Synergism between PGC-1α and estrogen in the survival of endometrial cancer cells via the mitochondrial pathway

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Abstract: Accumulating evidence shows that peroxisome proliferator-activated receptor gamma coactivator-1α (PGC-1α) is involved in the progression of hormone-related cancers, and there may exist an association between estrogen and PGC-1α. Notably, emerging evidence has led to considerable interest in the role of PGC-1α in endometrial cancer development. However, whether the synergism exists between PGC-1α and estrogen for regulating mitochondrial function to promote the development of endometrial cancer remains largely unknown. Here, we show that: 1) knockdown of PGC-1α attenuates the survival of endometrial cancer cells by inducing cell apoptosis through the mitochondrial pathway; 2) estrogen remedies the PGC-1α efficiency-induced decline of endometrial cancer cell viability; and 3) estrogen modulates the mitochondrial function to inhibit the PGC-1α deficiency-induced apoptosis in endometrial cancer cells. Collectively, these results demonstrated that the synergism between PGC-1α and estrogen was required for the survival of endometrial cancer cells, which was dependent on the mitochondrial pathway.

Keywords: PGC-1α, estrogen, endometrial cancer, survival, mitochondrial pathway

Introduction

Endometrial cancer is the most common gynecological malignancy. Approximately 80% of endometrial cancer cases are type I tumors, which are associated with a history of unopposed estrogen exposure or other hyperoestrogenic risk factors. Cell proliferation and apoptosis resistance might be mechanisms by which estrogens promote the progression of endometrial cancer. However, a series of reports have documented an association between antiestrogen therapy and the development of endometrial carcinoma. Therefore, an improved understanding of molecular pathogenesis of endometrial cancers may identify biomarkers for successful therapeutic strategies.

Estrogens have a broad spectrum of physiological functions and mediate their biological effects by binding to specific intracellular receptors, estrogen receptor-α (ERα) and ERβ. Recently, some interesting studies have indicated that ERs could localize to and play important roles in mitochondria. Estrogen protects the mitochondrial function to improve the function of heart. Furthermore, estrogen activates the mitochondrial biogenesis through the regulation of peroxisome proliferator-activated receptor gamma coactivator-1α (PGC-1α) in cardiomyocytes. PGC-1α is a multifunctional transcription coactivator that cooperates with nuclear receptors to regulate genes involved in mitochondrial function. It has been reported that PGC-1α could regulate ERα transcriptional activity in vitro and in vivo. Moreover, PGC-1α enhanced abnormal local estrogen production by stimulating aromatase activity in endometriosis. Therefore, these data provide a clue...
that there might exist an association between estrogen and PGC-1α in the regulation of mitochondrial function.

The critical role of PGC-1α in the regulation of cell metabolism and mitochondrial biogenesis has led to speculation that reduced PGC-1 function could be involved in the metabolic reprogramming undergone by cancer cells. Accumulated evidence indicated that PGC-1α was also involved in the progression of hormone-related cancer. Indeed, it was overexpressed in various estrogen-dependent diseases such as breast and ovarian cancers. Silencing of PGC-1α in breast cancer cells suspended their invasive potential and attenuated metastasis. PGC-1α could induce apoptosis in human epithelial ovarian cancer cells through a PPAR-γ-dependent pathway. Our previous study suggested that upregulation of PGC-1α and estrogen-related receptor γ (ERRγ) in endometrial cancer might be a requirement for cancer cell energy metabolism, which contributes to the development of endometrial cancer. Therefore, the similar effect of PGC-1α and estrogen on the mitochondrial function offered a potential mechanism that the synergism between PGC-1α and estrogen can be used for regulating mitochondrial function to promote the development of endometrial cancer. But the potential mechanism remained to be unknown.

In the present study, we evaluated the functional interaction between PGC-1α and estrogen and their impact on mitochondrial function in endometrial cancer cells. Our results suggested that the synergism between PGC-1α and estrogen was required for the survival of endometrial cancer cells, which was dependent on the mitochondrial pathway.

