestrus, while the ER β mRNA levels decreased. In this way it is suggestive that the PR expression is induced by the LH surge (Hild-Petito et al., 1988; Jo et al., 2002) and that PR rather than ER, is involved in the ovulatory process. The crucial role of PR in the ovulatory process has been demonstrated in PR-deficient mice, since such mice develop large follicles but fail to ovulate (Lydon et al., 1995). All these findings emphasize the pivotal role of progesterone and its receptor in the ovulation process.

In the present study, high numbers of attretic tertiary follicles were observed in all bovine ovaries during the different cycle stages. This observation supports the theory that follicle recruitment is mainly situated at the level of the tertiary follicles and that a large number of non-selected tertiary follicles regress (McGee and Hsueh, 2000). The number of apoptotic

cells in the different layers of atretic tertiary follicles showed significant differences. Cystic atretic follicles showed higher apoptotic levels compared to obliterative follicles. In cystic atretic follicles disruption of the intercellular contacts may stimulate apoptosis. In contrast, in obliterative follicles the granulosa and theca layers both collapse, while the cell-to-cell contacts remain intact and the intercellular integrity inhibits the induction of apoptosis. Ki-67 was detected at low levels in the obliterative atretic follicles, while in cystic atretic follicles very high levels, similar to these of vital tertiary follicles were observed. This indicates that cell proliferation continues even when the follicle becomes cystic atretic. Furthermore, it has been shown that atretic follicles may continue to secrete considerable quantities of steroid hormones (Jolly et al., 1994). Whether the cystic atretic follicles in the bovine ovary continue to produce steroids and consequently play a functional role remains to be elucidated.

Estrogens and progesterone are reported to be protective against apoptosis in the bovine tertiary follicles by binding to the specific receptors (Quirk et al., 2004; Emmen et al., 2005). However, in atretic tertiary follicles low to intermediate levels of PR and high levels of ER β mRNA were observed. This implicates that despite the presence of ER and PR and their putative protective effect, apoptosis is induced in these follicles. Therefore, it is suggestive that the effect of pro-apoptotic factors involved in the process of follicular atresia (e.g., tumor necrosis factor α , caspase-10 and β glycan) surpasses the protective effect of progesterone and estrogens in atretic tertiary follicles (Evans et al., 2004).

In the corpora hemorrhagica high levels of PR and intermediate to high levels of ER β mRNA were found. According to earlier studies of Quirk (2004) this observation may indicate that the protective activity of both hormones further continued in corpora hemorrhagica maintaining the cells vital.

In the present study PR immunostaining in the corpora lutea was lower than in most other ovarian structures. This can be due to the fact that the locally produced high levels of progesterone may downregulate the PR expression in corpora lutea cells (Revelli et al., 1996; Rueda et al., 2000). In the lutein cells and internal stroma cells a similar expression pattern for PR was observed. Furthermore, the PR expression in these cell types was higher than in the capsular cells. These observations suggest the influence of progesterone in the development of luteinized granulosa and theca cells in an autocrine and paracrine way (Sasano and Suzuki, 1997). In the corpus luteum ER α was exclusively observed during late

dioestrus and in this cycle stage the ERβ mRNA scores increased. These findings are suggestive for an up-regulation of both types of ERs by progesterone. It is of interest that the enhanced secretion of progesterone in the corpus luteum only leads to up-regulation of ER, but not of PR. In the corpora lutea high numbers of apoptotic lutein cells were found. Furthermore, the number of caspase-3 positive lutein cells was much higher than the number of TUNEL positive cells. This finding may indicate that in corpora lutea many caspase-3 positive cells disappear, probably by phagocytosis, before DNA fragmentation occurs. Additionally, the amount of caspase-3 positive cells increased from low numbers at oestrus to maximal levels at dioestrus/prooestrus. This is in accordance with the positive correlations observed between the plasma progesterone on apoptosis in the corpus luteum remains to be elucidated.

In corpora albicantia intermediate to high levels of ER β mRNA as well as low levels of ER α were observed. Therefore, it is possible that the simultaneous presence of ER α and ER β might lead to heterodimerization and possible involvement of ER α/β in the formation of the corpus albicans.

