Review Article

Multifaceted Functions of Prokineticins in Reproductive Cancers and Proposed Associated Therapies

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Abstract

Prokineticin 1 and prokineticin 2 are two secreted proteins that belong to the prokineticin family. The two ligands act via two G protein-coupled receptors, PROKR1 and PROKR2 to induce cell’s proliferation, migration, invasion and permeability. They mostly act on epithelial, endothelial and immune cells. Deregulations in the expression of the ligands and/or the receptors of this family have been associated with both tumor and non-tumor pathologies of the reproductive system. In these pathologies, prokineticins have been reported to be associated with strong angiogenic and inflammatory activities. While their direct involvement in some threatening reproductive non-tumor pathologies such as kallmann syndrome, polycystic ovaries, preeclampsia and fetal growth restriction are well established, their role and their consideration as potential targets in the tumors of the reproductive system are still debated. This review will address the multifaceted roles of prokineticins and their receptors in reproductive cancers. Especially, the review will address the role of the prokineticin system in two female cancers, the ovarian cancer and the gestational cancer, Choriocarcinoma. The two types of cancer differentially express the prokineticin ligands at their early developmental stages, leading to different overall responses to be adapted when antagonisation of the prokineticin system is proposed. The review will summarize recent advances in the understanding of the prokineticin system’s involvement in the reproductive cancers, discuss its multifaceted roles in relation to prokineticin’s actions and proposes associated therapies.

Keywords: Prokineticin; Reproductive cancer; Targeted therapies; Immune system; Prokineticin antagonists

1. Prokineticin Family

In mammals, the prokineticin family is composed of two members, the prokineticin 1 (PROK1) also called EG-VEGF (Endocrine Gland-derived Vascular Endothelial Growth Factor) and the prokineticin 2 (PROK2) also called Bv8 [1, 2]. EG-VEGF is transcribed from the prok1 gene composed of three exons located on chromosome 1 in 1p21 [3, 4]. The gene that transcribes PROK2 is located on the chromosome 3 at 3p13.

Prokineticins and their receptors are expressed in various tissues such as ovary, testis, adrenal gland, placenta, uterus, brain, digestive tract and bone marrow [5-7]. In many organs prokineticin’s expression has been shown to be dynamic throughout physiological processes such as, circadian rhythm [8] menstrual cycle [9] and throughout pregnancy [10, 11]. In these systems, prokineticins have been shown to be regulated by local effectors and/or endocrine hormones. In the ovaries PROK1 is regulated by β-hCG (human Chorionic Gonadotropin), estrogen and FSH [12-14]. In the placenta PROK1 has been shown to be regulated by hypoxia, hCG and PPARγ [10, 15, 16].

The prokineticins are secreted protein that act through their binding to two G protein-coupled receptors (GPCRs) [1, 2], the prokineticin receptor 1 (PROKR1) and the prokineticin receptor 2 (PROKR2). The prokineticin receptors were discovered simultaneously by three research teams [1, 2]. They belong to the GPCR family, with seven transmembrane domains that interact with heterotrimeric G proteins. PROKR1 and PROKR2 were initially called GPR73A and B, then receptors of prokineticin 1 and prokineticin 2, respectively. They share 85% homology in their amino acid sequence, but differ in their N-terminal parts [1, 2, 17].
Strong similarities were found in their transmembrane domains, suggesting similarities in their mechanisms of activation [17]. In human, the prokr1 and prokr2 genes include two exons and are located on two different chromosomes. The prokr2 gene is located on chromosome 2 at 2q14 and prokr1 gene is located on chromosome 20 at 20p13 [1]. The two prokineticins act as powerful ligands for both PROKR1 and PROKR2 receptors. Interestingly, differential and tissue-specific expressions were reported for the two receptors. PROKR2 is abundantly expressed in the brain and testes, while PROKR1 is mainly expressed in peripheral tissues such as the spleen, prostate, pancreas, heart and blood cells. In the reproductive system, high levels of both receptors have been reported in the testis, ovary, endometrium and placenta [6, 9, 12, 18, 19].

