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# Estrogen induces cell proliferation by promoting ABCG2-mediated efflux in endometrial cancer cells



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#### ABSTRACT

Recently, it has reported that overeating of lipid-food has led to increase the amount of estrogen in vivo and the incidence of endometrial carcinomas. It is well-known that ATP-binding cassette transporter sub-family G2 (ABCG2) is highly expressed in cancer stem cells (CSCs). CSCs possess the ability for differentiation, tumorigenesis, stem cell self-renewal, and the efflux of anti-cancer drug and these abilities affect malignancy of cancer cells. However, little is known about the relationship between the expression of ABCG2 and malignancy of cancer cells. The present study aimed at understanding the regulatory mechanism underlying 17- $\beta$ -estradiol (E<sub>2</sub>)-induced cell proliferation under the control of ABCG2. E<sub>2</sub> increased cell viability with a peak at 1  $\mu$ M and facilitated ABCG2 mRNA expression followed by the increase of ABCG2 expression level at plasma membrane. E<sub>2</sub>-induced cell proliferation was inhibited by reserpine, an inhibitor of ABCG2, and the ABCG2 siRNA treatment. Thus, these results imply that ABCG2 plays an important role in the promotion of E<sub>2</sub>-induced cell proliferation in Ishikawa cells.

#### 1. Introduction

One of the most important transporters for proteins, lipids, and ions efflux is ATP-binding cassette (ABC) transporter superfamily and these transporters are widely expressed in organelles such as plasma membrane, vesicle, and mitochondria. ABC transporters function as active transporters for multiple substrates across the cellular membrane by ATP hydrolysis [1]. Recently, it has reported that cancer stem cells (CSCs) are thought to play an important role in tumor recurrence and drug resistance, which result in therapy failure and tumor relapse [2]. Furthermore, CSCs possess the ability for self-renewal, differentiation, and tumorigenesis [3]. It is well-known that P-glycoprotein (ABCB1) [4], multidrug resistance-associated protein 4 (MRP4, ABCC4) [5] and ATP-binding cassette transporter sub-family G2 (ABCG2) are expressed in CSCs. Especially, ABCG2 is highly expressed in CSCs. CSCs acquire anti-cancer drug resistance ability and ABCG2 is used as marker of collecting CSCs from cancer cells [6,7].

Endometrial adenocarcinomas can be divided into two types: Type I, Ishikawa cells, has a hormone-derived etiology such as estrogen, while Type II is more aggressive and estrogen-independent [8]. Type I occurs more frequently in young people and develops via endometrial hyperplasia due to overexposure of estrogen. Phosphatase and tensin homolog (PTEN) and v-Ki-ras2 Kirsten rat sarcoma viral oncogene homolog (KRAS) mutations in Type I [9–12], and Tp53 mutation in Type II [13,14] play an important role in the carcinogenic process of endometrial cancer, however, the relationship between genetic mutations and the mechanism of the onset of endometrial carcinoma is not well understood.

In Ishikawa cells, it is not fully understood the mechanisms underlying E<sub>2</sub>-induced cell proliferation. The present study focused upon ABCG2 functions and investigated the role of ABCG2 on E<sub>2</sub>-induced cell proliferation. The results showed that 17- $\beta$ -estradiol (E<sub>2</sub>) (1  $\mu$ M) increases the number of Ki-67 positive cells, and upregulated the expression level of ABCG2 at plasma membrane. The E<sub>2</sub> effect was inhibited by reserpine and the ABCG2 siRNA, thus, ABCG2 activated by E<sub>2</sub> promoted unnecessary substances efflux and led to accelerate malignancy of Ishikawa cells.

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#### 2. Materials and methods

#### 2.1. Cell culture

Ishikawa cells were obtained from Japan Health Sciences Foundation (Tokyo, Japan). Cells in DMEM supplemented with 10% heat-inactivated fetal bovine serum without phenol red were cultured at 37 °C in 5%  $\rm CO_2$  incubator.

