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Endometriosis and Angiogenic Factors

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1. Introduction

Little is known about the pathogenesis of endometriosis. The prevailing hypothesis is that following retrograde menstruation, uterine endometrial tissue attaches, invades the peritoneal surface, and becomes vascularized.

The development of new blood vessels represents a crucial step during the establishment of endometriosis because endometriotic implants require neovascularization to guarantee oxygen and essential nutrient supply (Groothuis et al., 2005; McLaren, 2000). The interaction between the ectopic endometrium and the peritoneal tissue is a prerequisite for the induction of angiogenesis and the maintenance of endometriosis

At least, three processes appear to be critical to the establishment of endometriosis, according to the implantation theory: invasiveness, tissue remodeling and interactions between the ectopic endometrium and the surrounding peritoneal tissues (Giudice et al., 2008).

The establishment of endometriotic lesions needs a cascade of neoangiogenic factor, like the vasculatr endothelial factor, cytokines and metalloproteinases: this complex interrelation between factors permit sprouting of capillaries from pre-existing vessels and the subsequent supply for the development of ectopic implants (Hyder and Stancel, 1999).

2. Vascular endothelial growth factors family

2.1 Vascular Endothelial Growth Factor(VEGF)

2.1.1 VEGF family and its receptors

Vascular endothelial growth factors are important signaling proteins involved in both vasculogenesis (the *de novo* formation of the embryonic circulatory system) and angiogenesis (the growth of blood vessels from pre-existing vasculature). VEGF family comprises seven members: VEGF-A, VEGF-B, VEGF-C, VEGF-D, VEGF-E, VEGF-F, and PIGF. All members have a common VEGF homology domain. This core region is composed of a cystine knot motif, with eight invariant cysteine residues involved in inter- and intramolecular disulfide bonds at one end of a conserved central four-stranded-sheet within each monomer, which dimerize in an antiparallel, side-by-side orientation (Neufeld G, wt al 1999).

The human *VEGF* gene (*VEGFA*, OMIM 192240) is located on chromosome 6p12 (Zhao Z, et al 2008).

Vascular endothelial growth factor (VEGF) is one of the most potent and specific angiogenic factors. When VEGF binds to its targeted receptor, the VEGF receptor activation leads to a rapid increase in intracellular Ca2+ and inositol triphosphate concentrations in endothelial cells. The basic physiological function of VEGF is to induce angiogenesis, which allows the endometrium to repair itself following menstruation. It also modulates the characteristics of the newly formed vessels by controlling the microvascular permeability and permitting the formation of a fibrin matrix for endothelial cell migration and proliferation. This modulation may be responsible for local endometrical edema, which helps prepare the endometrium for embryo implantation. In endometriosis patients, VEGF is localized in the epithelium of endometriotic implants, particularly in hemorrhagic red implants. Moreover, the concentration of VEGF is increased in the PF of endometriosis patients. The exact cellular sources of VEGF in PF have not yet been precisely defined. Although evidence suggests that endometriotic lesions themselves produce this factor, activated peritoneal macrophages also can synthesize and secrete VEGF.

In women, physiological neoangiogensis is presented during the female reproductive cycle. VEGF-A is important in luteal angiogenesis. VEGF-A mRNA or protein is detectable in the granulosa cells of primordial and primary follicles, as they progressively become localized to the granulosa surrounding the oocyte and theca cells of the preovulatory follicle. After ovulation, VEGF-A mRNA and protein expression are observed in granulosa-derived luteal cells. VEGF-A expression in the corpus luteum appears highest early in the luteal phase and declines after the mid-luteal phase, with little or no expression in the late corpus luteum (Fraser HM, et al 2000). Gonadotrophic hormones, particularly luteinizing hormone (LH), appear to be major regulators of angiogenesis in the ovary (Hyder SM. Et al, 1999). The LH-stimulated luteinization of granulosa cells at the time of ovulation is associated with enhanced VEGF-A expression.

When VEGF is overexpressed, it can contribute to disease. Three VEGF tyrosine kinase receptors have been identified: The fms-like tyrosine kinase Flt-1 (VEGFR-1/Flt-1), the kinase domain region, also referred to as fetal liver kinase (VEGFR-2/KDR/Flk-1), and Flt-4 (VEGFR-3). Each receptor has seven immunoglobulinlike domains in the extracellular domain, a single transmembrane region, and a consensus tyrosine kinase sequence interrupted by a kinase insert domain (Ortega N et al, 1999). Binding its receptors R1, VEGF regulates development of tubular capillaries; binding R2 receptor, it promotes mesodermic cells differentiation into endothelial cells. In vivo and in vitro experiments indicate that steroid hormones, hypoxia and nitric oxide are potent inducers of vascular endothelial growth factor gene expression by enhancing hypoxia inducible factor 1-a activity and by the activation of the AKT/protein kinase B pathway (Kimura H, et al 2006).

2.1.2 VEGF role in endometriosis

Although the aetiology of endometriosis is unknown, it is generally accepted that the condition is a result of the implantation of exfoliated endometrium, deposited in the peritoneal cavity following retrograde menstruation (Sampson J 1927). When the exfoliated endometrium enters the peritoneal cavity and becomes attached to the mesothelial layer through attachment proteins like the cadherins, a process of angiogenesis is essential for further implantation and the development of peritoneal endometriosis (Nisolle M et al 1993). Angiogenesis is a fundamental process by which new blood vessels are formed and is considered as a major process in the pathogenesis of endometriosis. It is dependent on

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soluble factors released from cells. Many factors are involved in this complex mechanism, including FGF-a, FGF- b, PD-ECGF and VEGF, stimulate vascular endothelial cell growth *in vitro* and angiogenesis *in vivo* (Gordon J. Et al, 1995).

The development and maintenance of endometriosis is dependent on the recruitment of blood vessels to the endometriotic lesions from pre-existing ones to guarantee oxygen and essential nutrient supply. It has been shown that those endometriotic lesions recruit blood vessels by inducing angiogenesis. In a rat model, analyses of the assessed microvessel density demonstrated that angiogenesis is higher in ectopic endometriotic lesions compared with the eutopic endometrium (Machado DE et al, 2010).

Promoting angiogenesis, VEGF is involved in the etiology but also in the maintenance of peritoneal endometriosis. The endometrial tissue that has been migrated into the peritoneal space develops different angiogenic properties and invasive patterns, which could have a role in the implantation in the peritoneum of the pelvic cavity. It is likely that the action of VEGF in the implant of endometrial ectopic cells is to promote the differentiation of mesodermic cells into endothelial cells and to regulate the tubular capillary formation (Di Carlo C. Et al 2009).