In both deep and superficial stroma the expression pattern of PR is similar to that in the follicles, namely intermediate to high levels during oestrus, metoestrus and late dioestrus. Furthermore, high scores for ER β mRNA and ER α were found in cells of the deep and superficial stroma. All these observations may account for possible stromal-epithelial interactions and suggest that both ER and PR may mediate the regulation of follicular development. The possibility that the paracrine action of ovarian steroids on follicles is mediated through stromal steroid receptors has also been suggested by Revelli and co-workers (1996). Both types of stroma cells frequently expressed a higher hormonal sensitivity compared to the tunica albuginea cells. Furthermore, the relatively low numbers of apoptosis and the low proliferation rate in cells of the stroma and the tunica albuginea suggest a low turnover rate of the cells throughout the estrous cycle.

There seemed to be a temporal difference in receptor expression in the tunica albuginea compared to cells of the surface epithelium. In the latter, the expression of PR, ER α and ER β mRNA was obvious during oestrus, which may be suggestive for the involvement of these receptors and their hormones in the ovulatory process.

When verifying putative differences between the active and inactive ovaries, an ovary was considered to be active when a Graafian follicle or a mature corpus luteum was present. In 14 out of the 23 cows analyzed in this study, the left ovary was the most active (Chapter 1.1.2). This is in contrast to findings of Nation (1999) who found that in cows ovulation occurs mostly in the right ovary. Nevertheless, in bovine the follicle populations in both ovaries are the same (Erickson, 1966). Conclusions concerning the verification of the most active ovary should be considered with caution, as the number of analysed cows in the present study is very low.

Finally, correlations between the plasma progesterone levels on the one hand and the immunostaining scores of their receptors, apoptosis and proliferation on the other hand, were determined. Significant correlations between the plasma progesterone concentrations and PR or ER were always negative. These findings are in accordance with a study in bovine endometrium tissues (Boos et al., 1996) and indicate that progesterone may downregulate PR and ER expression in the bovine ovarian cell types. Generally, the plasma progesterone concentration was negatively correlated with the score for apoptosis and mitosis in the bovine ovarian tissues. However, in the lutein cells the plasma progesterone concentration was negatively correlated with the apoptotic score detected by the TUNEL technique, while positive using the caspase-3 assay.

The abundant expression of ER and PR in the different bovine ovarian cell types and the cyclic variations of ER and PR expression during the bovine estrous cycle stages indicate the importance of estrogens and progesterone in the regulation of the ovarian activity.

The topographic and numerical data of the present study may be useful for further studies regarding the role of estrogen and progesterone both in normal ovaries and in pathological conditions such as cystic ovarian disease.

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SUMMARY

During the bovine estrous cycle the different ovarian structures are subject to extensive tissue remodelling that consists of sequential phases of cell proliferation and cell death. Although the cyclic ovarian processes have been studied intensively, the regulatory mechanisms that are responsible for the cyclic changes remain unclear. A major role is attributed to the sex steroid hormones including estrogens and progesterone. Both hormones exert their functions by binding to specific receptors, i.e., the estrogen receptors (ER α and ER β) and the progesterone receptors (PRA and PRB).

The aim of the present study was to localize the estrogen receptors and progesterone receptors in the different ovarian cell types of normally cycling cows and to examine the variation of expression of ER and PR throughout the estrous cycle. Furthermore, the relation between the presence of the steroid receptors and the cellular dynamics was examined by determining the correlation between the plasma progesterone levels and the immunostaining scores of their receptors, apoptosis and proliferation.

The study was performed in 23 cows. According to their ovarian morphological characteristics and the corresponding plasma progesterone levels the animals were classified in 5 different stages, viz. oestrus, metoestrus, early dioestrus, late dioestrus and procestrus.

The first chapter of this thesis describes the localization of ER (ER α and ER β mRNA) and PR (PRA and PRB) in the bovine ovary at the different stages of the estrous cycle. In a preliminary study ER β mRNA was examined using *in situ* hybridization in the ovaries of randomly selected cows, irrespective of the cycle stage of the animals. A 602-bp fragment of ER β mRNA was cloned, sequenced and digoxigenin-labelled. Semi-quantitative evaluation in the different ovarian cell types demonstrated a cell-specific localization of ER β mRNA in several cell types.

