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Does IGF-1 play a role in the biology of endometrial cancer?

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ABSTRACT

Insulin-like growth factor 1 (IGF-1) is a mitogen which plays a key role in regulating cell proliferation, differentiation, and apoptosis. It belongs to the family of proteins also composed of insulin-like growth factor 2 (IGF-2), two types of membrane receptors (IGF-1R and IGF-2R), 6 binding proteins (IGFBP 1–6), hydrolyzing proteases, and reactive molecules binding proteins, which regulate the activity of growth factors. Disturbances in the functioning of IGFBP/IGF/1GF1R can lead to induction of carcinogenesis, which has been demonstrated in breast, prostate or colon cancers. Findings evaluating the role of IGF-1 in endometrial cancer biology are ambiguous and contradictory. Therefore, in the present study, we analyzed the role of IGF-1 in the process of carcinogenesis of endometrial cancer, based on the available literature.

Key words: endometrial cancer, insulin-like growth factor-1, carcinogenesis

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INTRODUCTION

Insulin-like growth factor-1 (IGF-1), also known as somatomedin 1, is a mitogen which plays a pivotal role in regulating cell proliferation, differentiation, and apoptosis. IGF-1 is a member of a protein family which also includes insulin-like growth factor 2 (IGF-2), two types of membrane receptors (IGF-1R and IGF-2R), 6 binding proteins (IGFBP 1-6), hydrolyzing proteases, and other reactive molecules binding proteins which regulate the activity of growth factors [1]. Disturbances in the functioning of the IGFBP/IGF/1GF1R system may lead to the induction of carcinogenesis, which has been demonstrated in various types of malignancies, i.e. breast, prostate, or colon cancer. Both, increased activity of the IGF-1/IGF-1R complex and lowered IGF-BP expression have been mentioned among the promoting factors for neoplastic processes [2]. IGF-1 is believed to exert its mitogenic effect on various cells by stimulating their proliferation and inhibiting their apoptosis via endocrine, paracrine and autocrine mechanisms [1]. Results of various studies which evaluated the role of IGF-1 in the biology of endometrial cancer (EC) are ambiguous and conflicting.

PHYSIOLOGICAL FUNCTION OF IGF-1

IGF-1 is predominantly produced by hepatocytes as a response to growth hormone (GH) activity and released to the serum of peripheral circulation blood. GH/IGF-1 cooperation is the main regulator of postnatal growth of the body, whereas IGF-2 plays a vital role during fetal development, and its synthesis is independent of GH. Insulin-like growth factors may also be synthesized de novo in the cells of various tissues (ovary, endometrium, breast, lung), where — being mitogenic factors — they participate in cellular growth processes and elicit anabolic effects. The primary role of IGF-1 consists in promoting cell and tissue growth and division [3-5]. The literature offers a significant amount of evidence that molecules from the insulin-like growth factors participate in proliferation and differentiation of normal endometrial cells, as well as embryo implantation. Various studies have demonstrated the presence of IGF-1, IGF-1, and all 6 IGFBP proteins in the endometrium during a normal menstrual cycle [6]. IGF-1 circulates in blood either free or bound to low- or high-IGF affinity binding proteins. Six IGF-1 binding proteins (from IGFBP-1 to IGFBP-6) have been identified so far. Over 90% of the circulat-

Corresponding author:

Dominika Majchrzak-Baczmańska Department of Surgical, Endoscopic and Oncologic Gynecology, Polish Mother's Memorial Hospital — Research Institute, Lodz, Poland Rzgowska St. 281/289, 93–338 Lodz, Poland tel.: +48 42 271 11 31, e-mail: majchrzak.dominika@gmail.com ing IGF-1 is bound to IGFBP-3 and ALS (acid-labile subunit), resulting in the formation of a 150 kDa molecular mass complex. The remaining part forms complexes with other proteins and less than 1% is unbound, which constitutes the active form of IGF-1. The effect of IGF-1 on cells of various tissues is executed via its binding to a specific receptor. There are two types of receptors which demonstrate affinity to IGF-1, namely IGF-1R and IGF-2R. However, IGF-2R binds IGF-1 to a lesser degree as compared to IGF-1R, thus revealing higher affinity to IGF-2 [1].