#### 2.2. Cell viability

Cell viability was measured using 3-(4, 5-dimethyl-2-thiazolyl)-2, 5diphenyl-2H-tetrazolium bromide (MTT) assay (Dojindo, kumamoto, Japan). MTT-reactive cells were quantified at an absorbance of 570 nm using a micro-plate reader Model 680 (Bio Rad, CA, USA), and percentage of independent basal levels was calculated.

#### 2.3. Western blotting

Lysates from Ishikawa cells were loaded onto 10% (w/v) or 12% (w/v) sodium dodecylsulfate polyacrylamide gel electrophoresis (SDS-PAGE) and transferred to polyvinylidene difluoride (PVDF). Blotting membranes were blocked by TBST [150 mM NaCl, 0.1% (v/v) Tween20, and 20 mM Tris] containing 5% (v/v) bovine serum albumin (BSA) and reacted with an anti-ABCG2 antibody (1:250, Santa Cruz Biotechnology, YX, USA) or an anti- $\beta$ -actin antibody (1:1000, Sigma-Aldrich, Tokyo, Japan) followed by Histofine Simple Stain MAX-PO (Nichirei Bioscience, Tokyo, Japan). Immunoreactivity was detected with a diaminobenzidine (DAB).

#### 2.4. Immunohistochemistry

Cells were fixed with 4% paraformaldehyde and were reacted with Ki-67 (1:20, Dako, CA, USA), ABCG2 (1:250), or SYTO 59 Red Fluorescent Nucleic Acid Stain (Thermo Fisher Science, MA, USA). Following an anti-Ki-67 antibody, cells were reacted with Histofine Simple Stain MAX-PO followed by the detection with a DAB. Cells stained by DAB were observed with a light microscope (Nikon, Tokyo, Japan) at 20 × objection lens. Cells were reacted with an anti-ABCG2 antibody followed by the reaction of Alexa Fluor 488 (Thermo Fisher Science, MA, USA). Fluorescence signals were obtained with Olympus Fluoview FV300 confocal microscope (Olympus, Tokyo, Japan) at 20 × objection lens.

#### 2.5. RNA interference

Small interfering RNAs (siRNAs) silencing the ABCG2 gene: sense, 5'-UGAGGAUGUUACCAAGUAU-3' and anti-sense, 5'-AUACUUGGUAA CAUCCUCA-3'; and sense, 5'-ACAGGGUCAUUCAAGAGUU-3' and antisense, 5'-AACUCUUGAAUGACCCUGU-3'; and sense, 5'-GUAUAGGAA UGGAGCUUAU-3' and snit-sense, 5'-AUAAGCUCCAUUCCUAUAC-3', and negative control (NC) siRNA were purchased from Nippon gene (Toyama, Japan). Each siRNA was transfected by Lipofectamine 3000 (Invitrogen, CA, USA).

#### 2.6. RT-PCR

Total RNAs were purified from Ishikawa cells using TRIZol reagent (Thermo Fisher Scientific, Kanagawa, Japan). Each cDNA (1  $\mu$ L) was amplified with rTaq polymerase (TOYOBO, Osaka, Japan) as follows: the first step, 94 °C for 2 min and the ensuring 30–35 cycles, 94 °C for 1 s, 62 °C for 15 s, and 72 °C for 30 s. We carried out using primers as follows: 5'- TTATGATGGTGGCTTATTCAGCCAGTTCC-3' and 5'- TTCG CCAGTACATGTTGCATAGTTACAAGG-3' for ABCG2; 5'- GAGGTGAAG AAGGGCCAGACGCTGGC-3' and 5'- GGCTTTGTCCAGGGCTTCTTGGA CAACC-3' for ABCB1; 5'- TGACAATGTGAACTTCATGTACAGTCC









**Fig. 1.**  $E_2$  increases cell viability in Ishikawa cells. A: Ishikawa cells were treated with  $E_2$  at concentration as indicated for 24 and 48 h, and cell viability was measured by MTT assay. Data showed the mean ( $\pm$  SEM) percentage of basal level (n = 5–8 independent experiments). \*\*, §§ P < 0.01 as compared with control, unpaired *t*-test. B: Immunohistochemistry was carried out in the presence or absence of  $E_2$ . After 24-h treatment, cells were reacted with an anti-Ki67 antibody followed by Histofine Simple Stain MAX-PO and detected with DAB. In the graph, indicative bar showed 50 µm. Results are from n = 3 independent experiments. C: The number of Ki-67 positive cells were counted in the consistent area (680 µm x 900 µm). In the graph, each value represents the mean ( $\pm$  SEM) number of Ki-67 positive cells (n = 3 independent experiments). \*\*P < 0.01, unpaired *t*-test.