VEGF-A is localized predominantly in the glandular epithelium of endometriosis lesions. Peritoneal fluid concentrations of VEGF have been demonstrated to be significantly higher in women with endometriosis than in the control patient and a positive correlation between the severity of endometriosis and the concentrations of VEGF in peritoneal fluid has been observed. Peritoneal fluid itself induced higher expression of the protein participating in the establishment and persistence of peritoneal endometriosis (Fasciani G. Et al, 2000).

In patients with endometriosis, high concentrations of VEGF in cystic and peritoneal fluids may be ascribed to a state of inflammation, where macrophages, which are the main source of this growth factor, play a central role.

It has been shown that cytokines released from immune cells play an important role in the pathogenesis of endometriosis, and many of these cytokines possess angiogenic activity. Pelvic implants and peritoneal fluid macrophages are the most likely source of VEGF-A in peritoneal fluid. McLaren (McLaren J, et al 2006) showed that peritoneal fluid macrophages express receptors for steroid hormones and secrete VEGF-A in response to ovarian steroids. The VEGF receptors VEGFR-1 and VEGFR-2 were also detected, suggesting an autocrine regulation. Peritoneal macrophages and activated lymphocytes seem to play an integral role in the secretion of proinflammatory/proangiogenic cytokines resulting in upregulation of VEGF from infiltrating neutrophils and macrophages (Machado DE et al, 2010).

Ovarian endometriotic cysts also over express VEGF in their cystic fluid with respect to follicular and serous cysts and to a similar degree as ovarian cystadenocarcinoma and a negative or positive correlation of VEGF expression has been reported with cyst diameter.

In ovary the process of angiogenesis is characterized by the existence of complex interrelations between the cell components of the ovarian cyst: diffuse VEGF expression in epithelial cells was associated with larger cysts; high VEGF expression in capsular fibroblast was associated with bilateral cysts; and expression of VEGF was found to be related in epithelial cells, capsular fibroblast and vessels, suggesting that neoangiogenesis might especially affect the outer cell cyst wall, thus contributing to cyst growth.(Goteri G. et al,

2004). The inner layers of cysts are characterized by high microvessel density but low expression of VEGF, whereas in the outer fibrosclerotic capsule, the vessels were less abundant, but Had a higher expression of VEGF and survive, thus activated to proliferate and protected from programmed cells death. Angiogenesis mediated by VEGF in the outer capsule contributed to the cyst growth and to the fibrosing process of adhesion (Goteri G. et al 2010). Antiangiogenetic drugs could act on the capsular vasculature and block the growth of ovarian cyst. The high VEGF levels could provoke an increase in the subperitoneal vascular network and facilitate implantation and viability of endometrial cells in the retroperitoneal space. Concerning sVEGFR-1, the highest levels of this protein were found in peritoneal fluids and cystic fluids of endometriosis patients with respect to both benign and malignant serous cysts. The soluble form of VEGFR-1, as already stated, should function as a modulator of VEGF's angiogenic activity. For this reason, it seems that sVEGFR-1 is secreted in proportionate amounts as VEGF itself. In benign cyst and peritoneal fluids, VEGF and sVEGFR-1 concentrations are proportionately low, while in endometriosis cyst and peritoneal fluids, VEGF and its soluble receptor are both expressed in much higher concentrations. Endothelial sVEGFR-1 is also known to be up-regulated by its ligand, VEGF-A, and the high levels of VEGF-A found in endometriomata compared with cystadenomas are likely to further contribute to up-regulation of sVEGFR-1. On the other hand, in patients affected by cystadenocarcinomas, there was discordance between the levels of VEGF and the levels of sVEGFR-1 in both cyst and peritoneal fluids. In fact, in malignant processes there seems to be an imbalance between pro-angiogenetic factors, represented by VEGF, and antiangiogenetic factors, represented by sVEGFR-1, leading to a disordered and exaggerated formation of blood vessels (Artini PG et al, 2008).

The microenvironment of endometriosis is a locale of important secretion of angiogenic factors that play a key role in the establishment and maintenance of endometriotic lesions, and suggest that the balance of these local pro-antiangiogenic factors and cytokines may determine whether endometriotic lesions develop and grow.

2.1.3 Genetic polymorphisms of VEGF genes

An important aspect of the correlation between endometriosis and VEGF, also for possible future therapeutic application is that the polymorphisms in vascular endothelial growth factor gene are associated with the risk of familial endometriosis. The human VEGF gene (VEGFA, OMIM 192240) is located on chromosome 6p12 (Zhao Z, et al 2008). VEGF messenger ribonucleic acid and protein were significantly higher in women with endometriosis, which supported a key role for VEGF in the pathological angiogenesis in endometriosis (Gilabert-Estelles J, 2007). In particular several transcription factor-binding sites are found in the VEGF 5' -untranslated region and variation within the region increases the transcriptional activity. A single family in two generations with four members who have histologically proven endometriosis showed that the circulating levels of VEGF were higher than the healthy control group, indicating a role for VEGF in disease susceptibility (Simpson JL et al 2003). In Chinese patients, the T allele of the VEGF gene -60 T/C (rs833061) polymorphism was associated with a higher risk of endometriosis. Study of the VEGF +405 G/C (rs2010963) polymorphism in a Korean population showed that the SNP was associated with the risk of advanced stage endometriosis. The analysis of both SNPs in an Indian population identified a haplotype associated with endometriosis In addition, the analysis of VEGF -460 T/C (rs833061), +405 G/C (rs2010963) and +936 C/T

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(rs3025039) polymorphisms in 147 endometriosis cases and 181 controls found a positive association between stages III–IV disease and the *VEGF* +936 T allele in a Japanese population. (Zhao Z, et al 2008). The first reported study in a Caucasian population of +405 G/C (rs2010963) in 203 Italian women affected with endometriosis and 140 controls reported a weak association of the C allele with endometriosis (Gentilini D et al, 2008).

2.1.4 Therapeutical approach

Therapy of endometriosis consists of surgical removal of implants or medical treatment such as analogues of Gonadotrophin releasing hormone or oral contraceptive or progestins. This therapeutic approach has been shown to be of limited benefit so new approaches need to be developed. Considering the importance of angiogenesis in developing and maintaining disease, and the role of vascular endothelial growth factor, anti-angiogenetic drugs could be very important.

Romidepsin, the Histone deacetylase (HDC) inhibitor, modulates the expression of a variety of genes by alteriting gormatin structure. It has been recently shown to inhibit proliferation and activate apoptosis in human epithelial endometriotic cells. In particle Imesch demonstrated that his epigenetically acting drug inhibits VEGF transcription at low nanomolar concentration with high efficency. It works at the transcriptional level down regulated VEGF expression. Romidepsin reduced the level of HIF- α protein, indicating that VEGF mRNA expression may be related to the reduction of HIF- α protein levels. The issue of whether VEGF transcription is the primary target of romidepsin or if it acts preventing deacytalation of HIF- α , must still be solved. However Romidepsin, acting at a transcriptional level, could be more effective than other angiogenetic drugs, which inhibit the VEGF active form in targeting angiogenesis, and it can be considered a novel therapeutic candidate to counter endometriosis (Imesch P et al, 2011).