The second section of the first chapter describes the localization of ER α and ER β mRNA in the various ovarian cell types during the 5 stages of the estrous cycle. ER α was detected by immunohistochemistry and the localization of ER β mRNA was examined using *in situ* hybridization. Cytoplasmic staining for ER β mRNA and nuclear staining for ER α were observed in different cell types. The immunostaining of ER α was low in the ovarian follicles, tunica albuginea and surface epithelium, but high in cells of the deep stroma and superficial stroma. These findings indicate a functional role of ER α in the stroma cells surrounding the follicles and suggest stromal-epithelial actions.

The predominant role of ER β mRNA over ER α was manifestly present in the bovine ovarian structures. The moderate scores of ER β mRNA in primordial and primary follicles and the high scores in tertiary follicles may indicate that estrogens are involved in follicular growth and act via ER β . In corpora lutea cyclic variations of ER β mRNA and low levels of ER α were observed. The simultaneous presence of both receptors in several ovarian cell types suggests colocalization of both ER α and ER β mRNA and may lead to ER α /ER β heterodimerisation. In general, the correlation between the plasma progesterone levels and the scores for ER α and ER β mRNA was low in all ovarian cell types.

The next section of the first chapter describes the immunolocalization of the progesterone receptor in the various bovine ovarian tissues during the 5 cycle stages. The PR immunostaining was observed in the nuclei of all ovarian cell types. Many differences were observed in the scores of PR between the different cell types and high variations were seen in the PR expression, concomitant with the bovine cycle stages. In primordial, primary and secondary follicles the score for PR in the follicle cells increased along with the development of the follicle. In both vital tertiary follicles and cystic atretic follicles a similar expression pattern and moderate scores of PR were seen, whereas in obliterative atretic follicles the expression pattern differed and the score was much lower. These observations might suggest that the physiologic processes continue in cystic atretic follicles as in vital tertiary follicles. In contrast, obliterative atretic follicles appear as a later stage or as a different type of follicular atresia. Generally, the scores for PR in all follicular structures were high at oestrus and decreased during the following stages. This finding is suggestive for a cell-specific and cycledependent influence of progesterone. In corpora lutea cells the staining pattern of PR was lower than in the other ovarian cell types, which can be due to a negative effect of the locally produced high levels of

progesterone. Furthermore, a negative correlation was found between the PR scores and the plasma progesterone concentration.

To examine the relation between the distribution of the estrogen and progesterone receptors and the cell dynamics, the cellular proliferative activity and cell death have been assessed in the ovaries of the cows.

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Chapter 2 describes the cell-specific localization of apoptosis in the bovine ovary at different stages of the estrous cycle. Apoptosis was localized in the different cell types during the various stages of the estrous cycle, using the detection of active caspase-3, in situ end labelling (TUNEL) and DNA fluorescent staining (DAPI). Each of the three techniques demonstrates a different aspect of the apoptotic process. The caspase-3 assay detects activated caspase-3 which is present at the onset of the apoptotic process, whereas the TUNEL assay and the DAPI staining demonstrate DNA fragmentation and chromatin condensation, respectively, which are only visible at later stages of the apoptotic process. In primordial, primary and secondary follicles no apoptosis was detected. In contrast, high numbers of attretic tertiary follicles were observed in all bovine ovaries during the different cycle stages. This observation supports the fact that follicle recruitment is mainly situated at the level of tertiary follicles and that a large number of non-selected tertiary follicles regress. Cystic atretic follicles showed higher apoptotic levels compared to obliterative atretic follicles. In cystic atretic follicles disruption of the intercellular contacts may stimulate apoptosis, whereas in obliterative attrict follicles the granulosa and theca layers both collapse, while the cell-tocell contacts remain intact and the intercellular integrity inhibits the induction of apoptosis. In all ovarian structures examined, variations in apoptotic scores were seen in the different cycle stages, suggesting a cycle-dependent influence on apoptosis, although correlation with plasma progesterone concentrations was low.

