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Targeting estrogen receptor beta (ER β) for treatment of ovarian cancer: importance of KDM6B and SIRT1 for ER β expression and functionality

Giulia Pinton¹, Stefan Nilsson² and Laura Moro¹

Abstract

Estrogen receptor (ER) β has growth inhibitory and chemo drug potentiating effect on ovarian cancer cells. We studied the dependence of ER β function on the presence of KDM6B and SIRT1 in human ovarian cancer cells in vitro. Activation of ER β with the subtype-selective agonist KB9520 resulted in significant inhibition of human ovarian cancer cell growth. KB9520-activated ER β had an additive effect on growth inhibition in combination with cisplatin and paclitaxel, respectively. Loss of KDM6B expression had a negative effect on ER β function as a ligand-dependent inhibitor of ovarian cancer cell growth. In contrast, loss or inhibition of SIRT1 deacetylase activity restored ligand-activated ER β functionality. Presented data suggest that selective targeting of ER β with an agonist potentiate chemotherapy efficacy for the treatment of ovarian cancer and that downregulation or inhibition of SIRT1 may further enhance its therapeutic effect.

Introduction

Ovarian epithelial carcinoma (OEC), believed to originate from the ovarian surface epithelium, is the fourth commonest cause of female cancer death in the developed world. The OEC incidence is estimated to more than 240,000 new cases and around 150,000 deaths per year worldwide. The highest rates are reported in Scandinavia, Eastern Europe, USA, and Canada^{1–4}.

The etiology of OEC is poorly understood but it is believed that nulliparity, high plasma levels of estrogen or long-term estrogen replacement therapy increases the risk for ovarian cancer, whereas pregnancy, lactation, and oral contraceptives decrease the risk^{1, 4–7}.

The standard of care for the management of advanced ovarian cancer has been unchanged for many years and

includes maximum cytoreductive surgery followed by platinum-based chemotherapy (carboplatin or cisplatin) in combination with paclitaxel^{2, 8}. Although the response rate for first-line carboplatin and paclitaxel is high, approximately 80% of patients with advanced OEC will experience recurrence and eventually become resistant to chemotherapy^{2,8, 9}. New investigational drugs targeting different pathways for improved disease control and for the treatment of platinum-resistant OEC are emerging¹⁰.

Both estrogen receptor (ER) subtypes, ER α and ER β , are expressed in normal ovarian tissue as well as in ovarian cancer cells^{7, 11}. There is, however, a falling trend in ER β expression as cells transition from normal to malignant state and with a further decline in its expression as the malignancy develops from early (stage I) to late disease stages (stage II–IV). No such trend in the expression of ER α has been observed^{7, 11}.

The levels of ER α are closely associated with estrogen-dependent growth and to increased metastatic potential of OEC by promoting epithelial–mesenchymal transition

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(EMT) through upregulation of Snail and Slug and downregulation of E-cadherin. In contrast, ER β mediates opposite effects to ER α in the presence of 17 β -Estradiol (E2), resulting in inhibition of EMT¹².

Although 40–60% of ovarian cancers express ER α , only a portion (15–18%) of them benefit from anti-estrogen (SERM) treatment, if at all. Tamoxifen produces only a modest effect in ovarian cancer^{1,11, 13}.

Introduction of ER β into the ER α -positive human epithelial ovarian cancer cell line BG-1 led to decreased basal and E2-induced cell proliferation in vitro. Moreover, ER β expression resulted in downregulation and inhibition of ER α activity, arrested the cells in the G2/M phase, and decreased the pAKT levels¹³.

Treatment of the ovarian cancer cell lines SKOV3 and OV2008 with the ER α -selective agonist, PPT, led to a significant stimulation of cell growth. In contrast, the ER β -selective agonist, DPN, significantly suppressed the growth of the two ovarian cancer cells. Moreover, the size of tumors in the in vivo SKOV3 xenograft model were significantly smaller in DPN-treated compared to vehicle-treated animals¹⁴. A recent study provided evidence that also natural ER β agonists have the potential to reduce cell viability and survival and to promote apoptosis of OEC cells¹⁵.

ER β expression in ovarian cancer cells has been found to be significantly associated with longer disease-free survival and overall survival (OS)¹¹. Moreover, patients with more than 30% of ER β -positive cancer cells were shown to respond well to chemotherapy and with increased progression-free survival and OS compared to patients with less than 30% of ER β -positive tumor cells¹. The analysis also demonstrated significantly longer OS time of patients with higher ER β immunoreactivity score after chemotherapy, compared with patients with low ER β score.