AGG-3' and 5'- CCAGTTGTCTTTGTCCAACACTAAAATTGG-3' for ABCC4; 5'-GACTTCAACAGCGACACCCACTCC-3' and 5'-AGGTCCACCA CCCTGTTGCTGTAG-3' for GAPDH. PCR products were separated by an electrophoresis on 2% (v/v) agarose gel and stained with ethidium bromide followed by visualization using UV Transilluminator (Funakoshi, Tokyo, Japan).

#### 2.7. Statistical analysis

Statistical analysis was performed by using unpaired *t*-test with the Stat View software. A value of P < 0.05 was considered significant.

#### 3. Results

#### 3.1. $E_2$ increases cell viability in Ishikawa cells

We examined the effect of  $E_2$  on cell viability in Ishikawa cells. In survival analysis, cell viability induced by  $E_2$  was increased in a bell shape at 24, 48 h, each reaching 129  $\pm$  10% and 144  $\pm$  7% with a peak at 1  $\mu$ M, respectively (Fig. 1A). Subsequently, we examined whether  $E_2$  induced cell proliferation in Ishikawa cells.  $E_2$  significantly increased the number of Ki-67 positive cells as compared with the control after 24-h treatment (Fig. 1B, C). These results suggest that  $E_2$  induces cell proliferation in Ishikawa cells.

## 3.2. $E_2$ upregulated the expression level of ABCG2 followed by leading to facilitate cell proliferation

It is well-known that E2 binds to estrogen receptors (ERs) followed by leading to transcribe E2/ER-targeted genes. To investigate whether E<sub>2</sub> facilitated *de novo* protein synthesis, we used cycloheximide (CHX), an inhibitor of protein synthesis. E2-induced cell proliferation was significantly inhibited by CHX (10 ng/mL) (Fig. 2A). This result suggests that E2 facilitated cell proliferation via de novo protein synthesis. We next asked a question as to what kind of gene expression underlie the E<sub>2</sub> action. We focused upon investigating the expression of ABC transporters such as ABCB1, ABCC4 and ABCG2. It is well-known that these transporters were widely expressed in various cancer cells and facilitated the efflux of exogenous toxins or anti-cancer drugs. So, we examined the expression level of these transporters using RT-PCR. E<sub>2</sub> did not affect the expression of ABCB1 mRNA, however, increased the expression of ABCC4 and ABCG2 mRNA transiently at 30-min treatment (Fig. 2B). In the present study, we focused upon ABCG2 expression, because ABCG2 was widely expressed in CSCs and was thought to play an important role in malignancy of cancer cells. E2 increased ABCG2 protein expression level (Fig. 3A) and the expression level of ABCG2 at plasma membrane after 24-h treatment (Fig. 3B). These results imply that E2 upregulates the expression level of ABCG2 at plasma membrane and activates ABCG2 function followed by promoting cell proliferation.

## 3.3. ABCG2 inhibitor and the ABCG2 siRNA attenuate $E_2$ -induced cell proliferation

We finally examined whether ABCG2 exhibits a promotive effect against cell proliferation. Reserpine, an inhibitor of ABCG2, significantly attenuated  $E_2$ -induced cell proliferation as compared with  $E_2$ treated cell group at 24 h (Fig. 4B). Furthermore, we constructed the siRNA silencing the ABCG2 gene. The expression of ABCG2 mRNA was clearly suppressed for cells transfected with the ABCG2 siRNA as compared with the expression for cells transfected with the negative control (NC) siRNA (Fig. 4A).  $E_2$ -induced cell proliferation was significantly blocked by the ABCG2 siRNA treatment (Fig. 4A). These results suggest that ABCG2 plays an important role in  $E_2$ -induced cell proliferation. Overall, these results indicate that  $E_2$  increases the expression level of ABCG2 at plasma membrane followed by activating ABCG2 function and increases cell proliferation via the facilitation of ABCG2-mdeiated efflux.