Chapter 3 describes the proliferation patterns in the bovine ovary at different stages of the estrous cycle. Proliferation was examined using immunohistochemical detection of the Ki-67 proliferation associated antigen. Our results are in accordance with studies in humans and mice showing that the recruited primordial follicles develop into primary follicles without numerical proliferation of the follicle cells. Follicle cell proliferation is initiated in primary follicles and increases progressively with the development of the follicle. A remarkable finding of the present study is that the scores for Ki-67 in vital tertiary and cystic atretic follicles were similar and high, whereas in obliterative atretic follicles the score was much lower. This finding implicates that the proliferative activity of vital tertiary follicles further continues in cystic atretic follicles but not in obliterative atretic follicles. In corpora lutea a different cyclic pattern was observed with high levels during metoestrus and early dioestrus, which might suggest that progesterone is involved in the proliferative activity of the lutein cells. However, correlation with the plasma progesterone concentrations was low.

In the final chapter the conclusions of the different studies are summarized and discussed. The abundance of estrogen receptors and progesterone receptors in the various bovine ovarian cell types indicates the importance of estrogens and progesterone in the regulation of the ovary activities. The topographic and numerical data of the present study may be useful for further studies on the role of estrogen and progesterone in the regulation of ovarian activity.

SAMENVATTING

Tijdens de oestrische cyclus van het rund zijn de verschillende ovariële structuren onderworpen aan uitgebreide weefselveranderingen. Deze veranderingen bestaan uit opeenvolgende fasen van celproliferatie (mitose) en celdood (apoptose). Alhoewel de ovariële processen reeds uitvoerig bestudeerd zijn, blijven de regulatiemechanismen die verantwoordelijk zijn voor de cyclische veranderingen onduidelijk. De steroïdale geslachtshormonen oestrogeen en progesteron spelen hierbij een grote rol. Beide hormonen oefenen hun functie uit door de binding aan specifieke receptoren, de oestrogeenreceptoren (ER α en ER β) en de progesteronreceptoren (PRA en PRB).

De eerste doelstellingen van deze studie bestonden uit het lokaliseren van de oestrogeenreceptoren en de progesteronreceptoren in de verschillende ovariële celtypes van normaal cyclerende runderen en het onderzoeken van mogelijke veranderingen in de receptordistributie tijdens de oestrische cyclus. Een volgende doelstelling was het nagaan van een mogelijke relatie tussen de aanwezigheid van de steroïdreceptoren en de cellulaire activiteit, met name proliferatie en apoptose. Een laatste objectief was het onderzoeken van de mogelijke correlatie tussen deze parameters en de progesteronconcentraties in het bloedplasma.

Dit doctoraatsonderzoek is uitgevoerd op 23 koeien. De runderen zijn over vijf verschillende groepen verdeeld naargelang hun progesteronconcentratie in het bloedplasma en de morfologische ovariële kenmerken. De vijf stadia waarin de oestrische cyclus van het rund werd opgesplitst zijn: oestrus, metoestrus, vroege dioestrus, late dioestrus en prooestrus.

Het eerste hoofdstuk van deze thesis beschrijft de lokalisatie van de oestrogeenreceptoren (ER α en ER β) en de progesteronreceptoren (PRA en PRB). In een voorafgaande studie is ER β mRNA onderzocht in de ovaria van een aantal koeien, ongeacht hun cyclusstadium. Ter bepaling van ER β mRNA werd een probe aangemaakt bestaande uit een ER β mRNA-fragment van 602 baseparen. Dit fragment werd gekloneerd, gesequeneerd en gemerkt met digoxigenine. De *in situ* hybridisatietechniek is op punt gesteld en toegepast op een beperkt aantal runderen, onafgezien hun cyclusstadium. Semi-kwantitatieve evaluatie van de verschillende ovariële celtypes toonde aan dat de lokalisatie van ER β mRNA celspecifiek was.