Several studies have been carried out to determine the affinity of the two ligands to their receptors. It clearly appears that PROK2 exhibits higher affinity for PROKRs (EC50 of 4.5 ± 0.8 nmol/L for PROKR1 and 6.4 ± 1.3 nmol/L for PROKR2) [20], whereas PROK1 exhibits a moderate affinity for both receptors (EC50 of 27.6 ± 8.2 nmol/L for PROKR1 and 52.2 ± 16.4 nmol/L for PROKR2). Distinct expression patterns of the prokineticins and their receptors have been reported in various tissues, providing the cue for their tissue-specific biological functions [5-7]. In addition, the differential pattern of expression of G-proteins plus the multiple types of G-proteins that can couple to these receptors further increase the functional complexity of the system. PROK1 and PROK2 bind to PROKR1 and PROKR2, which can be coupled to Gi, Gs and Gq to activate MAPK/Akt, induce cAMP accumulation and calcium mobilization [20].

2. Prokineticins and Diseases

Prokineticins were first identified as potent agents stimulating the muscle contraction of the gastrointestinal tract [14, 21]. Later, PROK1 was shown to mediate proliferation and differentiation of the enteric neural crest cells during the enteric nervous system (ENS) development [22]. Since their discovery, prokineticin proteins have been associated with two key biological functions, angiogenesis, including vascular remodeling, and inflammation [6, 23, 24]. Over the last two decades, research articles on the prokineticins have either studied their role in angiogenesis and neovascularization or in inflammation [6, 23, 24]. However, few studies have addressed combined actions of the prokineticins in a given physiological or pathological process.

A determinant factor that links the biology of prokineticins with vascular remodeling is the hypoxic environment which develops upon a vascular damage and which is known to up-regulate prokineticins. Importantly, both PROK1 and PROK2 promoter regions have response elements for the hypoxia inducible factor (HIF-1) [6, 23, 24]. In several studies, prokineticins have been reported as potent angiogenic factors and their receptors are highly expressed in both microvascular and macrovascular endothelial cells [6, 23].

PROK1 was reported as a potent angiogenic factor that promotes angiogenesis in various steroidogenic glands [4, 8, 10]. In relation to inflammatory processes, both ligands were reported to activate this process. However, unlike PROK1, PROK2 is more associated with this process as it is expressed in the bone marrow. PROK2 is also reported as the preferential prokineticin expressed by immune cells as it acts as a potent chemoattractant for monocytes and
neutrophils [24, 25]. It also induces survival, differentiation and activation of the granulocytic and monocytic lineages [26]. To fulfill the angiogenic and inflammatory functions, prokineticins and their receptors have been reported to be involved in the control of multiple processes including proliferation, differentiation, invasion and migration of the target cells. These effects are mainly represented by endothelial and epithelial cells for the angiogenic processes and by immune cells such as monocytes, macrophages and neutrophils for the inflammatory processes [26, 27].

While, the prokineticin system has been reported to be associated with numerous diseases, its association with the reproduction-associated pathologies is the most investigated [13, 28-30]. Since their discovery, multiple studies highlighted the direct involvement of the prokineticin system in the development of severe reproduction pathologies, such as the polycystic ovarian syndrome (PCOS), the congenital diseases, named Kallmann syndrome, the Hirschsprung disease [30-32] and multiple pregnancy-associated pathologies [6, 10, 16, 33-35]. Nevertheless, the majority of the studies in relation to the reproductive system emanated from their involvement in placental development, during pregnancy. In relation to pregnancy pathologies, PROK1 constitutes the master member. In fact, strong evidences established the direct involvement of PROK1/PROKRs system in physiological and pathophysiological aspects of pregnancy; these include clinical, in vitro, ex vivo, and in vivo studies [6, 10, 16, 33-36]. Through its receptors PROKR1 and PROKR2, PROK1 controls the proliferation and invasion of trophoblast cells to ensure the establishment of the fetomaternal circulation, along with direct effect on intravillous vasculature to ensure vascular branching and angiogenesis [6, 10, 11]. Beside its physiological effect, PROK1 and its receptors have been reported to be causative and/or compensatory factors in most of the threatening pathologies of pregnancy, such as preeclampsia and fetal growth restriction [37,38]. Also, PROK1 has been designated as a new noninvasive biomarker of successful embryo implantation and oocyte competence in conventional in vitro fertilization-embryo transfer [34, 39, 40].