**Fig. 2.**  $E_2$  increases the expression level of ABC transporters in Ishikawa cells. A: Cells were treated with 1 µM of  $E_2$  in the presence or absence of CHX (10 ng/mL) for 24 h, and cell viability was measured by MTT assay. Data showed the mean ( $\pm$  SEM) percentage of basal level (n = 6–8 independent experiments). \*\**P* < 0.01 as compared with control, unpaired *t*-test. B: Cells were treated with  $E_2$  (1 µM) for period time as indicated. Total mRNAs were extracted from Ishikawa cells and ABC transporter mRNAs (ABCB1, ABCC4, and ABCG2) were detected using RT-PCR. Note that a similar result was obtained with 3–5 independent experiments.

#### 4. Discussion

The results of the present study demonstrated that  $1 \mu M E_2$  facilitated cell proliferation via the new protein synthesis. Generally, it was reported that  $E_2$  bound to ERs followed by entering the nucleus and  $E_2$ /ERs complex transcribed various target genes, such as PTEN [15], IL-20 [16], and p21-activated kinase 4 (PAK4) [17]. Especially, PAK4 is a family of serine/threonine kinases that is first identified as effectors for Rho-family GTPases and are divided into group I (PAK1–3) and group II (PAK4–6) in humans [18,19]. PAK4 regulates cytoskeleton reorganization, directional motility, embryonic development, survival, invasion, metastasis, and gene regulation [20]. PAK4-signaling dependent cellular functions regulate disease processes such as cancer as PAK4 is overexpressed and/or hyper-stimulated in human cancer [21]. Thus,  $E_2$ /ER-signaling pathways might play crucial roles in malignancy of cancer cells.

ABC transporters are widely expressed in CSCs and promote the efflux of proteins, lipids, ions, and anti-cancer drug. The superfamily of ABC transporters consists of 48 members and ABCB1, ABCC4 and ABCG2 hold a major key in malignancy of cancers by mediating the efflux of many anti-cancer drug [22]. ABCG2 are also widely detected in CSCs and used as a surface marker of CSCs. We hypothesized that ABC transporters expressed in CSCs played an important role in keeping intracellular environment clear by facilitating the efflux of the unnecessary substances and might lead to induce cell proliferation. In Ishikawa cells,  $E_2$  upregulated ABCC4 and ABCG2 mRNA, but not ABCB1 mRNA. These results imply that not only ABCG2 but also ABCC4 plays an important role in  $E_2$ -induced cell proliferation. The present

Α

ABCG2

ABCG2 siRNA



**Fig. 3.**  $E_2$  increases the expression level of ABCG2 in Ishikawa cells. A: After 24-h treatment with  $E_2$ , proteins were collected from Ishikawa cells and Western blotting was carried out. The blotting membranes were reacted with an anti-ABCG2 antibody or an anti- $\beta$ -actin antibody followed by Histofine Simple Stain MAX-PO and detected with DAB. B: After 24-h treatment with  $E_2$ , immunohistochemistry was carried out. Ishikawa cells were fixed with 4% paraformaldehyde and were reacted with an anti-ABCG2 antibody followed by Alexa Fluor 488 (green). Subsequently, cells were stained with SYTO 59 Red Fluorescent Nucleic Acid Stain (red). Results are from 3 to 5 independent experiments (A, B).



We used reserpine, an inhibitor of ABCG2, and ABCG2 siRNA due to investigate the role of ABCG2 on E<sub>2</sub>-induced cell proliferation in Ishikawa cells. Reserpine and ABCG2 siRNA significantly inhibited E<sub>2</sub>induced cell proliferation. ABCG2 acts as a part of a self-defense mechanism for the organism [25]. E<sub>2</sub> might regulate the intracellular environment by promoting the unnecessary substances efflux from intracellular space via ABCG2, thus, accelerate malignancy of Ishikawa cells.