HYPOTHETICAL ROLE OF IGF-1 IN THE BIOLOGY OF ENDOMETRIAL CANCER

Research on the role of IGF-1 in the biology of endometrial cancer was initiated on the basis of a hypothesis which had its roots in the correlation between endometrial cancer and the following conditions: obesity, type II diabetes, and the mitogenic effect of estrogens on the endometrium. Type 1, whose development depends on abnormal stimulation of the endometrium by endogenous as well as exogenous estrogens, accounts for a significant number of sporadic EC cases [7]. Studies have demonstrated that estrogens (mainly E2) stimulate cell proliferation by inducing autocrine IGF-1 synthesis and expression of IGF-1 receptor in the uterus [8]. Additionally, IGF-1 encoding gene has been detected to undergo expression mainly in the follicular phase and early luteal phase of a menstrual cycle [9, 7]. Activation of the mitogenic effect of IGF-1 on the endometrium is the consequence of E2-dependent local production of IGF-1. Progesterone counteracts proliferation of endometrial

cells, which is induced by estrogens via stimulation of the local synthesis of 17β-hydroxysteroid dehydrogenase and estrogen sulfotransferase. As a result of their expression, E2 is converted to a less active estrone (E1) and estrone sulfates, which are quickly expelled from the cells and the body. Additionally, as a result of progesterone activity, the synthesis of the IGF-1 binding protein (IGFBP-1), which blocks IGF-1 activity in the endometrium by controlling IGF-1 availability for its cellular receptors, is stimulated. Estrogen-induced production of IGF-1 molecule, in the absence of IGFBP-1 synthesis which depends on progesterone expression, results in abnormal proliferation of endometrial cells [7, 10]. Recent reports in the literature have confirmed a correlation between EC and insulin resistance, hyperinsulinemia, and type II diabetes, which frequently constitute direct consequences of obesity. Insulin is believed to exert mitogenic effects on various cell types, especially cancerous cells, as a result of binding to a receptor or via other, non-receptor mechanisms, e.g. changing expression of IGF-1 and its binding proteins, or blocking of antiproliferative activity of progesterone. Insulin has been shown to stimulate mitogenic processes also by suppressing IGFBP-1 expression. Hyperinsulinemia may lead to lowered IGFBP-1 expression, even in the presence of progesterone, which probably results in a blockage of its protective activity [11].

IGF-1 EXPRESSION IN TUMOR CELLS AND RESULTS OF IN VITRO STUDIES

The results of *in vitro* studies demonstrate that IGF-1 participates in the process of EC carcinogenesis, both by induc-

Table 1. IGF-1 — in vitro findings										
Parameter	Cell population	Experiment	Result of the experiment	Role in carcinogenesis?	Ref.					
IGF-1	HEC-1B	IGF-1 incubation with HEC-1B	KCC induction KCC 1 expression	Yes	[12]					
mRNA: IGF-1 IGF-1R	RL95-2, KLE, HEC, ISH	Evaluation of IGF-1 synthesis in the cells Evaluation of mRNA expression: IGF-1 IGF-1R aIR3 incubation with the cells	Autocrine IGF-1 synthesis Autocrine stimulation of the cells by IGF-1 DNA synthesis induction in response to aIR3	Yes	[13]					
IGF-1	HEC-1A, KLE	IGF-1 incubation with HEC-1A and KLE	Induced mitogenic activity of the cells	Yes	[14]					
IGF-1 E2	ISH (ER+)	E2 incubation with ISH (ER+) Evaluation of IGF-1 activity after cell incubation with E2 IGF-1 incubation with ISH (ER+)	Incubation with E2: \rightarrow induction of cell proliferation \rightarrow induction IGF-1 secretion Incubation with IGF-1 \rightarrow induction of cell proliferation	Yes	[15]					
IGF-1 VEGF	HEC-1A (ER–)	IGF-1 incubation with HEC-1A	Incubation with IGF-1 $\rightarrow \uparrow$ mRNA VEGF activity	Yes	[22]					
IGF-1 IGF-1R IGFBPs	ISH, MCF-7	Cell incubation with tamoxifen	↑ISH proliferation Induction of IGF-1R phosphorylation	Yes	[26]					