2.1 Prokineticin’s and Reproductive Tumor Diseases

Due to their wide spectrum of actions, notably as potent angiogenic and immunoregulatory factors and to the fact that both PROK1 and PROK2 are activated by the most potent environmental cue of angiogenesis and tumorigenesis, the oxygen tension; investigators have searched for the PROKs/PROKRs system involvement in cancer development in a variety of tissues [4, 22, 36, 41-49]. While numerous studies reported deregulations of PROK1, PROK2 and/or their receptors in a wide range of cancers, few studies brought evidences for their direct involvement in cancer development and progression [22, 36, 41, 46, 50].
Table 1: Reports an updated list of the different type of tumors that were investigated in relation to the potential involvement of the prokineticin family in their growth, progression and/or aggressiveness.

As exemplified through this listing, the prokineticin system appears to be deregulated in many types of cancers. However the involvement of the ligands and/or their receptors is tightly dependent on the type of cancer, on its severity and its localization within the affected tissue. Because most of the examined cancers in relation to the prokineticin system emanate from descriptive studies involving cohorts enrolled at advanced stages of cancer, it...
is difficult to conclude on whether the deregulation of PROK system is a cause or consequence of the cancer. Nevertheless, most of the data confirmed the deregulation of prokineticins at these stages, suggesting their tight association with cancer’s aggravation. In these cases, their deregulations should rather be considered as consequences of the disease.

Because most advanced and far-reaching studies on prokineticins and cancer have been investigated in the reproductive system, the following sections will report the updated investigations on prokineticins in relation to prostate, testis, ovarian and placental cancers. These subsections will report the associated deregulated member, its role in the development and/or progression of the related cancer and will propose associated therapies, when applicable.

2.1 Prokineticins in the prostate and testicular cancers
PROKs and their receptors are expressed in the testis and the prostate. In the prostate, PROK1 is localized in the glandular epithelial cells and PROK2 as well, as the two receptors are expressed in the prostate epithelial cells [47]. In the testis, PROK1 is predominantly expressed in testosterone-producing Leydig cells, whereas PROK2 expression is restricted to primary spermatocytes [4, 6, 10]. The PROKRs are also expressed in vascular endothelial cells in the testis [4, 13]. Importantly, adenoviral delivery of PROK1 or PROK2 to the mouse testis resulted in a potent angiogenic response [4], suggesting that PROKs contribute with other factors, such as VEGF-A, in the maintenance of the integrity and the proliferation of the blood vessels in this organ. The exact role of each of these receptors and their implication in the physiology of prostate and testis functions remain to be elucidated.

In relation to prostate cancer, the PROK1 expression has been reported to be elevated in the prostate compared to healthy individuals. These levels were correlated to the cancer’s malignancy [47]. Thus, the levels of PROK1 could be useful for prostate cancer outcome evaluations and can be used as a target for prostate cancer treatment in the future. Regarding the testis cancer, few studies investigated prokineticins involvement in the development and or progression of this cancer. In a study comparing PROK1 levels of expression in different testicular tumors, including Leydig cell tumors, seminomas and non-seminomatous germ cell tumors, it has been reported that PROK1 is predominantly expressed in Leydig cell tumors. The PROKRs have also been reported to be expressed in the testicular endothelium, suggesting an involvement of the PROKs/PROKRs system in neoplastic testicular angiogenesis [48].