In conclusion,  $E_2$  increased ABCG2 mRNA expression and facilitated the expression level of ABCG2 at plasma membrane. Furthermore, ABCG2 promoted the unnecessary substances efflux followed by increasing cell viability of Ishikawa cells. ABCG2 plays an important role in malignancy of Ishikawa cells and is considered to be useful as a target for cancer treatment.

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#### Author contribution

1) Conceptualization: T.Y.; 2) Data curation: T.Y.; 3) Formal analysis: T.O. and T.Y.; 4) Funding acquisition: I have no fund.; 5)

Mock

siRNA

Fig. 4. ABCG2 inhibitor and the ABCG2 siRNA attenuate  $E_2$ -induced cell proliferation. A: Cells were transfected with the NC siRNA (Mock) or the ABCG2 siRNA. After 24-h treatment, total RNAs were purified and RT-PCR was carried out (n = 3 independent experiments). Note that no signal band for ABCG2 mRNA is visible in cells transfected with the ABCG2 siRNA. B: Cells were treated and untreated with reserpine (25  $\mu$ M) or were transfected with the NC siRNA or the ABCG2 siRNA. These cells were treated and untreated with  $E_2$  (1  $\mu$ M) for 24 h followed by MTT assay. Data showed the mean ( $\pm$  SEM) percentage of basal level (n = 6–8 independent experiments). \*\*P < 0.01 as compared with control, unpaired *t*-test.

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Investigation: T.O. and T.Y.; 6) Methodology: T.O. and T.Y.; 7) Project administration: T.Y.; 8) Resources: T.Y.; 9) Software: T.Y.; 10) Supervision:T.Y.; 11) Validation: T.Y.; 12) Visualization: T.Y.; 13) Writing - original draft: T.Y.; 14) Writing - review & editing: T.Y.

#### Appendix A. Transparency document

Supplementary data associated with this article can be found in the online version at doi:10.1016/j.bbrep.2018.10.005.