Gonadotrophin- releasing hormone agonist (GnHRa I) have been applied with success, in the treatment of endometriosis combined with the laparoscopic surgery. It leads to a reduction of ovarian hormone levels, to atrophy of endometriotic implants and it has antiproliferation and apoptotic effects. It reduce VEGF expression and it has been seen that after GnRHa treatment the concentration in peritoneal fluid of VEGF are significantly lower New molecule GnRHa II has been studied. Fengying et al demonstarted that this molecule can dose-dependently reduce VEGF protein secreted by ectopic and eutopic endometrial stromal cells cultured in vitro, and the inhibition effect is stronger than that of GnRHa I. GnRH II may reduce the secretion also of immune factors such as interleukin-8 and cyclooxigenase 2 relating to the incidence of endometriosis, suggesting for a antiproliferation and anti-inflammatory effect on endometrial cells (Fengying, H et al 2010).

Yilmaz showed (Yilmaz B et al, 2010.) the effect of metformin of endometriosis implants for his antioxidant characteristics and the beneficial effects on VEGF, and matrix metalloproteinases. In particular it reduces endometriotic implants in rats reducing VEGF levels.

Molecular therapies have been proposed as a treatment alternative for recurrent endometriosis. The use of concitionally replicative adenovirus (CRADs) has been explored for the therapy of disease. In particular Adenovirus constructed with the VEGF promoter controlling the expression of a marker gene have been evaluated in vitro culture of endometriotic cells. AdVEGFE1 replicates in a short-term culture of purified ectopic

endometriosis cells. The virus induces apoptosis in endometriotic cells in vitro (Rein DT. Et al, 2010). Ad VEGFE1 allowed specific replication and efficient killing of endometriotic cells.

Another approach, which has been recently been published (Essam –Eldin R et al, 2008) is the transfection of endometriotic cells by dominant negative estrogen receptor gene via Ad vector. Dominant negative mutants of the estrogen receptor are altered estrogen receptors forms that are unable to activate transcription of estrogen-responsive genes when estradiol binds them, resulting in decreased cell proliferation and increased apoptosis.

2.2 Endocrine Gland derived Vascular Endothelial Growth Factor (EG-VEGF)

Human endocrine gland derived vascular endothelial growth factor is a secreted angiogenetic mitogen growth factor expressed in the steroidogenic glands, ovary, testis, adrenal and placenta. It induces proliferation, migration and fenestration (formation of membrane discontinue) in capillary endothelial cells derived from endocrine glands. Human EG-VEGF is a 9.6 kDa protein consisting of 86 amino acid residues. Endocrine gland-derived VEGF (EG_VEGF) belongs to the prokineticin family. It is also known as prokineticn 1 (PK1). Although EG- VEGF is structurally distinct from VEGF, they induce similar angiogenic response in the ovary. The EG- VEGF acts through G-protein couplesd receptors, pkr1.

EG-VEGF was found to be expressed in non-endocrine tissues including endometrium: in human, it is highly expressed during the secretory phase of the menstrual cycle, when angiogenesis occurs. Lee et al evaluated the expression of EG-VEGF and its receptors in eutopic and ectopic endometrial tissues. A significant increase in molecule expression was found in the stromal cells of ectopic endometrium. It is possible that the stromal cells may synthesize EG-VEGF or that it is synthesize in the epithelial cells but is accumulated in the extracellular matrix of stroma. The endocrine gland-vascular endothelial growth factor, through its heparin- binding domain can, as VEGF, accumulated in the extracellular matrix (Lee K et al, 2010).

3. Cytokynes (IL-1, IL-6, IL-8, TNF- a)

Cytokines are small cell-signaling protein molecules that are secreted by several cells types and are a category of signaling molecules used extensively in intercellular communication. Cytokines can be classified as proteins, peptides, or glycoproteins; the term "cytokine" encompasses a large and diverse family of regulators produced throughout the body by cells of diverse embryological origin (Gilman et al., 2001). Virtually all nucleated cells, but especially endo/epithelial cells and resident macrophages (many near the interface with the external environment) are potent producers of IL-1, IL-6, and TNF-α (Boyle, 2005). Studies have reported elevated levels of several cytokines in the peritoneal fluid of women with endometriosis, thus implicating these proteins in the development and progression of endometriosis and endometriosis-associated infertility (Koninckx et al., 1998); (Harada et al.,2001); (Bedaiwy et al., 2003); (Kalu et al., 2007)._ Peritoneal fluid is derived from plasma transudate and ovarian exudates and in a small part from secretions of the mesothelial surface and tubal luminal fluid. Some studies suggest that the peritoneal fluid of women with endometriosis contains an increased number of activated macrophages and other immune cells that secrete various local products, such as growth factors and cytokines, which exert a paracrine action on endometriotic cells. (Harada et al., 2001)

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3.1 Cytokines and pathogenesis of endometriosis

Studies have shown the role of some cytokines in the implantation of ectopic endometrial tissue, and its progression and infiltration. In the implantation and growth of ectopic tissue, a primary role was attributed to several cytokines contained in the peritoneal fluid including interleukin IL-1,IL-6, IL-8, IL-12 and tumor necrosis factor-a (TNF-a) (Arici et al.,1996); (Iwabe et al., 1998); (Ho et al.,1997) Normally, peritoneal fluid contains leukocytes in concentrations of 0.5 to 2.0 3 106/mL, of which approximately 85% are macrophages (Syrop et al., 1987). Halme et al., (Halme et al., 1984) postulated that peritoneal macrophage activation may be a central contributor to the pathogenesis of endometriosis and activated macrophages in the peritoneal cavity of women with endometriosis (Vinatier et al., 1996) are potent producers of cytokines (Halme, 1989) ;(Fakih et al., 1987); (Rana et al., 1996). Thus, peritoneal fluid contains a rich cocktail of cytokines. Cytokines play a major role in the initiation, propagation, and regulation of immune and inflammatory responses. Immune cell activation results in a burst and cascade of inflammatory cytokines. These cytokines have pleiotropic and redundant activities that culminate in recruitment of numerous cell types to the site of inflammation (Harada et al, 2001). More, cytokines may regulate the actions of leukocytes in the peritoneal fluid or may act directly on ectopic endometrium, where they may play various roles in the pathogenesis and pathophysiology of endometriosis. Increased levels of cytokines in the peritoneal fluid of women with endometriosis may reflect increased synthesis of cytokines by peritoneal macrophages, lymphocytes, ectopic endometrial implants, or mesothelial cells of the peritoneum, all of which can produce cytokines (Tabibzadeh et al., 1989); (Betjes et al., 1993). The main source of cytokines is thought to be the macrophages, which originate in bone marrow, circulate as monocytes, and migrate to various body cavities. It seems that the cytokines playing the most important role in the endometriosis are : IL-1, IL-6, IL-8, IL-12, TNF- a.