2.1.1 Prokineticins in the prostate and testicular cancers

2.1.2 Prokineticins in the ovarian cancer
In the ovary, the representative and most abundant member is the PROK1, as numerous studies demonstrated that PROK2 is weakly or not expressed in the female reproductive organs [51]. PROK1, as well as its receptors have been reported to be strongly involved in physiological and pathophysiological processes of the ovary. In this organ, the expression of PROK1 fluctuates according to the ovarian cycle and is mainly associated with the secretary phase with a direct role in the maintenance of corpus luteum vascularization [12]. PROK1 is secreted during the follicular phase by granulosa cells of the primordial and primary follicles. Upon ovulation, PROK1 is secreted by granulosa lutein cells and its expression rises throughout the luteal phase with a peak during the very late stage [12]. Interestingly, PROK1 mRNA levels are also elevated in apoptotic tissues, especially during the corpus luteum regression or in the atretic follicles [18]. To fulfil this
pattern of expression in the ovary, PROK1 has been reported to be controlled by a panel of key ovarian regulatory factors, such as hCG, estrogen, FSH, oxygen tension and thrombin [12]. These events being associated with inflammation, it has been demonstrated that PROK1 triggers an onsite recruitment and activation of monocytes [18]. Study of gene expression profiles comparing VEGF, the major factor of the ovarian angiogenesis, and EG-VEGF revealed that their profiles rarely overlapped and seem to exhibit complementary type of functions during the ovarian cycle [18].

Given the above-described roles for PROK1 in the angiogenic and in the inflammatory processes in the ovaries, it was expected that deregulations in the PROK/PROKR system could be associated with the development of severe ovarian diseases. In this context, several studies reported prokineticins deregulation in the Polycystic ovarian syndrom (PCOS) [53]. Moreover, implication of prokineticins has also been investigated in ovarian cancer.

Ovarian cancer is the 5th cause of cancer deaths in women and it has the highest rate of death among gynecological cancers [54]. Malignant tumors that originate from the epithelium account for more than 90% of ovarian cancers. Among these is the high grade serous ovarian cancer; it accounts for nearly 70% of all ovarian cancers and is considered as the most threatening one [55]. Despite substantial survival benefits, chemotherapies are highly toxic and numerous patients develop resistance within few months after the first treatment. Therefore, there is still a need for efficient and less harmful treatment for this cancer [55].

Despite the fact that PROK1 is no longer detectable after menopause in the ovaries, some studies reported PROK1 positive cells at the tumors' periphery [56]. Two other studies reported significant correlations between PROK1 increased expression and the aggressiveness of the cancer, suggesting its potential consideration as a therapeutic target for this cancer [41,46]. On the contrary, Zhang et al. 2003 demonstrated that PROK1 mRNA levels decreased in the later stages of ovarian tumor development compared to premenopausal ovaries or early stages carcinomas [49]. Furthermore, in the same study, it was reported that PROK1 was secreted by the tumor microenvironment and especially by tumor-infiltrating T cells [49]. Importantly, the observation that PROK1 was mainly expressed by immune infiltrating cells in the ovary has been substantiated by other studies demonstrating that prokineticins could not only be expressed by tumor cells but also by immune cells in the tumor microenvironment [57]. Altogether, these findings strongly suggest that the balance between the angiogenic and the pro-inflammatory functions of the prokineticins in a given cancer may grandly contribute to the development and or progression of that cancer. While the role of the PROKs in the angiogenic process that accompany the early stages of ovarian cancer is strongly proposed, their expression by the local immune cells suggests that these proteins could be considered as specific targets to attenuate inflammatory associated reactions in the ovarian cancer.

2.1.3 Prokineticins in the placental cancer
Since the first ontology study on prokineticins, published by Ferrara et al in 2001, the human placenta appeared as a strong site of PROK1/PROKRs expression [3]. Publications following this first identification confirmed the strong expression of the PROKs, especially that of PROK1 and its receptors in the human placenta [13, 29, 58]. However,
most of these studies considered one gestational age that often corresponded to placentas collected at term pregnancies [29, 58]. Since 2006, more comprehensive studies [6, 10, 11, 23, 30, 33, 36, 37] reported complete profile, pattern, and regulations of the prokineticins during human pregnancy. Also, these studies investigated their involvement in the development of the most threatening pathologies of pregnancy [6, 10, 11, 30, 36, 37]. Briefly, those findings demonstrated that PROK1 was highly expressed during the first trimester of pregnancy, with the strongest localization in the syncytiotrophoblast and Hofbauer cells (placental macrophages). PROKR1 was mainly expressed in non-differentiated cells, while PROKR2 prevails in the differentiated cells, such as the syncytiotrophoblast and the extravillous trophoblasts (EVT). These studies also demonstrated that both PROKR1 and PROKR2 were expressed by placental micro- and macro-vascular endothelial cells [23]. Through its actions on PROKR1 and PROKR2, PROK1 has been reported to control trophoblast proliferation, migration and invasion. Its actions on the endothelial cells mainly concerned its effect on their proliferation and permeability [10, 11].