able 2. IGF-1 — tumor tissue expression								
No. of samples	Parameter	Expression in EC tissues	Additional information	Ref.				
26 — total	IGF-1R	⇔	↑ IGF-1R binding affinity in tumor tissues	[23]				
59 — EC 10 — EH 7 — NE	IGF-1R IGF-1 IGF-2R	↑ ↑	*	[20]				
152 — VM (7 — EC) 63 — NT	IGF-1R	ſ	*	[24]				
30 — EC 30— NE	mRNA IGF-1	↑	*	[16]				
15 — EC 45 — NE	IGF-1	⇔	*	[17]				
30 — EC 30 — NE	mRNA IGF-1 IGF-1 (p) IGF-1R (p)	↓ ↔	*	[21]				
11 — EC 42 — NE	mRNA IGF-1R mRNA IGF-1	↑ ↑ type l EC ↓ type ll EC	*	[18]				
13 — EC 10 — NE	IGF-1	↑ EC cell proliferation	Inc. of IGF-1 with cells from the investigated samples was performed	[19]				

⇔the same in cell tissues and normal healthy endometrial samples, ↑ increased, ↓ lowered, EC — endometrial cancer, EH — endometrial hyperplasia, inc. — incubation; NE — normal endometrium, VM — various malignancies, NT — normal tissues, (p) — protein product

ing proliferation of tumor cells and stimulating the process of neo-angiogenesis (Table 1) [12–15]. However, the results of in vivo studies are not as unambiguous (Table 2) [16–21].

Studies on the population of HEC-1B cells of EC revealed that induction of KCI-cotransport (KCC) is the result of IGF-1 activity. Also, IGF-1 was found to lead to KCC 1 expression, which affects tumor invasion ability [12]. Reynolds et al., in their study of EC cells (RL95-2, KLE, HEC, ISH), detected the presence of autocrine IGF-1 synthesis and mRNA IGF-1 and IGF 1R expression in all cell populations. Inhibition of DNA synthesis in response to cell incubation with IGF-1 receptor-blocking antibody (aIR3) was observed in three cell lines (HEC 1A, KLE, ISH). These authors found autocrine stimulation of EC cells by IGF-1 in three out of four populations of the investigated cells [13]. Pear et al., noted induction of time-dependent mitogenic activity of EC cells as a result of incubation of the cell populations (HEC 1A and KLE) with exogenous IGF-1. IGF-1 was found to play a regulatory role in the processes of EC cell growth. The authors postulated IGF-1 to mediate in estrogen-dependent proliferation of EC cells [14]. Kashima et al., in their study on EC cell population (ISH) expressing ER, concluded that dose-dependent E2-induced proliferation of the cancerous cells occurs via the MAPK3/1 signaling pathway as a result of autocrine stimulation of IGF-1. They also analyzed EC tissues and normal endometrial samples and reported MAPK3/1 expression and phosphorylation both, in tumor samples and healthy endometrium, although MAPK3/1 expression was statistically significantly higher in EC cells [15]. Bermont et al., conducted

an experimental study on estrogen receptor-negative EC cells in order to assess IGF-1 role in the process of tumor neoangiogenesis. They found a dose- and time-dependent increase in VEGF mRNA expression, and an increased concentration of its protein product after 24 and 48 hour-incubation of the HEC-1A cell line with exogenous IGF-1. These authors suggested that estrogen-dependent activation of the IGF-1 autocrine loop may result in tumor growth by stimulating the process of neoangiogenesis [22]. The presence of IGF-1R receptors were demonstrated both, in EC tissues and healthy endometrium (Table 2) [20, 23, 24]. Regardless, statistically significantly higher receptor binding affinity was found in EC tissues as compared to healthy endometrium. Also, IGF-1R binding affinity was proven to increase in response to decreasing tumor differentiation [23, 25]. Additionally, tamoxifen, which is used in breast cancer therapy and which is a risk factor for the development of EC, has been demonstrated to lead to EC cell proliferation by inducing IGF-1R phosphorylation [26]. Furthermore, a dysregulation of genes encoding the components of the IGF-1 autocrine system and the ER receptor in the endometrium was also detected, which in consequence might lead to the development of EC [27].