Interleukin-1 (IL1) is one of the major proinflammatory cytokine found in the peritoneal fluid of women with endometriosis (Mori et al., 1991); (Taketani et al., 1992); (Fakih et al., 1987). This multifunctional cytokine was shown to stimulate the production of angiogenic factors by ectopic endometrial cells and therefore play a role in ectopic endometrial cell growth (Lebovic et al., 2000). Other studies pointed to a possible role for IL1 in endometriosis-associated infertility (Fakih et al., 1987); (Sueldo et al., 1990). Both ectopic and eutopic endometrial cells of women with endometriosis display an increased sensitivity to IL1, which results in an enhanced production of angiogenic, growth, and proinflammatory factors (Lebovic et al., 2000); (Akoum et al., 1995); (Akoum et al., 2002).Some previous studies showed that the increased endometrial and endometriotic cell responsiveness to IL1 may in part be due to a deficiency in the expression of interleukin-1 receptor type II (IL1R2) revealed in eutopic and ectopic endometrial tissues (Akoum et al., 2001);(Kharfi et al., 2002). The soluble IL1R2 levels were found to be reduced in the peripheral blood of women with endometriosis, which may account for the activation of peripheral blood monocytes in them (Kharfi et al., 2002). More, the IL1R2 has no signaling properties in contrast to the functional signaling IL1R1, which mediates cell activation by IL1 (Bossu et al., 1995); (Dinarello 2004); (Colotta et al., 1993). However, the membrane form of this receptor and the soluble form, which is shed by proteolysis from the cell surface (Cui et al., 2003); (Orlando et al., 1997), bind to IL1 and with higher affinity to IL1b, which is the circulating and the preferential ligand for IL1R2, in particular for its soluble form (Bossu et al., 1995). This inhibits the interaction of IL1 with its functional receptor type I and, consequently, IL1-mediated cell activation (Bossu et al., 1995); (Colotta et al., 1993); (Subramaniam et al., 2004);(Symons et al, 1995). Akoum et al. (Akoum et al., 2008) showed an imbalance in IL1/soluble IL1R2 levels in women with endometriosis suffering from infertility and pelvic pain and a relationship with endometriosis initial stages and infertility. This is in keeping with other findings showing a reduced expression of IL1R2 in the eutopic endometrial tissue of women with endometriosis, particularly in those who were infertile, and provide evidence for a deficiency in the regulation of IL1 actions at the local peritoneal level in initial endometriosis stages, which may result in increased cell reactivity and contribute to endometriosis development and the manifestation of its clinical symptoms.

Interleukin-6 (IL-6) is a pleiotropic cytokine that is produced by a variety of cell types, including monocytes, lymphocytes, fibroblasts, endothelial cells, and mesangial cells. It is said to mediate numerous physiological and pathogenic processes and acts on a wide variety of cells. IL-6 may also have important functions in reproductive physiology, including the regulation of ovarian steroid production, folliculogenesis and early events related to implantation. (Jacobs et al., 1992); (Akoum et al., 1996). Both eutopic and ectopic endometrium are known to produce IL-6 (Harada et al., 1997). Infact IL-6 belongs to the group of cytokines produced in increased amount by endometriotic cells both in basal and cvtokine-stimulated conditions (Akoum et al., 1996), (Tsudo et al., 2000). IL-6 in turn is able to increase the secretion of several other cytokines and promotes the activation of immune cells (Iwabe et al., 2002). Examining eutopic endometrium from patients with endometriosis, it was found an increased basal- and IL-1b stimulated production of IL-6 compared with patients without endometriosis. This suggests that the endometrial cells of women who develop endometriosis may function differently from those in women who do not develop this condition. Endometrial stromal cells were considered the critical cells in endometrial attachment to the mesothelial surface of the peritoneum and that endometrial epithelial cells fail to attach to the mesothelium (Scott et al., 1953). It has also been suggested that cellular adhesion itself stimulates chemokine expression (Smith et al., 1997). More interestingly IL-6 family cytokines, such as IL-6, IL-11, leukemia inhibitory factor, and oncostatin M, were shown to be potent stimulators of aromatase expression in adipose stromal cells in culture (Zhao et al., 1995). Yoshioka et al. (Yoshioka et al., 1999) reported that IL-6 inhibits proliferation of endometrial stromal cells derived from the secretory phase but not from the proliferative phase. In contrast, negative regulation by IL-6 was not observed in the stromal cells of endometriotic tissues, suggesting that the biological characteristics of endometriotic cells differ from those of eutopic endometrial cells. Bedaiwy et al. (Bedaiwy et al., 2002) reported that serum IL-6 and peritoneal fluid TNF- a, could be used to discriminate between patients with and without endometriosis with a high degree of sensitivity and specificity. On the other hand, other authors, such as Kalu (Kalu et al, 2007) have failed to confirm this. They did not find any significant differences in the concentration of IL-6 in the sera of women in the two groups and more, they try to explain this result, considering that the measurement of cytokine concentrations is complicated by the fact that they have very short half-lives and are never produced in isolation, but as a mixture which may have similar or opposing effects. So, it is possible that the circulating IL-6, IL-1 b, and TNF- a found by some investigators may be non-functional or antagonized by anti-inflammatory cytokines or cytokine inhibitors. While this may also be true of peritoneal fluid cytokines, the presence of activated macrophages does suggest that at least some of the cytokines are functionally active.

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• Interleukin-8 (IL-8) is a potent angiogenic cytokine produced by mesothelial cells, macrophages, and endometrial and other cells. Its concentration in the peritoneal fluid of patients with red endometriosis is found to increase as the size and number of active lesions increase (Iwabe et al.,1998). It stimulates adhesion of endometrial stromal cells to extracellular matrix proteins, matrix metalloproteinase activity and endometrial stromal cell proliferation in a dose-dependent manner, all of which can help to promote the implantation and growth of ectopic endometrium (Garcia-Velasco et al.,1999); (Ryan et al.,1995); (American Society for Reproductive Medicine, 1997); (Harada et al.,1997); (Punnonen et al.,1996); (Jolicoeur et al., 1998); (Akoum et al.,2000); (Pizzo et al., 2002) ; (Arci et al., 1998).