Because PROK1 is a circulating protein, its levels were examined throughout pregnancy and used to compare normal and pathological pregnancies. Circulating levels of PROK1 in non-pregnant women are about 50 pg/ml. These amounts increase five times during the first trimester of pregnancy to reach 250 pg/ml and significantly fall during the second and third trimesters of pregnancy to reach 70 pg/ml [23]. Importantly, deregulated PROK1 levels were observed in threatening pathologies of pregnancy such as preeclampsia and fetal growth restriction [11, 37]. In relation to these conditions, PROK1 is proposed as a new diagnostic and/or prognostic biomarker [11, 37].

Given the direct involvement of PROK1 and its receptors in the control of key aspects of placental development during the first trimester of pregnancy and its deregulation in pregnancy pathologies, the PROK1/PROKRs appeared as potential targeted proteins. While no study has yet addressed this topic in relation to pregnancy pathologies, such as preeclampsia and FGR, a recent study reported the involvement of PROK1 and its receptors in the development and progression of gestational choriocarcinoma (CC) and proposed therapeutic options through the targeting of the receptors [36].

Gestational CC is a malignant trophoblastic tumor that can develop from a normal or abnormal pregnancy [59]. Most often, this malignant lesion appears following complete (CHM) or partial hydatidiform mole (PHM), a spontaneous abortion, or an ectopic pregnancy [59]. Patients who acquire CHM or PHM have a high risk of developing CC. However, this risk is much higher for CHM (20%) compared to PHM (1.5%). It is well established that the progression of MHC towards a CC is governed by the acquisition by trophoblastic cells of a hyperproliferative character [60]. This phenomenon is reflected in a second step by the appearance of cellular clusters, which acquire a migratory and invasive phenotype. CC is a very metastatic tumor due to the invasive and intrinsic property of trophoblastic cells. Diagnosis and monitoring of CC progression are currently based on the measurement of circulating levels of β-hCG (human chorionic gonadotropin), secreted by syncytiotrophoblast cells [61].

As reported above, PROK1 acts as a novel placental growth factor which controls trophoblast proliferation, invasion and is upregulated by hypoxia and by hCG, two key parameters of tumorigenesis. Altogether, these data suggested that the PROK system might be involved in the development and or
progression of this cancer. In this context our laboratory recently deciphered the role of PROK1 and its receptors in the development of this cancer through, a clinical study using a distinctive cohort of CHM and CC patients; an in vitro study using the human choriocarcinoma cell line, the JEG-3; and an in vivo study using a recent animal model of CC. This study demonstrated that both placenta and circulating PROK1 levels were significantly increased in CHM and CC; that PROK1 increased JEG-3 proliferation, migration and invasion in 2D and 3D culture systems and that these observations were substantiated using the developed animal model [36]. To better characterize the type of receptor involved in these effects, non-peptide specific antagonists for PROKR1 and PROKR2 were employed [8, 62]. Importantly, our data showed that the antagonization of PROK1 receptors reduced JEG-3 tumorigenesis in vitro and in vivo and that the PROKR2 antagonist was more potent in mediating these effects [36]. Because, the actual measurements of circulating β-hCG have been reported to be associated with the generation of false positives, requiring some patients to undergo invasive and unnecessary therapeutic procedures such as chemotherapy, hysterectomy, and other surgeries [50], the study by Traboulsi & al offers the possibility to consider circulating PROK1 as potential diagnostic biomarker for CC progression. Also, this study proposes the antagonization of its receptors as potential therapy to treat gestational CC. This may provide safe and less toxic therapeutic option compared with the currently used multi-agent chemotherapies.