#### References

- C.B. Jeong, H.S. Kim, H.M. Kang, J.S. Lee, ATP-binding cassette (ABC) proteins in aquatic invertebrates: evolutionary significance and application in marine ecotoxicology, Aquat. Toxicol. 185 (2017) 29–39.
- [2] B.C. Lu, J. Li, W.F. Yu, G.Z. Zhang, H.M. Wang, H.M. Ma, Elevated expression of Nrf2 mediates multidrug resistance in CD133 + head and neck squamous cell carcinoma stem cells, Oncol. Lett. 12 (6) (2016) 4333–4338.
- [3] N. Bansal, V. Yendluri, R.M. Wenham, The molecular biology of endometrial cancers and the implications for pathogenesis, classification, and targeted therapies, Cancer Control. 16 (1) (2009) 8–13.
- [4] X.Z. Tong, F. Wang, S. Liang, X. Zhang, J.H. He, X.G. Chen, Y.J. Liang, Y.J. Mi, K.K. To, L.W. Fu, Apatinib (YN968D1) enhances the efficacy of conventional chemotherapeutical drugs in side population cells and ABCB1-overexpressing leukemia cells, Biochem. Pharmacol. 83 (5) (2012) 586–597.
- [5] L. Oevermann, J. Scheitz, K. Starke, K. Köck, T. Kiefer, G. Dölken, J. Niessen, A. Greinacher, W. Siegmund, M. Zygmunt, H.K. Kroemer, G. Jedlitschky, C.A. Ritter, Hematopoietic stem cell differentiation affects expression and function of MRP4 (ABCC4), a transport protein for signaling molecules and drugs, Int. J. Cancer 124 (10) (2009) 2303–2311.
- [6] A.F. Prasanphanich, D.E. White, M.A. Gran, M.L. Kemp, Kinetic modeling of ABCG2 transporter heterogeneity: a quantitative, single-cell analysis of the side population assay, PLoS Comput. Biol. 12 (11) (2016), https://doi.org/10.1371/journal.pcbi. 1005188.
- [7] G.F. Guan, D.J. Zhang, Y. Zheng, L.J. Wen, D.J. Yu, Y.Q. Lu, Y. Zhao, Abnormal Wnt signaling and overexpression of ABCG2 contributes to drug efflux properties of side population cells in nasopharyngeal carcinoma, Mol. Med. Rep. 12 (3) (2015) 4352–4357.
- [8] A.A. Suarez, A.S. Felix, D.E. Cohn, Bokhman redux: endometrial cancer "types" in the 21st century, Gynecol. Oncol. 144 (2) (2017) 243–249.
- [9] L.J.M. van der Putten, R. van Hoof, B.B.J. Tops, M.P.L.M. Snijders, S.H. van den Berg-van Erp, A.A.M. van der Wurff, J. Bulten, J.M.A. Pijnenborg, L.F.A.G. Massuger, Molecular profiles of benign and (pre)malignant endometrial lesions, Carcinogenesis 38 (3) (2017) 329–335.
- [10] R.D. Burk, Z. Chen, C. Saller, K. Tarvin, A.L. Carvalho, C. Scapulatempo-Neto, H.C. Silveira, J.H. Fregnani, C.J. Creighton, M.L. Anderson, P. Castro, S.S. Wang, C. Yau, C. Benz, A.G. Robertson, K. Mungall, L. Lim, R. Bowlby, S. Sadeghi D. Brooks, P. Sipahimalani, R. Mar, A. Ally, A. Clarke, A.J. Mungall, A. Tam, D. Lee, E. Chuah, J.E. Schein, K. Tse, K. Kasaian, Y. Ma, M.A. Marra, M. Mayo, M. Balasundaram, N. Thiessen, N. Dhalla, R. Carlsen, R.A. Moore, R.A. Holt, S.J. Jones, T. Wong, A. Pantazi, M. Parfenov, R. Kucherlapati, A. Hadjipanayis, J. Seidman, M. Kucherlapati, X. Ren, A.W. Xu, L. Yang, P.J. Park, S. Lee, B. Rabeno, L. Huelsenbeck-Dill, M. Borowsky, M. Cadungog, M. Iacocca, N. Petrelli, P. Swanson, A.I. Ojesina, X. Le, G. Sandusky, S.N. Adebamowo, T. Akeredolu, C. Adebamowo, S.M. Reynolds, I. Shmulevich, C. Shelton, D. Crain, D. Mallery, E. Curley, J. Gardner, R. Penny, S. Morris, T. Shelton, J. Liu, L. Lolla, S. Chudamani, Y. Wu, M. Birrer, M.D. McLellan, M.H. Bailey, C.A. Miller, M.A. Wyczałkowski, R.S. Fulton, C.C. Fronick, C. Lu, E.R. Mardis, E.L. Appelbaum, H.K. Schmidt, L.A. Fulton, M.G. Cordes, T. Li, L. Ding, R.K. Wilson, J.S. Rader, B. Behmaram D. Uyar, W. Bradley, J. Wrangle, A. Pastore, D.A. Levine, F. Dao, J. Gao, N. Schultz, C. Sander, M. Ladanyi, M. Einstein, R. Teeter, S. Benz, N. Wentzensen, I. Felau, J.C. Zenklusen, C. Bodelon, J.A. Demchok, L. Yang, M. Sheth, M.L. Ferguson, R. Tarnuzzer, H. Yang, M. Schiffman, J. Zhang, Z. Wang, T. Davidsen, O. Olaniyan,
  - C.M. Hutter, H.J. Sofia, D.A. Gordenin, K. Chan, S.A. Roberts, L.J. Klimczak, C. Van Waes, Z. Chen, A.D. Saleh, H. Cheng, J. Parfitt, J. Bartlett, M. Albert, A. Arnaout, H. Sekhon, S. Gilbert, M. Peto, J. Myers, J. Harr, J. Eckman, J. Bergsten, K. Tucker, L.A. Zach, B.Y. Karlan, J. Lester, S. Orsulic, Q. Sun, R. Naresh, T. Pihl, Y. Wan, H. Zaren, J. Sapp, J. Miller, P. Drwiega, A.I. Ojesina, B.A. Murray, H. Zhang, A.D. Cherniack, C. Sougnez, C.S. Pedamallu, L. Lichtenstein, M. Meyerson,