• Arici et al. reported that IL-8 is produced in the human endometrium in vivo, mainly in glandular cells (Arici A, et al.,.) and that this interleukin induces proliferation of endometrial stromal cell as a potential autocrine growth factor (Arici et al., 1998). Iwabe et al. (Iwabe et al.,1998) also found that peritoneal fluid levels of IL-8 significantly enhanced proliferation of stromal cells derived from ovarian endometriomas. Expression of IL-8 receptor type A messenger RNA was detected in endometriotic stromal cells These results suggest that IL-8 may promote the progression of endometriosis.

Tumor necrosis factor α (**TNF-** α is a cytokine involved in systemic inflammation and is • a member of a group of cytokines that stimulate the acute phase reaction. The primary role of TNF is in the regulation of immune cells. TNF is able to induce apoptotic cell death, to induce inflammation and to inhibit tumorigenesis and viral replication. It is considered the most representative cytokine involved in the pathogenesis of endometriosis: a clear positive association between the content of TNF- a in peritoneal fluid and the severity of endometriosis has been demonstrated (Overton et al., 1996). Numerous studies indicate that both activated macrophages and the ectopic endometrium itself are responsible for the abnormal production of TNF- a in the peritoneal fluid and that this cytokine is involved in the proliferation of endometriotic stromal cells (Iwabe et al., 2000). It acts through both direct and indirect mechanisms, and by mediating the proliferative effect of IL-8 (Sakamoto Y et al., 2003). Moreover, gene and protein expression of IL-8 in the stromal cells of endometriotic tissues are up-regulated by TNF- α (Iwabe et al., 2000), and TNF- α stimulates the proliferation of the endometriotic stromal cells. This stimulatory effect of TNF- a was abolished by adding anti-TNF- α antibody or anti-IL-8 antibody. Therefore, TNF- α may act on stromal cells by mediating the proliferative effects of IL-8. Expression of type I and type II receptors for TNF- a was observed in endometriotic stromal cells. This evidence suggests that TNF- a action mediated by IL-8 may not only be an initiating factor that facilitates adhesion of endometrial cells to the peritoneum but may also contribute to development and progression of endometriosis. Thus, the differential response of endometrial cells to TNF- a in women with and without endometriosis may reflect differential regulation of TNF-receptor expression or signaling by this cytokine. Braun et al. (Braun et al., 2002) published data suggesting that in women without endometriosis, endometrial cells do not implant in ectopic locations because normal apoptotic mechanisms are activated by TNF- α through the TNFR1 receptor and because the proliferation enhancing effects of TNF- α are inhibited by down-regulation of the TNFR2 receptor. Disruption or dysregulation of the normal, cyclical expression of these two TNF- a receptors on endometrium from women with endometriosis could create cells that can grow in the presence of high concentrations of TNF- α a possibility for which evidence is avalable (Ding et al., 2000). Harada et al. (Harada et al., 1997) and Iwabe et al. (Iwabe et al., 1998) found that the extent of superficial red

endometriotic lesions was related to increased levels of IL-6, IL-8, and TNF- a in the peritoneal fluid. Red lesions, such as red flame-like lesions, gland-like lesions, and red vesicles, were classified as active lesions of endometriosis because angiogenesis is more pronounced in red lesions than in black or white lesions (Wiegerick et al., 1993) and because early red lesions invade extracellular matrix (Spuijbroek et al., 1992). Braun et al. (Braun et al., 2002) also suggested that ectopic growth of endometrial cells and the physiological consequences of that growth in women with endometriosis may be retarded by agents that block the effects of TNF- a. Presumably, this could be achieved by blocking TNF- a production (e.g., administration of pentoxifylline or ciprofloxacin) or by blocking the effects of TNF- a on target tissues (e.g., administration of etanercept). The attenuation of the proliferation-enhancing activity in peritoneal fluid from women with endometriosis by etanercept that the athors observed, and results in an animal model of endometriosis using a recombinant human TNF-binding protein (D'Antonio et al., 2000;), support this idea. Blocking the effects of TNF- α on target tissues might be especially appropriate in patients with extensive or intractable disease and might be useful in the postsurgical adjuvant setting to reduce the likelihood of recurrence. Given the potential of TNF- α to play a prominent role in both the etiology and the pathogenicity of endometriosis, studies of such treatment are warranted

Interleukin-12 (IL-12) is naturally produced by dendritic cells (Kaliński Pet al., 1997), macrophages and human B-lymphoblastoid cells in response to antigenic stimulation. It is known as a T cell-stimulating factor, which can stimulate the growth and function of T cells. It stimulates the production of interferon-gamma (IFN-γ) and tumor necrosis factor-alpha (TNF-a) from T and natural killer (NK) cells, and reduces IL-4 mediated suppression of IFNγ. It acts on T and NK cells, inducing cytokine production, enhancing NK cell cytotoxic activity, and finally favoring generation of Th1 cell response, (Kobayashi et al., 1989) ;(Wysocka et al., 1995) so it seems to play an important role, in the controll of the endometriosis. Harada et al., (Harada et al., 2001) demonstrated that the concentrations of IL-12 in the peritoneal fluid are low, but detectable, regardless of the presence or absence of endometriosis (Zeyneloglu et al., 1998). Administration of IL-12 significantly prevented ectopic endometrial implantation in a murine model of endometriosis (Somigliana et al.,1999). A direct growth inhibitory effect on endometrial cells seems unlikely because endometrial cells do not express receptors for IL-12. A potential explanation for these results is that IL-12 enhances the growth and augments the cytolytic activity of both NK and T cells. These data support the idea that manipulation of cytokine activity in the peritoneal fluid is a novel management approach to controlling the establishment of endometriosis.

3.2 Cytokines and endometriosis related to infertility

Muscato et al. (Muscato et al., 1982) demonstrated that peritoneal macrophages phagocytized sperm in vitro and that macrophages from women with endometriosis were more active than those from women without the disease. Peritoneal fluid diffusing into the tubal and endometrial environment may affect sperm and their interaction with the oocyte. Studies showed that the peritoneal fluid of patients with endometriosis has detrimental effects on sperm function. Sperm motility (Curtis et al., 1993) ;(Drudy et al., 1994), acrosome reaction (Arumugam et al., 1994), gamete interaction (Coddington et al, 1992.), and ovum capture by tubal fimbriae (Suginami et al., 1988) have been studied. Aeby et al. (Aeby et al., 1996), using a penetration assay, showed that peritoneal fluid from patients with

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endometriosis impaired gamete interaction. In their study, the mean number of eggs penetrated by sperm mixed with peritoneal fluid from patients with endometriosis was significantly less than that observed in controls. These data suggest that substances in the peritoneal fluid of patients with endometriosis contribute to infertility by impairing sperm function. Peritoneal fluid from patients with endometriosis has frequently been shown to be toxic to the preimplantation embryo. A study demonstrated that medical treatment of endometriosis eliminated the embryotoxicity of the peritoneal fluid (Keenan et al., 1995). In this study, the levels of IL-1 and TNF- a were markedly reduced in the peritoneal fluid of women who had undergone medical treatment (danazol or intranasal buserelin for 4 to 6 months) of endometriosis. This finding supports the hypothesis that increased levels of cytokines in peritoneal fluid may be involved in the pathogenesis of endometriosisassociated infertility. One study demonstrated that the addition of human recombinant IL-6 to culture medium suppressed the rate of blastocyst formation of mouse embryos (Harada et al., 1997), suggesting that increased IL-6 in the peritoneal fluid of endometriosis patients may contribute to infertility by adversely affecting embryonic development. Moreover, other authors (Minici et al., 2008) showed that in endometriosis, the milieu surrounding the uterine cavity may be involved in impaired eutopic endometrial stromal cell decidualization, partially due to increased peritoneal levels of TNF- a. So they concluded that in endometriosis either intrinsic defects of endometrial stromal cell differentiation or the biochemical environment of the uterine cavity could concur to compromise the normal decidualization required for optimal implantation.

