Roles of Interferons in Human Endometrial Function

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Endometriosis is an estrogen-dependent inflammatory disease that develops in reproductive-aged women who experience pelvic pain and infertility. Even though endometriosis is not a new disease, its molecular etiology has not been clearly elucidated. Defects in the immune system might be one of the factors that promote endometriosis progression. For example, elevated levels of proinflammatory cytokines are associated with endometriosis. Interferon is one of the cytokines that is elevated in endometriotic tissues compared with normal endometrium.

endometriosis endometrium interferon

1. Introduction

Endometriosis is the growth of endometrial lesions outside the uterus, such as in the ovaries and peritoneal cavities ^[1]. Three different phenotypes of endometriosis are found in the pelvic cavity: superficial peritoneal endometriosis (SUP), ovarian endometrioma (OMA), and deeply infiltrating endometriosis (DIE) ^{[2][3]}. Red lesions are found on the surface of the peritoneum in SUP, while endometriotic tissues are invaginated into the ovary in OMA. In DIE, endometriotic tissue invades other organs, including the cervix and rectum ^[2]. Endometriotic lesions that are shown in red are early, active lesions with vascularization. The lesions become black after inflammatory reaction and scarification and eventually turn white ^[2]. Up to 5-10 percent of women of reproductive age experience symptoms of endometriosis [1]. Symptoms include mild to severe pelvic pain, infertility, painful urination or bowel movements, and abnormal menstrual flow. Surgical resection of the lesions is the primary treatment for endometriosis, but surgery cannot prevent disease relapse 4. The 2-year and 5-year post-operative recurrence rates are about 21.5 percent and 40–50 percent, respectively ^[5]. The median time of post-operative recurrence is about 30 months [6]. Additionally, endometriosis is an estrogen-dependent disease [1]. Therefore, hormonesuppressive drugs that block the synthesis or activity of estrogens (E_2), such as elagolix ^[8], progestin ^[9], and Danazol ^[10], have been applied to relieve endometriosis symptoms. Hormonal suppression treatment after conservative surgery significantly decreases the risk of endometriosis recurrence and its related symptoms [11]. However, hormone-blocking drugs cause severe adverse effects, such as postmenstrual symptoms, and off-target effects in other hormone-responsive organs, including bone and the brain, in endometriosis patients [4][12]. Therefore, alternative endometriosis treatments to replace hormonal therapy are in high demand.

Endometriosis is an estrogen-dependent proinflammatory disease ^{[1][7][13]}. Although the exact causes of endometriosis remain unknown, the theory of retrograde menstruation, which is an efflux of menstrual blood and cells via the fallopian tubes, is the well-accepted hypothesis by which endometriosis develops and progresses ^[1].

However, while 90% of reproductive-aged women experience retrograde menstruation, only 10% of them are diagnosed with endometriosis [14]. Therefore, in addition to retrograde menstruation, other factors are likely involved in the pathogenesis of endometriosis. Although the exact etiology of endometriosis has not yet been elucidated, the heritability of endometriosis has been estimated at 50 percent. In addition, a few twin studies have shown that the risk of endometriosis is increased in monozygotic twins compared with dizygotic twins, which indicates the possible contribution of specific genes or genetic alternations to endometriosis [15][16]. Meta-analyses of genome-wide association studies have identified single nucleotide polymorphisms (SNPs) associated with endometriosis risk. For example, SNPs in 11 different loci located in or near IL1A (interleukin 1 alpha), ETAA (ETAA1 activator of ATR kinase), RND3 (Rho family GTPase 3), NFE2L3 (nuclear factor, erythroid 2 like 3), WNT4 (Wnt family member 4), ID4 (inhibitor of DNA binding 4), CDKN2B-AS1 (cyclin-dependent kinase inhibitor 2B antisense RNA), VEZT (vezatin, adherens junctions transmembrane protein), GREB1 (growth regulating estrogen receptor binding 1), and FN1 (fibronectin 1) are correlated with the risk of endometriosis 17. Among them, IL1A and NFE2L3 are the genes associated with inflammatory pathways ^[17]. In addition, eight loci in the *IL1A* gene are related to the risk of endometriosis in Japanese population, which supports the involvement of immune and inflammatory responses in the development of endometriosis [18][19]. While those genes have not yet been validated as potential targets for endometriosis treatment, a recent study has identified and validated a gene named Neuropeptide S receptor 1 (NPSR1) as a potential target through DNA sequencing ^[20]. This study showed that deleterious low-frequency coding variants in NPSR1 are overrepresented in patients with familial endometriosis, especially in moderate/severe stages. NPSR1 is strongly expressed in glandular epithelial cells in eutopic and ectopic endometrium, while its ligand NPS is mostly found in the stroma. Inhibition of NPSR1 with the small molecular inhibitor SHA 68R prevents the release of proinflammatory Tumor Necrosis Factor (TNF) α by monocytes in vitro. In addition, the administration of SHA 68R relieves inflammation and pain in mouse models of endometriosis. NPSR1 expression is increased in several inflammatory diseases, such as inflammatory bowel disease and asthma ^{[21][22]}. In addition, NPSR1 is found in macrophages and T lymphocytes ^[23]. Sundman et al. show that stimulating monocytic THP-1 cells with proinflammatory cytokines TNF α and Interferon (IFN)y significantly increases NPSR1 isoform expression, suggesting that NPSR1 may be a key factor of proinflammatory cytokine signaling [24].