Materials and methods

Cell lines and culture

Human endometrial cancer (HEC-1-A) and Ishikawa cell lines were purchased from the Shanghai Institute of Cell Biology of Chinese Academy of Sciences (Shanghai, People’s Republic of China). The HEC-1-A cells were respectively cultured as adherent monolayers in McCoy’s 5A medium (Thermo Fisher Scientific, Waltham, MA, USA) containing 10% fetal bovine serum (HyClone, Logan, UT, USA) at 37°C and 5% CO2 in a saturated humidified incubator. The Ishikawa cells were cultured as adherent monolayers in Roswell Park Memorial Institute-1640 medium (HyClone) containing 10% fetal bovine serum at 37°C in a 5% CO2 humidified incubator. The study protocols were approved by the Ethics Committee of China Medical University, Shenyang, People’s Republic of China.

Transfection

Two small interfering RNAs (siRNAs) targeting different sites of human PGC-1α mRNA (GenBank accession no NM_013261.3) were designed and synthesized by the Genepharmacy Inc. (Jiangsu, People’s Republic of China) to knock down the PGC-1α expression, and a no-silence siRNA was synthesized as a negative control to eliminate the possibility of nonspecific silencing effects. All the siRNAs mentioned previously are shown in Table 1.

The cells were seeded into six-well plates and incubated overnight to allow for full extension and adherence before transfection. The cells were transfected by using DharmaFECT transfection reagent (Thermo Fisher Scientific) according to the manufacturer’s instruction. In all, 10 μL of 5 μM siRNA was added to 190 μL of serum-free medium. At the same time, 5 μL of DharmaFECT transfection reagent was added to 195 μL of serum-free medium. Then the contents of each tube were mixed and incubated for 5 minutes at room temperature. The content of siRNA was added to the DharmaFECT transfection reagent, and then 1,600 μL of antibiotic-free complete medium was added. Finally, the culture medium was removed from the wells of the six-well plate and 2 mL of the appropriate transfection medium was added to each well. The cells were harvested 72 hours after transfection for future experiments. For estrogen treatment, 48 hours after transfection, the cells were treated with 10 nmol/L 17β-estradiol for the indicated time followed by the individual experiments.

Table 1 siRNA sequence used in this study

<table>
<thead>
<tr>
<th>Name</th>
<th>siRNA sequence</th>
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<tbody>
<tr>
<td>siRNA1</td>
<td>Sense: 5′-GUCCGAGUCACAACACUAATT-3′</td>
</tr>
<tr>
<td></td>
<td>Anti-sense: 5′-UAGUGUUGUGACUGCGACTT-3′</td>
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<tr>
<td>siRNA2</td>
<td>Sense: 5′-GGACAGUGAUAUCAAGUATT-3′</td>
</tr>
<tr>
<td></td>
<td>Anti-sense: 5′-AUUACUGAAAUCAGUGUCCTT-3′</td>
</tr>
<tr>
<td>No-silence</td>
<td>Sense: 5′-UCUCCGAAACGUGACGUGTT-3′</td>
</tr>
<tr>
<td></td>
<td>Anti-sense: 5′-ACGUGACAGCUCCGAATT-3′</td>
</tr>
</tbody>
</table>

Abbreviation: siRNA, small interfering RNA.

Cell proliferation assay

Cell proliferation was determined by Cell Counting Kit-8 (CCK-8; Dojindo Molecular Technologies, Kumamoto, Japan). A total of 2×10⁴ cells/well were seeded into the 96-well plates and treated with siRNAs with or without estrogen. In all, 10 μL of CCK-8 solution was added to each well and incubated for 2 hours. The absorbance of each well was measured at 450 nm using a microplate reader.

Annexin V–fluorescein isothiocyanate/propidium iodide double staining assay

Cell apoptosis of endometrial cancer cells was analyzed by flow cytometry using an Annexin V–fluorescein isothiocyanate
(AV-FITC) apoptosis detection kit (Beyotime Biotechnology, Shanghai, People’s Republic of China). After treatment with estrogen for 24 hours, different groups of cells were stained with the apoptosis detection kit according to the manufacturer’s instructions. The cells from each sample were suspended and incubated in 195 μL of AV-FITC binding buffer and 5 μL of AV-FITC for 20 minutes at room temperature in the dark. Then the samples were centrifuged at 1,000×g for 5 minutes and resuspended in 190 μL of binding buffer and 10 μL of propidium iodide (PI). The samples were analyzed using FACScan flow cytometry (BD Biosciences, San Jose, CA, USA) with at least 10,000 events recorded for each condition.