On the other hand, Soufla et al., in a study of 30 postoperative EC preparations and 30 normal endometrial samples from the same patients, found no statistically significant differences in the mRNA IGF-1 transcription levels in both groups. On further analysis, these authors detected both, overexpression and lowered mRNA IGF-1 expression (39% and 56% of the cases) in tumor tissues as compared to the transcript level of that factor in normal healthy endometrium. Regardless, they emphasize that the presence of mRNA IGF-1 overexpression constitutes an important element in the oncogenesis of EC because it is associated with the process of tumor neoangiogenesis [16]. Toole et al., observed lowered IGF-1, ERa, ERb, and PR (progesterone receptor) expression in EC tissues as compared to healthy endometrial samples [17]. Maiorano et al., in a study of postoperative EC preparations and healthy endometrial samples, found a lowered mRNA IGF-1 expression in tumor cells as compared to healthy tissues. In turn, the level of IGF-1 and IGF-1R protein products, measured by immunohistochemistry, was the same in tumor cells and healthy endometrium. These authors postulate that perhaps the IGF-1 system is not directly involved in the carcinogenesis of EC [21]. Roy et al., reported mRNA IGF-1R overexpression in all of the investigated EC tissues, whereas increased mRNA IGF-1 was detected only in cancers developing on the basis of an endometrial hyperplasia with atypia — type 1 EC. Lowered mRNA IGF-1 expression was observed in type 2 EC. Thus, these authors postulated the existence of two different mechanisms which activate IGF-1R in endometrial cancer: ligand-dependent in type 1 EC and ligand-independent in

type 2 EC [18]. Reynolds et al., in their study of 13 EC preparations and 10 healthy endometrium samples, found that incubation of tumor tissues with IGF-1 results in higher EC cell proliferation. Interestingly, these authors observed that tissue proliferation of healthy endometrium is inhibited by IGF-1, EGF and TGF-alfa, and stimulated by cell incubation with Ab528 and a IR3 antibodies. The situation was reversed in case of EC cells. Based on their findings, these authors claimed the existence of a different, growth factor-dependent mechanism of tumor cell and heathy endometrium proliferation [19]. Pavelić et al., observed an increased IGF-1, IGF-1R, and IGF-2R expression in EC tissues, especially in FIGO stage III and IV, as compared to healthy endometrium, and endometrial hyperplasia with atypia. They found that specific inhibition of IGF-1R, and IGF-2R activity resulted in inhibition of tumor cell proliferation. They implied that the local IGF-1 autocrine loop may play an important role in tumor progression [20].

SERUM IGF-1 AND ENDOMETRIAL CANCER

Researchers, inspired by the results of EC cell and tissue tests, started to investigate serum IGF-1 in women with endometrial cancer. However, these studies are scarce and the results remain inconsistent (Table 3) [10, 28–35].

Table 3. Serum IGF-1 concentration								
No. of patients	IGF-1 concentration	Additional information	Study oversight	Ref.				
166 — EC 315 — controls	÷	Detailed Prospective study	*	[10]				
288 — EC 392 — controls	⇔	Detailed analysis	*	[29]				
73 — EC 108 — controls	⇔	Detailed analysis	*	[28]				
174 — EC 136 — controls	÷	Detailed analysis	*	[30]				
692 — EC 1723 — controls	*	No link between IGF-1 gene polymorphism and risk for EC	*	[36]				
32 — EC 18 — controls	Ļ	Impaired glucose tolerance and hyperinsulinemia in EC women might have affected the results	CVD not included	[32]				
250 — EC 465 — controls	Ļ	↑ insulin levels in EC women might have affected the results	CVD not included	[51]				
84 — EC 84 — controls	Ļ	*	CVD not included	[33]				
56 — EC 40 — controls	Ļ	*	CVD, impaired glucose tolerance, diabetes not included	[31]				
24 — EC 12 — EH with atypia 31 — EH without atypia 52 — controls (2 groups)	î	*	CVD, HRT, HC not included	[35]				
18 — EC 8 — controls	î	*	CVD not included	[48]				