Genetic mutations in the exon-coding region of genes involved in cell adhesion and chromatin-remodeling complexes associated with endometriosis progression were identified ^[25]. There is no direct evidence showing how retrograde menstrual debris found outside the uterus is cleared in healthy women. However, the defective cytotoxic activity of Natural Killer (NK) cells and decreased phagocytic macrophage activity in endometriosis patients indirectly support the hypothesis that the impaired immunes surveillance system attributes to the pathogenesis of endometriosis ^{[26][27]}. In the peritoneal cavity of endometriosis patients, the immune cells are recruited and secrete excessive levels of proinflammatory cytokines (interleukin (IL)-1, IL-6, TNF, and IFNy), which promote disease development and progression ^{[1][28][29]}. Additionally, dysregulation of growth factors is involved in endometriosis. For example, the serum levels of vascular endothelial growth factor (VEGF), fibroblast growth factor (FGF), hepatocyte growth factor (HGF), insulin-like growth factor-I (IGF-I), granulocyte-macrophage colony-stimulating factor (GM-CSF), epidermal growth factor (EGF), and platelet-derived growth factor (PDGF) are higher in women

with advanced endometriosis than in women without endometriosis ^{[30][31][32][33]}. In addition to genetic mutations, alterations in immune and growth factor signaling pathways are also causal factors for endometriosis progression.

During retrograde menstruation, refluxed apoptotic endometrial cell debris in the peritoneal area is recognized by dendritic cells. Dendritic cells presenting endometrial autoantigens express increased type I IFNs to promote monocyte differentiation, dendritic cell survival, and cytotoxic T-cell activity, which may attack endometrial cells that present autoantigens in healthy women ^{[34][35]}. However, type 1 IFN-mediated cell death signaling is dysregulated in endometriotic cells in endometriosis patients, and thus, these cells can evade the host immunosurveillance system. Type II IFN (IFNy) fails to induce apoptosis in ectopic endometrial stromal cells, especially ovarian endometriotic cyst stromal cells, unlike in eutopic endometrial stromal cells with endometriosis have been well studied, the IFN signaling pathways in endometriosis are not fully understood ^{[1][35][37]}. IFN signaling also plays a significant role in normal endometrium, especially during pregnancy ^[38].

2. Interferons in Human Endometrial Function

2.1. Type I IFNs

The endometrium is the innermost lining of the uterus and changes during the menstrual cycle in response to ovarian hormones. The endometrium consists of columnar epithelial cells, the underlying stroma containing numerous embedded glands, immune cells, and vascular endothelial cells. This lining becomes thickened during the proliferative phase (i.e., before ovulation) and prepares for a possible pregnancy during the secretory phase (i.e., after ovulation). The thickened endometrium is shed during menstruation when progesterone and estrogen levels fall ^[39]. In conjunction with ovarian hormones, IFNs modulate pregnancy ^{[38][40]}.

Type I IFNs are highly expressed in the preimplantation blastocyst, trophoblast, and decidua trophectoderm during decidualization in mice ^[41]. In addition, preimplantation embryos secrete IFN α in vitro ^[42]. Therefore, type I IFNs might be crucial in embryo implantation and decidualization progression (**Figure 1**).



Figure 1. Role of Type I IFN in human pregnancy. Type I IFNs (IFN α , IFN β , and IFN ϵ) differentially modulate the downstream signals for successful pregnancy. This figure was created with BioRender.com.