**Measurement of mitochondrial membrane potential**

After treatment with estrogen for 24 hours, the changes in mitochondrial membrane potential were assessed by Mitochondrial Membrane Potential Assay Kit with JC-1 (Beyotime Biotechnology). After treatment, the cells were washed with phosphate-buffered saline (PBS) and incubated with the medium containing 20 μg/mL JC-1 at 37°C for 20 minutes. The cells were collected and washed twice with PBS and then analyzed by flow cytometry.

**TUNEL assay**

Cell apoptosis was also examined using TUNEL (terminal deoxynucleotidyl transferase dUTP nick end labeling) assay. The one-step TUNEL Apoptosis Assay Kit (Beyotime Biotechnology) was used to stain cells according to manufacturer’s instructions. Cells were seeded into six-well plates containing disinfected and sterilized glass coverslips. After 72 hours of transfection, cells were fixed with 4% paraformaldehyde for 20 minutes at room temperature. Then the samples were centrifuged at 1,000×g for 5 minutes and resuspended in 190 μL of binding buffer and 10 μL of propidium iodide (PI). The samples were analyzed using FACScan flow cytometry (BD Biosciences, San Jose, CA, USA) with at least 10,000 events recorded for each condition.

**Statistical analysis**

Each experiment was repeated at least three times, and data were expressed as the mean ± standard deviation values. SPSS Version 17.0 software (SPSS Inc., Chicago, IL, USA) was used for statistical analysis. χ² analysis and Student’s t-test were appropriately used for different categories of data. P<0.05 was considered statistically significant.

**Results**

**Knockdown of PGC-1α impaired the viability of endometrial cancer cells**

Our previous study suggested that PGC-1α could actually support cell proliferation by the regulation of energy metabolism in endometrial carcinoma tissues. To further ascertain the role of PGC-1α in the progress of endometrial cancer, the expression of Ishikawa cell, an ER-positive endometrial cancer cell line, was downregulated by using two different PGC-1α siRNAs. The levels of PGC-1α were detected by Western blot to determine the effect of RNA interference. As shown in Figure 1A, two different PGC-1α siRNAs effectively reduced the endogenous expression of PGC-1α. Next, the effect of PGC-1α knockdown on cell viability was measured by CCK-8. The results showed that there was a significant reduction in cell viability of PGC-1α knockdown cells (Figure 1B). Furthermore, the apoptosis in the Ishikawa cell was analyzed by flow cytometry with AV-FITC/PI double staining (Figure 1C and D). There was a significant difference in the apoptotic cell population between control and PGC-1α siRNAs groups. Compared with the control, knockdown of PGC-1α led to an increase in apoptotic rate in the early
Figure 1 Effect of PGC-1α knockdown on cell proliferation and apoptosis in Ishikawa cells.

Notes: (A) Western blot analysis for the expression of PGC-1α in Ishikawa cells treated by no-silence siRNA or two siRNAs targeting PGC-1α. (B) CCK-8 analysis to detect cell proliferation. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α were seeded in 96-well plates, and cell proliferation was detected by CCK-8. Data are presented as mean ± SD values. *P<0.05 versus the no-silence group and **P<0.01 versus the no-silence group. (C and D) Flow cytometric analysis to detect the early cell apoptosis and the later cell apoptosis. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α were collected and stained with AV-FITC and PI before flow cytometric analysis. Data are presented as mean ± SD values. ***P<0.01 versus the no-silence group. (E) TUNEL analysis to detect cell apoptosis. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α were stained with TUNEL assay (red) and DAPI (blue) before imaging by microscopy. The arrows show TUNEL-positive cells. There was an increase in the number of TUNEL-positive cells in the absence of PGC-1α.

Abbreviations: PGC-1α, proliferator-activated receptor gamma coactivator-1α; siRNA, small interfering RNA; CCK-8, Cell Counting Kit-8; AV-FITC, Annexin V-fluorescein isothiocyanate; PI, propidium iodide; DAPI, 4′,6-diamidino-2-phenylindole; OD, optical density; hrs, hours; TUNEL, terminal deoxynucleotidyl transferase dUTP nick end labeling.
apoptotic phase as well as later apoptotic phase, which could be further confirmed by the detection of TUNEL apoptosis (Figure 1E). These results suggested that knockdown of PGC-1α attenuates the survival of endometrial cancer cells by inducing cell apoptosis.