M.S. Noble, D.I. Heiman, D. Voet, G. Getz, G. Saksena, J. Kim, J. Shih, J. Cho, M.S. Lawrence, N. Gehlenborg, P. Lin, R. Beroukhim, S. Frazer, S.B. Gabriel, S.E. Schumacher, K.M. Leraas, T.M. Lichtenberg, E. Zmuda, J. Bowen, J. Frick, J.M. Gastier-Foster, L. Wise, M. Gerken, N.C. Ramirez, L. Danilova, L. Cope, S.B. Baylin, H.B. Salvesen, C.P. Vellano, Z. Ju, L. Diao, H. Zhao, Z. Chong, M.C. Ryan, E. Martinez-Ledesma, R.G. Verhaak, L. Averett Byers, Y. Yuan, K. Chen, S. Ling, G.B. Mills, Y. Lu, R. Akbani, S. Seth, H. Liang, J. Wang, L. Han, J.N. Weinstein, C.A. Bristow, W. Zhang, H.S. Mahadeshwar, H. Sun, J. Tang, J. Zhang, X. Song, A. Protopopov, K.R. Shaw, L. Chin, O. Olabode, A.I. Ojesina, P. DiSaia, A. Radenbaugh, D. Haussler, J. Zhu, J. Stuart, P. Chalise, D. Koestler, B.L. Fridley, A.K. Godwin, R. Madan, G. Ciriello, C. Martinez, K. Higgins, T. Bocklage, J.T. Auman, C.M. Perou, D. Tan, J.S. Parker, K.A. Hoadley, M.D. Wilkerson, P.A. Mieczkowski, T. Skelly, U. Veluvolu, D.N. Hayes, W.K. Rathmell, A.P. Hoyle, J.V. Simons, J. Wu, L.E. Mose, M.G. Soloway, S. Balu, S. Meng, S.R. Jefferys, T. Bodenheimer, Y. Shi, J. Roach, L.B. Thorne, L. Boice, M. Huang, C.D. Jones, R. Zuna, J. Walker, C. Gunderson, C. Snowbarger, D. Brown, K. Moxley, K. Moore, K. Andrade, L. Landrum, R. Mannel, S. McMeekin, S. Johnson, T. Nelson, E. Elishaev, R. Dhir, R. Edwards, R. Bhargava, D.G. Tiezzi, J.M. Andrade, H. Noushmehr, C. Gilberto Carlotti, D.P. Tirapelli, D.J. Weisenberger, D.J. Van Den Berg, D.T. Maglinte, M.S. Bootwalla, P.H. Lai, T. Triche, E.M. Swisher, K.J. Agnew, C.S. Shelley, P.W. Laird, J. Schwarz, P. Grigsby, D. Mutch, J. Schwarz, P. Grigsby, D. Mutch, Integrated genomic and molecular characterization of cervical cancer, Nature 543 (7645) (2017) 378-384.