In het tweede deel van dit hoofdstuk is de lokalisatie van ER α en ER β mRNA beschreven in de verschillende ovariële celtypes gedurende de vijf stadia van de oestrische cyclus. ERa werd bepaald door middel van immunohistochemie en de lokalisatie van ERB mRNA is onderzocht gebruik makende van de in situ hybridisatietechniek. ERa is in de celkernen aangetroffen terwijl ERB mRNA gelokaliseerd werd in het cytoplasma. De score voor ERa was laag in de ovariële follikels, de tunica albuginea en het oppervlakte-epitheel, maar was hoger in de cellen van het omringende diepe en oppervlakkige stroma. Deze bevindingen duiden op een functionele rol van ERa in de stromacellen die de follikels omringen en kan wijzen op stroma-epitheliale interacties. De middelmatige scores van ERB mRNA in primordiale en primaire follikels en de hoge scores in tertiaire follikels kunnen erop wijzen dat oestrogenen en ERß betrokken zijn in de follikulaire groei. In cellen van het corpus luteum werden cyclische schommelingen voor ERB mRNA gezien en is tevens de aanwezigheid van ERα aangetoond. De aanwezigheid van ERβ mRNA was duidelijk groter dan die van ERα in alle follikulaire structuren van het rund. Daarnaast zou de gelijktijdige aanwezigheid van beide receptoren kunnen wijzen op co-lokalisatie van ERα en ERβ mRNA wat kan leiden tot ERα/ERβ heterodimerisatie. In het algemeen was de correlatie tussen de progesteronconcentraties in het bloedplasma en de scores voor ERa en ERß mRNA zwak en negatief in alle ovariële celtypes.

Het laatste deel van het eerste hoofdstuk beschrijft de immunolokalisatie van de progesteronreceptor in de verschillende ovariële celtypes gedurende de vijf stadia van de oestrische cyclus. De immunokleuring voor PR was zichtbaar in de celkernen van alle ovariële structuren. De PR-scores varieerden sterk tussen de verschillende celtypes, doch opmerkelijk was het steeds voorkomen van hetzelfde cycluspatroon. De score voor PR in primordiale, primaire en secundaire follikels nam toe naargelang de ontwikkeling van de follikel. Vitale en cysteuze tertiaire follikels vertoonden hetzelfde expressiepatroon voor PR met middelmatige scores. Daarnaast was het expressiepatroon van PR in oblitererende follikels verschillend van dat van de vitale en cysteuze follikels en was de score veel lager. Deze bevindingen kunnen erop wijzen dat cysteuze follikels ontstaan na atresie van vitale tertiaire follikels. Dit impliceert de continuïteit van de fysiologische processen zelfs wanneer de follikel cysteus wordt. Anderzijds zou een oblitererende follikel kunnen beschouwd worden als een laat stadium van een cysteuze follikel, of zoals beschreven in de literatuur, als een ander type van atresie. Over het algemeen was de score voor PR hoog in alle follikulaire structuren tijdens de oestrus en daalde deze score gedurende de volgende stadia. Deze

bevinding zou kunnen wijzen op een celspecifieke en cyclusafhankelijke invloed van progesteron. In de cellen van de corpora lutea was het expressiepatroon lager dan in de andere ovariële celtypes. Dit kan te wijten zijn aan een negatief effect van de lokaal geproduceerde hoge progesteronconcentraties in het corpus luteum. Daarenboven was de correlatie tussen de PR-scores en de progesteronconcentraties in het bloedplasma negatief.

Om de relatie te bestuderen tussen de distributie van de oestrogeenreceptor en de progesteronreceptor enerzijds en de cellulaire dynamiek anderzijds werden proliferatie en apoptose nagegaan in de verschillende celtypes van de rundovaria.

Hoofdstuk 2 beschrijft de celspecifieke lokalisatie van apoptose in het rundovarium gedurende de vijf verschillende stadia van de oestrische cyclus. Hiervoor werd gebruik gemaakt van de detectie van actieve caspase-3, *in situ* end labelling (TUNEL) en de DNA fluorescent staining techniek (DAPI). Elk van deze technieken toont een verschillend aspect aan van het apoptotisch proces. De caspase-3 techniek detecteert actief caspase-3, dat aanwezig is tijdens het apoptotisch proces. De TUNEL techniek en de DAPI-kleuring tonen DNA-fragmentatie en chromatinecondensatie aan, die alleen waarneembaar zijn tijdens de latere stadia van het apoptotisch proces.