3. Antagonisation of the Prokineticin System as a Therapeutic Opportunity for Reproductive Cancers

The prokineticin system is equally involved in angiogenic and inflammatory processes, which are two key routes that drive cancer development and progression. In relation to angiogenesis, PROKs and especially PROK1 inhibition has been proposed as combinatorial option of therapies that target the most angiogenic factor, the VEGF. This was very attempting, because the effects of both factors in regards to the angiogenic processes were somewhat complementary. Beyond this important aspect, patients treated with anti-VEGF therapies show strong refractoriness [63], suggesting that exploration of new anti-angiogenic therapies that target less ubiquitous and more tissue-specific factors such as prokineticins might lead to better results. This can be applicable for reproductive cancers that affect the ovary and the placenta.

In regards to the prokineticin system involvement in inflammation, growing literature is addressing their local role in regulating immune responses. To date, these studies mainly reported the involvement of PROK2 in these functions [64]. In the case of ovarian cancer, PROK2 was shown to be expressed by components of the immune system, especially the myeloid cells and macrophages [49, 63]. These cells have been reported to represent a crucial cell subpopulation responsible for the refractoriness to anti-VEGF treatment [49, 63]. These data strongly suggest that therapies targeting not only PROK1 for its pro-angiogenic effects, but also PROK2 for its mediated refractoriness responses can advance the treatment of cancers that overexpress these proteins, such as the ovarian cancer. Figure 1 reports a cartoon that illustrates these conclusions. The panel A shows an active vascularized tumor secreting, among other angiogenic and pro-inflammatory factors, the prokineticin ligands (PROK1 and PROK2) and expressing their receptors. The increased secretion of prokineticin is potentiated by the tumor hypoxic environment. The secreted prokineticins are part of the first signals that allow immune cells mobilization towards the tumor’s
microenvironment. Once at the tumor site, the immune cells will in turn, contribute to the exacerbation of prokineticins secretion, inducing tumor cells proliferation, migration and invasion. Panel B shows the same phenomenon but in the presence of PROKR1 and/or PROKR2 antagonists. In this case, the prokineticin receptors will be occupied by the antagonists, which consequently leads to a significant decrease in all prokineticins’ mediated effects. In this schematic, blocking antibodies are also proposed as targeted therapy.

Figure 1: Reports a proposed model for tumor development in the absence or presence of targeting molecules (antagonists or antibodies) of PROK receptors.

Importantly, several studies have shown that the use of antagonists or antibodies against PROK2 in vivo, significantly decreased tumor growth in poorly and highly vascularized cancers [49, 63]. These effects were similar to those obtained with anti-VEGF treatments and the combination of both treatments was sometimes even more efficient [49, 63]. Through the inhibition of the prokineticin signaling, the tumor vasculature is no longer expanding; it acquires necrotic properties and exhibits a decrease in the production and secretion of pro-inflammatory cytokines.

Altogether these findings, strongly propose that therapies targeting prokineticins will not only be efficient to attenuate their pro-angiogenic activities but also to decrease their exacerbation of inflammatory responses within the tumor.

4. Conclusion and Perspectives
To date, the proposed therapies that target the prokineticin system are mainly based on the antagonization of its receptors and have only been used in vitro, ex vivo and in

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animal models [30, 65, 66]. Although more in vivo and translational studies are still needed to consider the use of these antagonists in human, they have shown beneficial effects in in vivo models. In fact, both antagonists exhibit analgesic effects and control the nociceptive process that is mediated by inflammatory pathologies [67]. More importantly, the treatment of the gravid CC mice with antagonists allowed the maintenance of the gestation with fewer resorbed fetuses compared to non-treated mice [36]. Altogether, these data advocate a strong therapeutic potential for these two antagonists when used solely or in combination with actual anti-angiogenic or anti-inflammatory molecules.

Because growing literature demonstrate that angiogenic factors are often involved in local immunosuppressive effects in the tumor microenvironment, one can speculate that this can also be the case for prokineticsins as they are strong angiogenic factors and are highly expressed by monocytes that generate myeloid dendritic cells, involved in immnossuppressive effects. Thus, the antagonistiaon of the prokinetin signaling can serve to control two arms of tumor development, angiogenesis and immunosuppression.

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**Conflicts of Interest**

The authors declare no potential conflicts of interest.

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