**Lack of PGC-1α induced the apoptosis of endometrial cancer cells through the mitochondrial pathway**

Alteration in mitochondrial membrane permeability is also regarded as one of the crucial events in cellular apoptosis. Therefore, we first measured the potential using a method of staining with JC-1, and then the cells were subsequently analyzed by flow cytometry (Figure 2A). The aggregate JC-1 was dramatically decreased, while the green fluorescent JC-1 monomer was increased (Figure 2B). These data suggested that PGC-1α deficiency induced an obvious alteration in mitochondrial membrane permeabilization.

It is well recognized that mitochondrial membrane permeability is controlled by the balance between anti-apoptotic protein Bcl-2 and proapoptotic protein Bax. Furthermore, the mitochondrial pathway of apoptosis is initiated by mitochondrial membrane permeabilization, leading to the release of apoptogenic factors such as Smac. Hence, the changes in these proteins were determined by Western blot in the process. The results showed that knockdown of PGC-1α significantly induced a reduction in the levels of Bcl-2 and an increase in the levels of Bax and Smac (Figure 2C and D). These data indicated that the depletion of PGC-1α triggered the mitochondrial pathway of apoptosis to attenuate the survival of endometrial cancer cells.

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**Figure 2** Effect of PGC-1α knockdown on mitochondrial membrane potential and the expression of Bcl-2, Bax, and Smac in Ishikawa cells.

**Notes:** (A and B) Flow cytometry was performed using JC-1 to determine the mitochondrial membrane potential and apoptotic cell death rate. Data are presented as mean ± SD values. **P<0.01 versus the no-silence group.***P<0.01 versus the no-silence group. (C) Western blot analysis to detect the Bcl-2 and Bax protein levels. Data are presented as mean ± SD values. **P<0.01 versus the no-silence group.***P<0.01 versus the no-silence group. (D) Western blot analysis to detect the Smac protein level. Data are presented as mean ± SD values. **P<0.01 versus the no-silence group.***P<0.01 versus the no-silence group. **Abbreviations:** PGC-1α, proliferator-activated receptor gamma coactivator-1α; siRNA, small interfering RNA.
Estrogen remedied the PGC-1α efficiency-induced decline of endometrial cancer cells’ viability

Some reports suggested that there might exist an association between estrogen and PGC-1α in the regulation of mitochondrial function. Therefore, the two endometrial cancer cell lines, Ishikawa cells (ER positive) and HEC-1-A cells (ER negative), were selected to further investigate the adaptation of PGC-1α-deficiency cells under estrogen stimulation. First, the results from the CCK-8 analysis showed that estrogen could counteract the effects of PGC-1α deficiency on the cells viability of Ishikawa cells (Figure 3A). However, there were no alterations in HEC-1-A cells’ viability in the presence or absence of estrogen (P>0.05; Figure 3B). Subsequent results from the analysis of flow cytometry with Annexin V-FITC/PI double staining displayed similar results in Ishikawa cells (P<0.05; Figure 3C and D) and HEC-1-A cells (P>0.05; Figure 3E and F). Notably, the downregulation of PGC-1α could augment estrogen-induced apoptosis resistance in endometrial cancer cells. Meanwhile, estrogen primarily inhibited the early cellular apoptosis induced by PGC-1α deficiency (Figure 3C and D). Collectively, these data suggested that there was a compensatory effect between PGC-1α deficiency and estrogen treatment.