- [11] A.D. Cherniack, H. Shen, V. Walter, C. Stewart, B.A. Murray, R. Bowlby, X. Hu, S. Ling, R.A. Soslow, R.R. Broaddus, R.E. Zuna, G. Robertson, P.W. Laird, R. Kucherlapati, G.B. Mills, Cancer Genome Atlas Research Network, J.N. Weinstein, J. Zhang, R. Akbani, D.A. Levine, Integrated molecular characterization of uterine carcinosarcoma, Cancer Cell 31 (3) (2017) 411–423.
- [12] N. Bansal, V. Yendluri, R.M. Wenham, The molecular biology of endometrial cancers and the implications for pathogenesis, classification, and targeted therapies, Cancer Control 16 (1) (2009) 8–13.
- [13] Y. Suga, T. Sugai, N. Uesugi, T. Kawasaki, T. Fukagawa, E. Yamamoto, K. Ishida, H. Suzuki, T. Sugiyama, Molecular analysis of isolated tumor glands from endo metrialendometrioid adenocarcinomas, Pathol. Int. 65 (5) (2015) 240–249.
- [14] E. Stelloo, T. Bosse, R.A. Nout, H.J. MacKay, D.N. Church, H.W. Nijman, A. Leary, R.J. Edmondson, M.E. Powell, E.J. Crosbie, H.C. Kitchener, L. Mileshkin, P.M. Pollock, V.T. Smit, C.L. Creutzberg, Refining prognosis and identifying targetable pathways for high-risk endometrial cancer; a TransPORTEC initiative, Mod. Pathol. 28 (6) (2015) 836–844.
- [15] P.A. Perez, J.P. Petiti, F. Picech, C.B. Guido, L. Del Valle Sosa, E. Grondona, J.H. Mukdsi, A.L. De Paul, A.I. Torres, S. Gutierrez, Estrogen receptor β regulates the tumoral suppressor PTEN to modulate pituitary cell growth, J. Cell Physiol. (2017), https://doi.org/10.1002/jcp.26025.
- [16] J.Y. Lee, Y.J. Park, N. Oh, K.B. Kwack, K.S. Park, A transcriptional complex composed of ER (α), GATA3, FOXA1 and ELL3 regulates IL-20 expression in breast cancer cells, 2017. <a href="https://dx.doi.org/10.18632/oncotarget.17459">https://dx.doi.org/10.18632/oncotarget.17459</a>>.
- [17] Tao Su, Jun-Jie Qu, Kai Wang, Bi-Lan Li, Dong Zhao, Yi-Ping Zhu, Lei Ye, Wen Lu, Xiao-Ping Wan, Cross-talk between p21-activated kinase 4 and ERα signaling triggers endometrial cancer cell proliferation, Oncotarget (2017), https://doi.org/ 10.18632/oncotarget.19188 (PMCID: PMC5620238).
- [18] J. Eswaran, M. Soundararajan, R. Kumar, S. Knapp, UnPAKing the class differences among p21-activated kinases, Trends Biochem. Sci. 33 (2008) 394–403.
- [19] B.H. Ha, E.M. Morse, B.E. Turk, T.J. Boggon, Signaling, regulation, and specificity of the type II p21-activated kinases, J. Biol. Chem. 290 (2015) 12975–12983.
- [20] Y.G. Shao, K. Ning, F. Li, Group II p21-activated kinases as therapeutic targets in gastrointestinal cancer, World J. Gastroenterol. 22 (3) (2016) 1224–1235.
- [21] Y. Li, Y. Shao, Y. Tong, T. Shen, J. Zhang, Y. Li, H. Gu, F. Li, Nucleo-cytoplasmic shuttling of PAK4 modulates β-catenin intracellular translocation and signaling, Biochim. Biophys. Acta 1823 (2) (2012) 465–475.
- [22] H. Cui, A.J. Zhang, M. Chen, J.J. Liu, ABC transporter inhibitors in reversing multidrug resistance to chemotherapy, Curr. Drug Targets 16 (12) (2015) 1356–1371.
- [23] L. Zhang, X. Guo, D. Zhang, Y. Fan, L. Qin, S. Dong, L. Zhang, Upregulated miR-132 in Lgr5+ gastric cancer stem cell-like cells contributes to cisplatin-resistance via SIRT1/CREB/ABCG2 signaling pathway, Mol. Carcinog. (2017), https://doi.org/10. 1002/mc.22656.
- [24] A. Yu, W. Dang, Regulation of stem cell aging by SIRT1 Linking metabolic signaling to epigenetic modifications, Mol. Cell Endocrinol. 455 (2017) 75–82, https://doi.org/10.1016/j.mce.2017.03.031.
- [25] M.H. Hasanabady, F. Kalalinia, ABCG2 inhibition as a therapeutic approach for overcoming multidrug resistance in cancer, J. Biosci. 41 (2) (2016) 313–324.