IFNα stimulation synergically induces the expression of IFN-regulated gene 1 (*IRG1*) in humans at periimplantation along with E₂, although the underlying mechanisms are still unknown ^[43]. The transcription factor IFN regulatory factor 1 (IRF1) promotes implantation and is expressed in the human endometrium throughout the menstrual cycle and during peri-implantation in response to type I IFN exposure ^{[44][45]}. The levels of *IFNAR1* and *IFNAR2* are increased in the menstrual stage than in the proliferative phase in human endometrium ^[46]. IFNAR1 and IFNAR2 expressions are mostly found in the glandular epithelial cells but not in endometrial stromal cells, suggesting that glandular cells are the main cell type mediating type I IFN signaling in human endometrium ^[46]. In addition, ISGs, including interferon regulatory factor 8 (*IRF8*), interferon α-responsive protein (*IARP*), interferonactivated gene 202B (*IFI202b*), interferon γ-inducible GTPase 1 (*IIGP1*), interferon-stimulated gene 12 (*ISG12*), and *ISG15*, are upregulated in decidual tissues to enhance the decidualization of endometrial stromal cells in mice (**Figure 1**) ^[47].

Stimulation of human uterine epithelial cells with poly (I:C), a viral double-stranded RNA (dsRNA) mimic, increases IFN β and ISG expressions ^[48]. In addition, inhibition of IFN β signaling either with IFN β -neutralizing antibody or IFNAR2-blocking antibody partially prevents the increases of ISG expressions, including MxA, OAS2, and PKR upon poly (I:C) stimulation, indicating that epithelial IFN β exerts antiviral responses (**Figure 1**) ^[48].

While IFNE expression is widely found in the female reproductive tract, its expression in the endometrium is hormonally regulated during the menstrual cycle ^[49]. Unlike the vagina and ectocervix, in which IFNE is expressed in the basal and parabasal layers of the epithelium, IFNE is also detected in the surface of the endometrial luminal

epithelium, suggesting that IFN ϵ may play a role in immune protection from infection in the endometrium ^[49]. In female mice, IFN ϵ -gene-deleted mice (IFN $\epsilon^{-/-}$) are more susceptible to Herpes Simplex Virus (HSV)-2 and *Chlamydia muridarum* bacteria than wildtype mice ^[50].

In addition, vaginal concentrations of IFNɛ are lower in genital HSV-infected pregnant women than in healthy pregnant women ^[51]. The amniotic fluid concentration of IFNɛ is elevated in women undergoing spontaneous preterm labor with intra-amniotic infection than in women with no infection or sterile intra-amniotic inflammation ^[52]. The IFNɛ expression level is significantly elevated in the endometrium during the luteal phase than during the follicular phase. Progesterone receptor (PR) and its ligand progesterone regulate IFNɛ expression, in which progesterone stimulation interferes with the activation of IFNɛ promoter in endometrial epithelial cells in vitro and ex vivo ^[49]. In contrast, uterine IFNɛ expression is increased in estrogen-treated ovariectomized mice, which suggests hormonal regulation in IFNɛ expression ^[50]. The excessive expression of IFNɛ in the endometrium and its change during the menstrual cycle suggests that IFNɛ plays an essential role during pregnancy ^[49]. The vaginal concentration of IFNɛ is increased during pregnancy ^[51]. IFNɛ is gradually expressed in the myometrium from mid-to late gestation ^[52]. Therefore, IFNɛ exerts antiviral and anti-bacterial effects in the endometrium and during pregnancy (**Figure 1**).

2.2. Type II IFNs

IFNy, the only type II IFN, is highly expressed in human trophoblast cells in the first trimester, while its receptor IFNGR is found in the placenta throughout the pregnancy ^[53]. Decidualized human endometrial stromal cells treated with conditioned media from human trophoblasts upregulate IFN-related genes, including *IFNGR1* and *JAK2* ^[54]. IFNy is expressed in luminal and glandular uterine epithelial cells during the estrus stage in mice ^[55]. Additionally, IFNy expression is highly increased within implantation sites in mice, which suggests the possible involvement of IFNy in the implantation process ^[56]. Human uterine NK cells secrete cytokines, including IFNy, and IFNGR is found in endometrial epithelial cells ^{[57][58]}. Ablation of NK cells, IFNy, IFNGR, or STAT1 in mice causes defects in the decidual remodeling of spiral arteries that supply blood to the placenta. Moreover, the administration of IFNy to NK-cell knockout (KO) mice led to the recovery of decidual artery remodeling ^[59]. Therefore, IFNy may play a critical role in decidualization during the implantation process in humans.

2.3. Type III IFNs

In addition to type I and II IFNs, type III IFNs also modulate normal endometrial functions. For example, IFN λ 1 has an essential role in the immune defense of the placenta against viral pathogens, as human uterine epithelial cells and fibroblasts secrete IFN λ 1 after exposure to the synthetic dsRNA viral ligand poly (I:C) ^[60]. Sex hormones also modulate IFN λ 1 signaling because estrogen suppresses, but progesterone stimulates, IFN λ 1-induced ISG expression in uterine epithelial cells ^[60]. Therefore, IFN λ 1 might differentially modulate endometrial function based on menstrual cycle stage, pregnancy, and menopausal status.

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