The recent discovery of alterations in genes encoding histone modifiers suggests their possible roles in cancer development. The lysine-specific demethylase 6B (KDM6B), also called Jumonji domain-containing protein D3 (Jmjd3), is a member of the family of JmjC histone demethylases that specifically catalyzes the demethylation of di-methylated or tri-methylated Lys 27 in histone H3 (H3K27me2/3). The role of KDM6B has been extensively studied in development, cell plasticity, immune system, neurodegenerative disease, and cancer¹⁶.

While, H3K27me3 is considered a repressive epigenetic mark and is recognized as a determining factor in promoting tumorigenesis and tumor progression, the mechanisms underlying KDM6B expression and function in cancer is still controversial¹⁷.

Accumulating evidence indicates that SIRT1, involved in a variety of cellular processes, is a key player in oncogenesis and cancer progression. SIRT1 belongs to a family

of seven class III histone deacetylases (HDACs) enzymes, which are highly conserved enzyme homologs of the yeast Sir2 protein, with nicotine adenine dinucleotide NAD⁺-dependent protein deacetylase activity¹⁸. Over-expression of SIRT1 has been recently associated with poor outcome and chemoresistance of OEC patients^{19–22}.

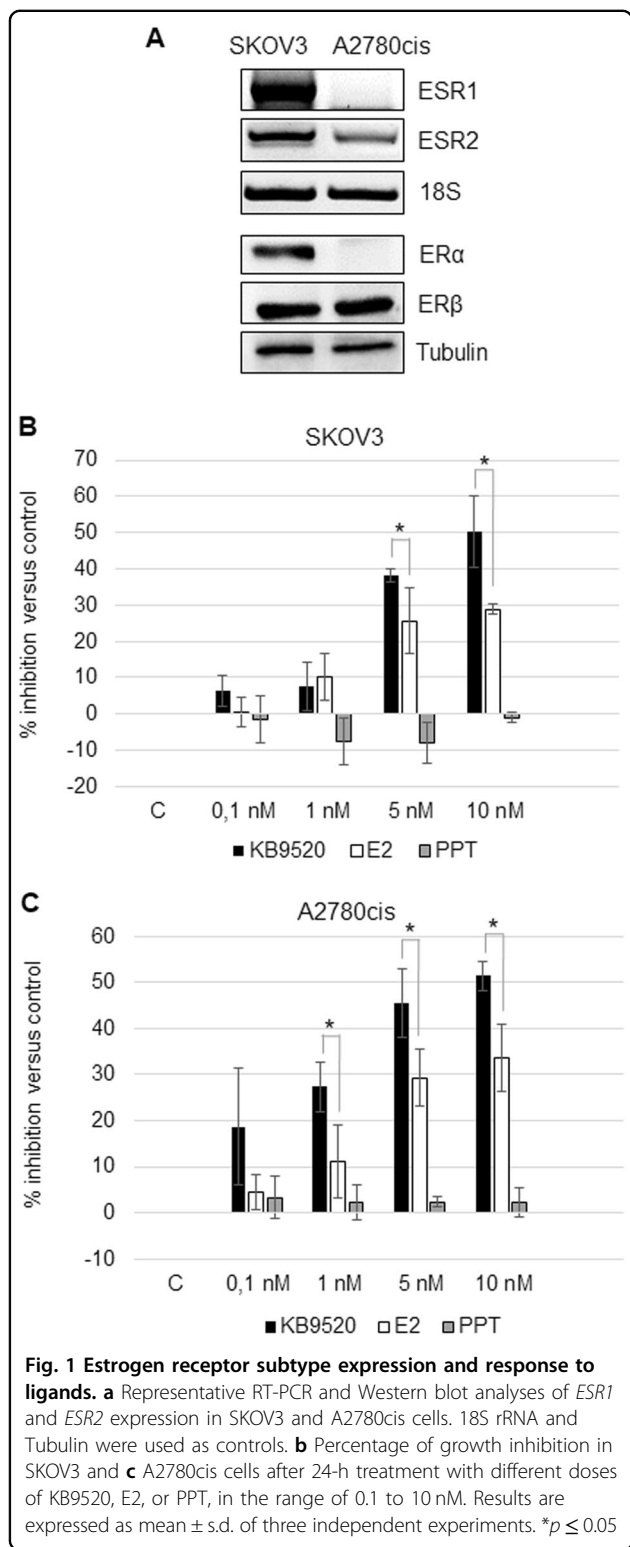
In this report we have studied the role and effect of ER α and ER β in response to the natural hormone estradiol (E2) and the subtype-selective agonists PPT (ER α -selective) and KB9520 (ER β -selective) in the human ovarian cancer cell lines SKOV3 and A2780cis. Furthermore, we have characterized the role of KDM6B and SIRT1 for the tumor inhibitory mechanism of ER β in the presence of its selective agonist KB9520.