⇔the same in cell tissues and normal healthy endometrial samples, ↑ increased, ↓ lowered, EC — endometrial cancer, EH — endometrial hyperplasia, CVD — cardiovascular diseases, HC — hormonal contraception, HRT — hormone replacement therapy

Four studies found no association between serum IGF-1 and EC biology [10, 28-30]. Lukanova et al., in a multi-center prospective study in 481 women (166 - study group, 315 — controls), reported no link between the risk for EC and serum IGF-1 levels. They found no statistically significant differences between serum IGF-1 concentration and EC as compared to healthy women, regardless of their menopausal status at the time of blood collection (most women were postmenopausal at the time of testing) [10]. Three other authors, in detailed analyses of large groups of patients, also found no statistically significant differences between serum IGF-1 concentration of women with EC as compared to healthy controls, or an association between that parameter and the risk for EC [28-30]. McGrath et al., in a large multi-center study including 2014 subjects (692 — women with EC, 1723 — controls) found no statistically significant correlation between IGF-1 gene polymorphism and the risk for EC. These authors imply that genetic variability in IGF-1 has no effect on the occurrence of EC [36].

Importantly, serum IGF-1 is synthesized mainly in the liver, and its level is affected by numerous endogenous — genetic predispositions, and exogenous factors — diet, physical exercise, medication use, including HRT, BMI, chronic disease (DM, cardiovascular and other conditions), as well as age [37–47]. Authors of the abovementioned publications adjusted their results to the factors which might potentially affect the results of their analyses. Also, the studies took into account the menopausal status of the study participants as the analysis of serum IGF-1 levels in women with breast and ovarian cancers demonstrated a correlation between elevated serum IGF-1 levels and the risk for developing these cancers only in young patients, i.e. < 55 years of age, while no such association was found for postmenopausal women [48–50].

Some studies detected statistically significantly lower or higher serum IGF-1 levels in women with EC as compared to healthy controls [31–35, 51].

Four teams found statistically significantly lower IGF-1 levels in subjects with EC as compared to healthy women [31–35, 51], but they did not account for the possibility of cardiovascular diseases in the investigated populations (common in older, postmenopausal patients), what might be a serious study limitation due to a proven link between lowered serum IGF-1 level and these pathologies [40, 42].

Additionally, Rutanen et al., in their study on total serum IGF-1 level, found a statistically significantly higher incidence of impaired glucose tolerance and hyperinsulinemia in women with EC as compared to controls, which — even according to these authors — might have affected the findings of the study [32]. Gunter et al., also reported elevated serum insulin levels in patients at higher risk for developing EC cancer, which also might have influenced the final results of their analysis [51]. Studies indicate lowered serum total IGF-1 levels in women with hyperinsulinemia and impaired glucose tolerance [43, 44]. Serum free IGF-1 levels in women with type II diabetes are statistically significantly higher as compared to healthy populations [44].