Een belangrijke bevinding in deze thesis was de afwezigheid van apoptose in primordiale, primaire en secundaire follikels. Daarentegen werden vele atretische tertiaire follikels waargenomen in alle runderovaria tijdens de verschillende cyclusstadia. Deze observatie bevestigt het feit dat follikelrekrutering voornamelijk plaatsgrijpt op het niveau van de tertiaire follikels en dat een groot aantal niet-geselecteerde tertiaire follikels regresseren. Cysteuze follikels vertonen hogere apoptotische scores in vergelijking met oblitererende follikels. In cysteuze follikels zou het teloorgaan van intercellulaire contacten apoptose stimuleren. In oblitererende follikels daarentegen blijven de celcontacten behouden en dit zou de inductie van apoptose afremmen. In alle onderzochte ovariële structuren werden variaties in de apoptotische scores gezien tijdens de verschillende cyclusstadia. Dit suggereert dat apoptose een cyclusafhankelijk proces is, ook al was de correlatie tussen apoptose en de progesteronconcentraties in het bloedplasma laag.

In hoofdstuk 3 werd de proliferatieve activiteit in het rundovarium tijdens de verschillende ovariële stadia beschreven. Proliferatie is onderzocht door middel van immunohistochemische

detectie van het Ki-67 nucleaire antigen. De resultaten bekomen uit dit onderzoek zijn in overeenstemming met studies bij de mens en muis en impliceren dat de gerekruteerde primordiale follikels zich ontwikkelen tot primaire follikels zonder toename van het aantal follikelcellen. Proliferatie wordt geïnitieerd in de primaire follikels en neemt toe naarmate de follikel zich verder ontwikkelt. Een opmerkelijke bevinding van de huidige studie is dat in vitale en cysteuze tertiaire follikels even hoge scores voor Ki-67 werden gevonden, terwijl de score in oblitererende follikels duidelijk lager was. Deze bevinding impliceert dat de proliferatieve activiteit van de vitale tertiaire follikels aanhoudt in cysteuze follikels maar niet in oblitererende follikels. In corpora lutea werd een verschillend cyclisch patroon teruggevonden met hoge scores voor Ki-67 tijdens metoestrus en vroege dioestrus. Dit zou kunnen betekenen dat progesteron de proliferatieve activiteit in de luteïnecellen beïnvloedt. Nochtans is de correlatie tussen de score van Ki-67 en de progesteronconcentraties in het bloedplasma laag.

De uitgebreide distributie van oestrogeen- en progesteronreceptoren in de verschillende ovariële celtypes van het rund illustreert het belang van oestrogenen en progesteron in de fysiologie van het rundovarium. De topografische en numerieke gegevens bekomen uit deze studie kunnen van nut zijn voor verder onderzoek naar de specifieke rol van oestrogenen en progesteron in de regulatie van de ovariële activiteit. Mylène D'Haeseleer werd geboren op 5 december 1963 te Aalst.

Zij behaalde het diploma van Gegradueerde in de Farmaceutische en Biologische Technieken aan het Paramedisch Hoger Instituut te Anderlecht in 1986 met onderscheiding. Daarna trad zij achtereenvolgens in dienst bij de vakgroep Plantengenetica, de vakgroep Algemene Plantkunde en Natuurbeheer en de vakgroep Ecologie aan de Vrije Universiteit Brussel. Zij behaalde het diploma van Licentiaat in de Moleculaire Biologie aan de Vrije Universiteit Brussel in 2000 met onderscheiding.

Geboeid door het wetenschappelijk onderzoek, startte zij in 2001 een doctoraatsstudie bij de vakgroep Morfologie aan de Faculteit Diergeneeskunde. Deze studie werd gefinancierd door het Bijzonder Onderzoeksfonds van de Universiteit Gent. Tevens behaalde zij het diploma van de Doctoraatsopleiding in de Diergeneeskundige Wetenschappen aan de Universiteit Gent in 2006.

Mylène D'Haeseleer is auteur of mede-auteur van meerdere wetenschappelijke publicaties en rapporten en nam actief deel aan verschillende nationale en internationale congressen.

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