Estrogen modulated the mitochondrial function to inhibit the PGC-1α deficiency-induced apoptosis in endometrial cancer cells

The previous results had indicated the involvement of the mitochondrial pathway in PGC-1α deficiency-induced cellular apoptosis in endometrial cancer cells. To further verify the interaction between PGC-1α and estrogen, the alterations in mitochondrial membrane permeability were measured under estrogen stimulation. As shown in Figure 4, estrogen could remarkably abolish PGC-1α deficiency-induced increase in mitochondrial membrane permeability of Ishikawa cells (P<0.05; Figure 4A and B), whereas PGC-1α deficiency-induced increase remained unchanged in the HEC-1-A cells (P>0.05; Figure 4C and D) after treatment with estrogen. Further results from Western blot also showed that the decreased levels of antiapoptotic protein Bcl-2 were elevated in PGC-1α-deficiency Ishikawa cells after treatment with estrogen (Figure 5A). On the contrary, the increased levels of proapoptotic proteins Bax and Smac were obviously lessening (Figure 5B and C). Taken together, these data suggested that estrogen modulated the mitochondrial function to inhibit the PGC-1α deficiency-induced apoptosis in endometrial cancer cells.

Discussion

The PGC-1α-dependent pathway of mitochondrial biogenesis has been reported in human endometrial tissue. Our previous clinical study indicated that the upregulation of PGC-1α and ERRγ in the endometrial cancer tissue was related to tumor energy metabolism, which contributed to the development of endometrial cancer. Here, we have identified a novel synergism mechanism between PGC-1α and estrogen in endometrial cancer cells: PGC-1α was required for the survival of endometrial cancer cells, depletion of PGC-1α could induce the mitochondrial-dependent apoptosis in endometrial cancer cells, and the addition of estrogen protected the mitochondrial function to counteract the effect of PGC-1α deficiency in endometrial cancer cells.

Although a growing body of evidence points toward an important role for PGC-1α in cancer, there exists a contradiction about the roles of PGC-1α in different cancer types. An improved understanding of the role of PGC-1α in different tumor types will be important in determining whether this pathway will be amenable to therapeutic intervention. In the present study, our findings revealed that PGC-1α had protumorigenic function in the development of endometrial cancer, that is, PGC-1α was required for the survival of endometrial cancer. Our results indicated that the endometrial cancer cells’ knockdown of PGC-1α was inclined to apoptosis. Given its role as a “master regulator” of mitochondria, a question was raised whether PGC-1α was involved in the progress of the mitochondrial-mediated apoptotic pathway. Subsequently, this hypothesis was verified by the enhancement in mitochondrial membrane permeabilization, the reduction in antiapoptotic protein Bcl-2, and the increase in the proapoptotic proteins Bax and Smac in PGC-1α-deficiency endometrial cancer cells. It had been reported that PGC-1α could interact with different tissue-specific transcription factors, thus driving distinct genetic programs in different cancer types. The heterogeneous response of PGC-1α in various tumors was likely dictated at least in part by tumor microenvironment. Therefore, we postulated that the roles of PGC-1α in endometrial cancer were likely affected by estrogen stimulation, and then, whether the estrogen response overlaid the effects of PGC-1α to enhance the mitochondrial function and then led to the outcome of apoptotic resistance. Given this, we further investigated the effects of estrogen on the mitochondrial function under PGC-1α-deficiency condition.

Accumulating studies have shown that estrogen could rescue the mitochondrial function. For example, estrogen amelioration of Aβ-induced defects in mitochondria is mediated by mitochondrial signaling pathway. Estrogen
Synergism between PGC-1α and estrogen in endometrial cancer cells

Figure 3 Cell proliferation and apoptosis of endometrial cancer cells in the presence and absence of estrogen.

Notes: (A) CCK-8 analysis to detect cell proliferation. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were seeded in 96-well plates, and cell proliferation was detected by CCK-8. Data are presented as mean ± SD values. *P<0.05. (B) CCK-8 analysis to detect cell proliferation. HEC-1-A cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were seeded in 96-well plates, and cell proliferation was detected by CCK-8. Data are presented as mean ± SD values. *P>0.05. (C and D) Flow cytometric analysis to detect the early cell apoptosis and the later cell apoptosis. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were collected and stained with AV-FITC and PI before flow cytometric analysis. Data are presented as mean ± SD values. *P<0.05, **P<0.01, ***P<0.001. (E and F) Flow cytometric analysis to detect the early cell apoptosis and the later cell apoptosis. HEC-1-A cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were collected and stained with AV-FITC and PI before flow cytometric analysis. Data are presented as mean ± SD values. *P>0.05.