The study by Baloglu et al., generated particularly interesting results as these authors not only did not observe lower serum total IGF-1 levels in women with EC as compared to controls, but also noted decreased concentrations of that factor as the disease progressed [serum free IGF-1 levels were lower in women with myometrial infiltration (24 cases) as compared to infiltration-free women (32 cases)]. The results were surprising even for the authors, who stated they might have reflected disturbances in the functioning of the autocrine IGF-1 system in the uterus [31]. However, Lukanova is of the opinion that the locally produced IGF-1, whose synthesis is dominated by estrogens, participates in tumor biology, and serum IGF-1 level produced mainly in the liver as a response to GH activity is not a good marker of autocrine IGF-1 synthesis in the endometrium [10]. Besides, numerous factors affect the levels of serum IGF-1, including cardiovascular diseases, impaired glucose tolerance, elevated insulin levels, which often precede symptomatic diabetes type II, and that was not taken into consideration by these authors. Therefore, it remains unclear whether the obtained results indeed reflect the lowering IGF-1 level as the disease progresses, or whether they are caused by other reasons. The main limitation of the study by Baloglu et al., was the fact that they did not include patients with advanced-stage EC in their analysis, so the question whether the tendency for the serum IGF-1 level to drop would also be confirmed in more advanced disease, or disappear altogether, remains unanswered. Moreover, bearing in mind that most studies have confirmed the role of local IGF-1 loop in the biology of EC, the inverse correlation between serum IGF-1 level and disease advancement is unlikely to result from the process of endometrial carcinogenesis.

Two teams observed elevated serum IGF-1 levels in EC women as compared to controls [34, 35]. Waksmański et al., noted statistically significantly higher serum IGF-1 levels in women with EC and hyperplasia with atypia as compared to patients with normal endometrium. However, these authors failed to consider the possibility of HRT or hormonal contraceptive use, or the existence of cardiovascular diseases, which might have significantly affected the obtained results. That oversight questions the reliability of their findings [35]. Various authors have demonstrated that oral HRT leads to statistically significant decrease in serum IGF-1 levels, probably as a result of direct inhibition of its synthesis in the liver by the orally ingested estrogen [47]. Similarly to Waksmański et al., Ayabe et al., reported statistically significantly higher serum IGF-1 levels in post-

menopausal Japanese women with EC (18 cases) as compared to controls (8 cases). They did not observe statistically significant differences in IGF-1 levels among premenopausal women with EC ac compared to controls (5 — study group, 19 — controls). However, they did not take into account the possibility of coexisting cardiovascular diseases, and the sample size was particularly small. Therefore, their study may not constitute a reliable proof for undisputed role of serum IGF-1 in the biology of EC. Furthermore, as was suggested by the authors themselves, higher serum IGF-1 levels in postmenopausal Japanese women with EC might have been the result of varying parameter values between different races [48]. Several studies have reported that serum IGF-1 levels may be determined by ethnicity [52]. The majority of the abovementioned studies investigated Caucasian women, and only the study by Gunter et al., included 24 Asian subjects (7 — study group, 17 — controls) out of 715 all participants, although these authors did not mention any differences in IGF-1 concentration depending on the race [51]. Moreover, there is no consensus among researchers as to whether differences in serum IGF- levels are dependent on ethnicity [52].

CONCLUSIONS

On the basis of the abovementioned analysis of the literature, it seems safe to conclude that serum IGF-1 most probably does not participate in the process of endometrial carcinogenesis. The IGF-1 autocrine loop, which is present in the uterus, is believed to play a more significant role. Endometrial cancer affects mainly older, postmenopausal women, which might be an analogy to ovarian and breast cancers where, in case of postmenopausal patients, local IGF-1 system, rather than the factor present in the serum of the peripheral circulation, plays a role in tumor biology. Hypothetically, one cannot exclude the possibility that some women with EC present elevated serum IGF-1 levels, de facto resulting from the disease, as compared to normal healthy women, in which case that factor may participate, together with locally produced IGF-1, in the process of tumor oncogenesis. The presence of low serum IGF-1 concentration in EC women as compared to controls, which is supposed to result from the neoplastic process and show a dysregulation of IGF-1 system functioning in the uterus, seems to be rather unlikely, especially taking into account tumor biology, as well as the results of various studies which demonstrate the mitogenic effect of that factor on EC cells. On the other hand, some in vivo studies of EC tissue preparations dismissed the possibility that insulin-like growth factor 1 plays a significant role in tumor biology. Thus, it is possible that neither serum IGF-1 nor local autocrine system, which is involved in the physiological processes of normal healthy endometrium, participate in the oncogenesis of endometrial cancer.

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