In conclusion, cytokines, which are produced by many cell types in peritoneal fluid, play a diverse role in constructing the peritoneal environment that induces the development and progression of endometriosis and endometriosis-associated infertility.

Intense basic research into the specific role of these cells and soluble factors may improve our understanding of endometriosis and result in novel therapies for endometriosis.

4. Transforming growth factor beta (TGF-ß)

Transforming growth factor beta (TGF- β) is a protein that controls proliferation, cellular differentiation and other functions in most cells. TGF- β is a secreted protein that exists in at least five isoforms called TGF- ß 1, TGF- ß 2 and TGF- ß 3. It was also the original name for TGF- ß 1, which was the founding member of this family. TGF-beta acts as an antiproliferative factor in normal epithelial cells and at early stages of oncogenesis. Some cells that secrete TGF- ß also have receptors for TGF- ß. This is known as autocrine signalling. Cancerous cells increase their production of TGF-β, which also acts on surrounding cells.Oosterlynck et al. (Oosterlynck et al.,1994) found increased TGF-b activity in the peritoneal fluid of women with endometriosis. Transforming growth factor-b may be a cytokine that inhibits NK activity in the peritoneal fluid of women with endometriosis (Oosterlynck et al., 1994). It may play a major role in the biological processes leading to establishment and maintenance of endometriosis., in fact TGF- ß is implicated in the gene expression, cell motility, proliferation, apoptosis, differentiation, immune responses and tumorigenesis (Derynck et al., 2001). TGF-b is abundantly and differentially expressed in the endometrium and is secreted by endometrial cells and macrophages into the uterine fluid where interaction with the preimplantation embryo is suspected (Jones et al., 2006).

Secretion of TGF-b into peritoneal fluid of women suffering from endometriosis suggests that they may be crucial in establishment and/or maintenance of endometriosis. Omwandho et al., showed that all TGF- ß and their high-affinity receptors were stagespecifically expressed in the human endometrium with highest levels around menstruation . Many researchers have reported staining of TGF-b1 and 3 in stromal and glandular cells (Chegini et al., 1994); (Gold et al., 1994); (Johnson et al., 2005); (Komiyama et al., 2007); (Gaide Chevronnay et al., 2008) and for TGF-b1 also in nerve fibres (Tamburro et al., 2003) and inflammatory cells specially in macrophages (Chegini et al., 1994); (Tamura et al., 1999); (Komiyama et al., 2007). TGF- ß 2 is more strongly expressed in stromal compared with glandular cells (Gold et al., 1994); (Bruner et al., 1995); (Gaide Chevronnay et al., 2008), although opposite staining intensity has been reported (Chegini et al., 1994). Localization of TbRII and RI was observed in both cellular compartments of the endometrium (Chegini et al., 1994; (Gaide Chevronnay et al., 2008) with stronger expression of TbRII than TbRI (Gaide Chevronnay et al., 2008) suggesting that TbRI might be a limiting factor for signal transduction in the endometrium or during endometriosis. TGF-b1 was found in the stromal cells (Johnson et al., 2005) and expression increased in the epithelial cells of endometriotic cysts (Tamura et al., 1999) and endometriotic nerve fibers (Tamburro et al., 2003). The TGF-b signal transducers Smad3, pSmad3, Smad4 (SMADs are intracellular proteins that transduce extracellular signals from TGF beta ligands to the nucleus where they activate downstream TGF- ß gene transcription) and the inhibitory Smad7 proteins were also observed in the endometrial stromal and epithelial cells (Luo et al., 2003). These observation suggest a role of the TGF-ß s in the normal function of the human endometrium. In fact is well known that TGF- ß sprevent the breakdown of the endometrial tissue (Tabibzadeh, 2002). This assumption is based on the observation that Lefty-2/EBAF (endometrial bleeding associated factor), a member of the TGF-b family, is dramatically up-regulated during endometriosis (Kothapalli et al., 1997) and antagonized TGF- ß signaling by inhibiting phosphorylation of Smad2 downstream of the TbRI (Ulloa et al., 2001). That Lefty-2 was noticeably more abundant in patients with endometriosis who did not conceive compared with those who became pregnant, suggested a role in implantation (Tabibzadeh et al., 2000). More, it has been shown that TGF- ß 1also induces contractions of decidual stromal cells (Kimatrai et al., 2003) and inhibits motility of stromal endometrial cells (Nasu et al., 2005). It is important to show how TGF- & 1stimulated DNA synthesis in epithelial cells at low concentrations, but inhibited DNA synthesis at higher concentrations in women with and without endometriosis (Meresman et al., 2003). Additional evidence showed that TGF- ß 1induces expression of FasL mRNA and protein in endometrial stromal cells (Garci-Velasco et al., 1999), possibly preventing apoptosis during transit to the peritoneal cavity. Another very important analysis showed that TGF- ß 1 represses the immune system (Shull et al., 1992) and the escape from immune surveillance is also important for adhesion of endometriotic cells in the peritoneum. Finally TGF- ßs participate in the initiation of menstruation via vasoconstriction, in menstrual tissue repair and in endometriosis. In a classic experiment, Luo et al. (2003b) demonstrated that the pretreatment with a GnRH antaonist resulted in further suppression of Smad3 in endometrial stromal cells but co-treatment with GnRH and TGF- ß 1or pretreatment with TbRII antisense partially inhibited TGF- ß 1-activated Smad3. Taken collectively, these observations suggest that GnRH may prevent endometriosis by altering expression and activation of Smads and interrupting TGF- ß receptor signaling.