Abbreviations: CCK-8, Cell Counting Kit-8; siRNA, small interfering RNA; PGC-1α, proliferator-activated receptor gamma coactivator-1α; AV-FITC, Annexin V-fluorescein isothiocyanate; PI, propidium iodide; OD, optical density; hrs, hours.
could modulate metabolic adaptations by mitochondrial pathways in breast cancer cells. Some studies previously reported that estrogen acted through ERs to upregulate expression of nuclear-encoded mitochondrial proteins to modulate the mitochondrial function. In Ishikawa cells, ER-positive endometrial cancer cells, the addition of estrogen restored the effects of PGC-1α deficiency on the mitochondrial function, including enhanced mitochondrial membrane permeabilization, decreased antiapoptotic protein Bcl-2, and increased proapoptotic proteins Bax and Smac. However, in the HEC-1-A cells, ER-negative endometrial cancer cells, there were no effects on these alterations in the presence of estrogen. These data demonstrated that the rescue effects of estrogen in the process were ER-dependent pathways. More interestingly, upon only estrogen stimulation, lack of PGC-1α yet induced the apoptosis of all endometrial cancer cells, which was not limited with the existence of ERs. This disconnection suggests that estrogen protected the mitochondria function to inhibit cell apoptosis by a pathway distinct from PGC-1α. Based on this, our findings suggested that PGC-1α did not mediate effects of estrogen on ERs and mitochondrial function in endometrial cancer cells, while PGC-1α coactivates both ERs. A recent study has indicated that estrogen modulates several master transcriptional regulators of mitochondrial function, NRF-1, TFAM, and PGC-1 isoforms, in the blood vessels of brain. The pattern of a
**Figure 5** The Bcl-2, Bax, and Smac protein levels of endometrial cancer cells in the presence and absence of estrogen.

**Notes:** (A) Western blot analysis to detect the Bcl-2 protein level. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were harvested, and cell lysates were prepared and used for Western blot. Data are presented as mean ± SD values. **P<0.05.** (B) Western blot analysis to detect the Bax protein level. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were harvested, and cell lysates were prepared and used for Western blot. Data are presented as mean ± SD values. **P<0.05.** (C) Western blot analysis to detect the Smac protein level. Ishikawa cells transfected with no-silence siRNA or two siRNAs targeting PGC-1α in the presence and absence of estrogen were harvested, and cell lysates were prepared and used for Western blot. Data are presented as mean ± SD values. **P<0.05.

**Abbreviations:** siRNA, small interfering RNA; PGC-1α, proliferator-activated receptor gamma coactivator-1α.
decrease in PGC-1α together with an increase in PGC-1β, PRC, NRF-1, and TFAM appeared to underlie protective effects of estrogen on cerebral blood vessel mitochondria. Conversely, another study indicated that the increased mitochondrial biogenesis in type I endometrial cancer was associated with the upregulation of PGC-1α signaling pathway, which might transactivate NRF-1 leading to TFAM content increase and increase in mitochondrial biogenesis.28 These contradictory results may reflect context-specific actions of estrogen and the particular role of mitochondria in supporting the different metabolic requirements of tissue function. In the present paper, we provided experimental evidence to demonstrate that PGC-1α signaling-mediated mitochondrial apoptosis resistance paralleled with the mitochondrial effects of estrogen in endometrial cancer cells, but there was a synergistic effect on the protection of mitochondrial function to resist mitochondria-dependent apoptosis.

Conclusion
Our study revealed that PGC-1α was required for the survival of endometrial cancer cells and downregulation of PGC-1α is one of the major contributors of apoptosis in endometrial cancer cells. Additionally, there was a synergism between PGC-1α and estrogen, that is, estrogen stimulation from tumor microenvironment could exaggerate the effects of PGC-1α, which was separate from the PGC-1α signaling. Our study supported a crucial and potential role for PGC-1α in endometrial cancer and suggested that it might be a potential target for therapeutic strategy in endometrial cancer.

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Author contributions
All authors made substantial contributions to conception and design, acquisition of data, or analysis and interpretation of data; took part in drafting the article or revising it critically for important intellectual content; gave final approval of the version to be published; and agree to be accountable for all aspects of the work.

Disclosure
The authors report no conflicts of interest in this work.

References