5. Plasminogen activator (uPA) and matrix metalloproteinase systems (MMS)

5.1 Plasminogen activator system (PA)

The PA system includes a wide cluster of proteolytic enzymes for plasmin generation. Plasminogen is activated to plasmin by two types of activators, urokinase-type PA (uPA) and tissue- type PA (tPA). Whereas tPA is involved in the role in the control of intravascular fibrin degradation, uPA is mainly implicated in cellular proteolysis and migration. The activity of the PAs is regulated by specific PA inhibitors (PAIs). (Kruithof et al., 1995; Grancha et al., 1996; Heeb et al., 1987).

The PA system and its specific plasminogen activator inhibitors (PAIs) exert physiological and pathophysiological functions such as fibrinolysis, tissue remodelling and tumor invasion, signal transduction, cell adherence and cell migration (Harbeck et al., 2001).

Fernández-Shaw et al., firstly reported high levels of urokinase and plasminogen in ectopic endometrium as a more invasive nature of the endometriotic implants in the peritoneal cavità (Fernández-Shawet al., 1995); afterwards, Sillem confirmed an altered activation of plasminogen in endometrium from women with endometriosis that could lead to a higher proteolytic potential of retrogradely menstruated endometrial fragments with consecutive development of endometriotic foci (Sillem et al., 1997).

In situ hybridization studies performed by Bruse et al. showed that uPA mRNA seems to be up-regulated in both endometriotic glands and endometrial stroma from women with endometriosis (Bruse et al., 2005).

Moreover, Lembessis and coworkers reported an increase in uPA mRNA expression in endometriotic lesions compared to eutopic endometrium (Lembessis et al. (2003).

Despite contrasting data in vitro culture model (Guan et al., 2002) ; recently, Cosin reported an increase in uPA antigenic levels in endometrium from women with endometriosis (Cosin et al., 2010)

In relation to PA levels in eutopic endometrium from women with endometriosis, it has been suggested that a higher concentration of uPA in the endometrium might result in endometrial fragments with a higher potential to degrade the extracellular matrix after the implantation at ectopic sites (Spuijbroek et al., 1992; Bruse et al., 1998, 2004; Kobayashi, 2000).

5.2 Matrix metalloproteinase systems (MMPs)

Matrix metalloproteinases (MMPs) are a class of zinc-dependent endopeptidases involved in extracellular matrix remodelling (Matrisian, 1992).

Members of this family share high level of structural analogy and are secreted by several cell types as zymogens. In relation to substrate preference and protein-domain considerations, MMP family members have been categorized into subgroups that include gelatinases, stromelysins, collagenases, membrane-type (MT)-MMPs and 'other MMPs'.

The activity of metalloproteinases is tightly regulated, as these molecules are potent proteolytic enzymes, at differente steps: transcriptional level (by cytokines, chemicals, and

growth factors), post-translation modification and by a family of inhibitors: the tissue inhibitors of metalloproteinases or TIMPs (Matrisian, 1990).

Elevated cytokines may play a role in the establishment of ectopic endometrium in the peritoneal cavity by stimulating MMPs to remodel the mesothelial lining of the peritoneum thus allowing for tissue invasion.

MMPs are stimulated by cytokines and also by the protein Extracellular Matrix Metalloproteinase Inducer (EMMPRIN). Braundmeier et al., showed that IL-1beta stimulated MMP-1 protein secretion and mRNA levels in a time dependent manner (P < 0.05), MMP-2 mRNA in a time dependent manner and MMP-3 in a time and dose dependent manner. TNF-alpha stimulated MMP-1 and -3 protein secretion in a time dependent manner). Neither IL-1beta nor TNF-alpha treatment affected MMP-2 protein secretion. TGF-beta-1 inhibited MMP-1 and MMP-2 mRNAs at the highest treatment dose after 24 hr but there was no effect on protein secretion. TGF-beta-1 exerted no effect on MMP-3 mRNA or protein secretion (Braundmeier et al., 2010).

MMPs have been implicated in the endometrial remodelling during the menstrual cycle with higher levels during menstrual and proliferative phases and decreased levels during the secretory phase (Salamonsen and Woolley, 1996);

Montly, in the absence of pregnancy, degradation of the ECM is a critical step in the initiation of tissue breakdown that leads to menstruation (Marbaix *et al.*, 1996; Salamonsen and Woolley, 1996).

During the proliferative phase of the natural cycle, MMP-1, MMP-3 and MMP-9 are downregulated in the stroma (Hulboy *et al.*, 1997), presumably to allow endometrial stable growth. The expression of MMPs then decline in the early secretory phase and then increase during the late secretory phase in anticipation of the next proliferative phase. These modification are related to serum progesterone levels, which has led to the suggestion that endometrial expression of MMPs is under gonadal steroid hormone control. Critically, MMP-9 expression is highest in the menstrual phase endometrium when tissue breakdown occurs.

Moreover, several repots suggest that these proteases are also involved in the ectopic invasion of endometriotic cells associated with endometriosis (Cox et al., 2001).

Deregulation of peritoneal fluid cytokines levels of women with endometriosis show indicate that an altered immune system may play an important role in the pathogenesis of endometriosis. The invasion of ectopic endometrium into peritoneal mesothelium, in association of different angiogenic factors, requires matrix metalloproteinases (MMPs) for tissue remodeling. Several MMPs are differentially expressed in human uterine endometrium with menstrual endometrium showing the highest level of expression. (Braundmeier et al., 2010)

MMP systems closely interact with PA system, because plasmin is an active enzyme, which degrades a variety of extracellular matrix proteins and activates MMPs and growth factors (Murphy et al., 2000).

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The catalytic domains of all MMPs share high amino acid similarity and their active sites are extensively conserved (Lauer-Fields et al., 2009). As a consequence, differenciate between different MMPs activities is extraordinarily difficult. However, some members of these proteases showed a role in the pathogenesis of endometriosis.

Matrix metalloproteinase 7 (MMP7): MMP7 is secreted mostly from the endometrial epithelium cells during the receptive phase localized to endometrial glandular and luminal epithelium (Yanaihara et al., 2004; Zhang et al., 2005)

Moreover, MMP-7 has been shown to be the dominant metalloproteinase during the initial development of endometriosis in a baboon model (Fazleabas et al., 2002). In addition, a recent study clearly demonstrated that MMP-7 mRNA was identified in host peritoneal tissues during the development of endometriosis in a nude mouse model (Hull et al., 2008)

MMP-7 protein expression in epithelial cells was significantly higher in red peritoneal lesions compared with that of deep infiltrating endometriosis, ovarian endometriosis and black peritoneal lesions, in all phases of the menstrual cycle. MMP-7 protein expression may be down-regulated during the evolution of peritoneal endometriotic implants, as active red lesions transition into inactive black lesions (Matsuzaki et al., 2010).

Matrix metalloproteinase 5 (MMP5): The strongest MMP5 staining was seen in luminal epithelial cells, whereas endometrial glands frequently showed partial expression.

Both the gene chip expression analyses as well as PCR indicated strongly elevated transcript levels in most peritoneal endometriosis lesions. Moreover enhanced MT5-MMP expression has been detected in the eutopic endometrium from patients suffering from endometriosis. (Gaetje et al., 2007)

Matrix metalloproteinase 3 (MMP3): IMMP-3 is hormonally regulated during the menstrual cycle, with the highest levels of expression occurring during menses local regulation that is absent in the in vitro cultures. (Hulboy et al., 1997;.)

MMP-3 has not been well studied in endometriosis, however, studies suggest that retrogradely shed menstrual fragments, the putative precursors of endometriotic lesions, express high levels of MMP-3 (Koks et al., 2000).

Cox et al, demonstrated in a rat model that elevated MMP-3 expression by endometrial tissue leads to the establishment and progression of ectopic endometrial tissue growth. (KE. Cox, et al., 2001).

Significant expression differences were obtained for MMP3 in the ovarian endometriomas. The deregulation of the different genes, included MMP3 genes, may be responsible for the loss of cellular homeostasis in endometriotic lesions (Meola et al., 2010)

Matrix metalloproteinase 2 (MMP): Overexpression of stromal MMP-2 may play a role in the development of adenomyosis (Tokyol et al., 2009)

Matrix metalloproteinase 9 (MMP9): Eutopic endometrium of women with endometriosis compared with normal women showed an increased release of MMP-9, and a decreased release of its natural inhibitor, TIMP-1, at both the protein and the mRNA levels (Chen *et al.* 2004; Collette et al., 2004; Collette et al., 2006).

5.2.1 Genetic polymorphism of MMPs family

Genetic polymorphisms located in the promoter region of the MMP genes could lead to increased gene expression and could be associated with predisposition to endometriosis (Ye, 2006). Nevertheless, the genetic susceptibility of endometriosis in relation to MMPs polymorphism is very complex, because for several polymorphisms, allele frequencies were found to be significantly different according to ethnic origin (MMP2.1, MMP2.2, MMP3 and MMP12).

Borghese et al., investigated the role of MMP1, MMP2, MMP3, MMP7, MMP12 and MMP13 polymorphisms as endometriosis risk factors in a case – control study of patients affected by superficial, deep infiltrating or endometrioma in the Caucasian population. The study found a potential role for MMP12 -82 A/G and MMP13 -77 A/G combined polymorphisms, which modulate transcriptional activity, in superficial endometriosis. As no association was found with deep infiltrating endometriosis, this combination of polymorphisms might protect from a more in-depth penetration of tissues (Borghese et al., 2008). On the other hand, they did not find any correlation between endometriosis and MMP1, MMP2, MMP3, MMP7 polymorphism (Borghese et al., 2008). Data regarding the lack of association between MMP-1 e MMP-3 polymorphism and endometriosis susceptibility were confirmed in another study concerning the Italian population (Ferrari et al., 2006).

A case-control study in women of caucasian origin, evaluated the potential associations of MMP-2 and MMP-9 gene promoter region polymorphisms as well as MMP-2 promoter haplotypes with susceptibility to endometriosis. The results demonstrated that polymorphisms in MMP-2 (-735 C/T) and MMP-9 (-1562 C/T) were associated with elevated risk of endometriosis and that certain MMP-2 promoter haplotypes were more common in control group (Saare et al., 2010).

A genetic study regarding North Chinese women on three polymorphisms in the MMP-2 (MMP-2; -1306C-->T and -735C-->T) and TIMP-2 (TIMP-2; -418G-->C) genes found that the TIMP-2 -418C/C homozygote may be a protective factor against the development of endometriosis (Kang et al., 2008). A analogous study in the same population showed that MMP-7-181A/G polymorphism has a potential to be a susceptibility factor for endometriosis and adenomyosis while MMP-9-1562C/T polymorphism may not provide a useful marker to predict susceptibility to endometriosis and adenomyosis (Shan et al., 2006). On the other hand, an increase in the distribution of the MMP-9R279Q/P574R (2678G>A/4859C>G) and -1562C>T/R668Q (-1562C>T/5546G>A) haplotypes was significantly associated with endometriosis (Han et al., 2009).

5.2.2 Relevance of MMPs serum/urinary levels as diagnostic markers

The balance between MMPs and their inhibitors is preserved in the serum of women with endometriosis; however MMP-3 mRNA seems to be a promising peripheral blood marker that discriminates between patients with endometriosis and healthy subjects. Circulating mRNA for MMP-3 is significantly higher in patients with endometriosis than in control patients, regardless of the degree of severity. (De Sanctis et al., 2010).

Conversely, the clinical relevance of MMP-2 and MMP-9 as markers of endometriosis is controversial; some data report that serum concentrations of MMP-2, MMP-9, TIMP-1 and TIMP-2 cannot be considered to represent a valid measure of the severity of endometriosis

(Salata et al., 2008; De Sanctis et al., 2010). On the other hand a prospective, blinded, longitudinal study show that MMP-2, MMP-9, and MMP-9/neutrophil gelatinase-associated lipocalin were significantly more likely to be detected in the urine of patients with endometriosis than in controls (Beker et al. 2010).

Bruner-Tran et al. described progesterone treatment inhibits expression of MMP-3 and -7 in human endometrium and prevents the establishment of ectopic lesions in a nude mouse model (Bruner-Tran et al., 2002).

A study concerning human endometrium intraperitoneally transplanted into nude mice, demonstrated a significant suppression of MMP-2 transcription by all progestins tested, and a significant down-regulation of MMP-3 by dydrogesterone (Mönckedieck et al., 2009).

In conclusion, angiogenesis is proposed as an important mechanism for the pathogenesis of endometriosis. Different evidences support the hypothesis that the endometrium of women with endometriosis has an increased capacity to proliferate, implant and grow in the peritoneal cavity. Further studies are needed to better understand critical steps of the pathogenesis of endometriosis; nevertheless excessive endometrial angiogenesis suggests novel new medical treatments.

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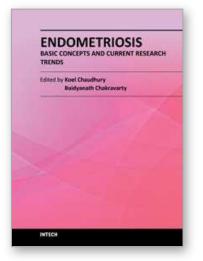
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This book provides an insight into the emerging trends in pathogenesis, diagnosis and management of endometriosis. Key features of the book include overviews of endometriosis; endometrial angiogenesis, stem cells involvement, immunological and hormonal aspects related to the disease pathogenesis; recent research reports on infertility, endometrial receptivity, ovarian cancer and altered gene expression associated with endometriosis; various predictive markers, and imaging modalities including MRI and ultrasound for efficient diagnosis; as well as current non-hormonal and hormonal treatment strategies This book is expected to be a valuable resource for clinicians, scientists and students who would like to have an improved understanding of endometriosis and also appreciate recent research trends associated with this disease.

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