Altogether, the preclinical and clinical data on OEC suggest that selective targeting of ER β may provide a significant improvement of existing therapy for the treatment of ovarian cancer, at least for the patient population that expresses ER β .

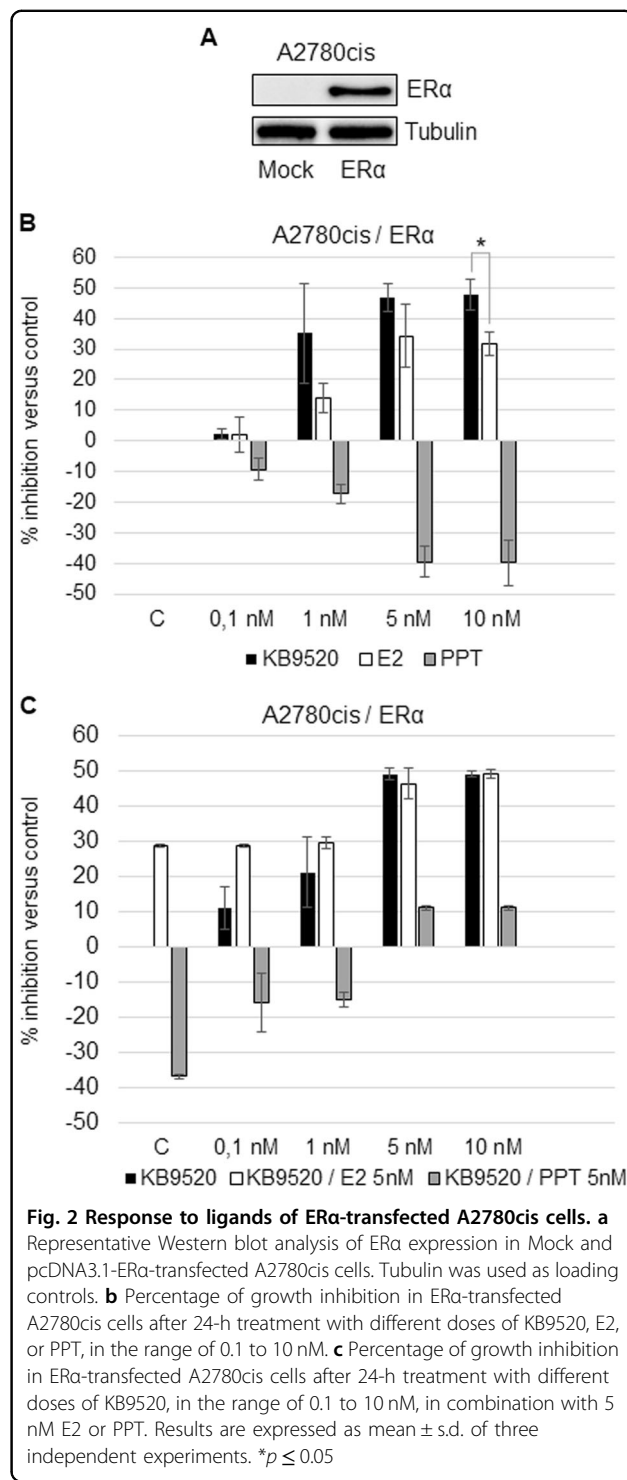
Results

ER subtype expression and response to ligands

The expression of the two ER subtypes, ER α and ER β , was investigated in the human ovarian cancer cell lines SKOV3 and A2780cis, respectively. The SKOV3 cells were shown to express both ER subtypes at gene and protein levels, whereas the A2780cis cells only expressed ER β (Fig. 1a). The effect of the non-subtype-selective agonist E2 and the two ER subtype-selective ligands KB9520 and PPT on cell growth was explored (Fig. 1b, c). The ER β -selective agonist KB9520 resulted in a robust, concentration-dependent, inhibition of cell growth of both SKOV3 and A2780cis cells, with a maximum at 5–10 nM. The ER α -selective agonist PPT caused only a weak concentration-dependent stimulation of SKOV3 cell growth, whereas the growth of the ER α -negative A2780cis cells was unaffected. E2 inhibited the growth of A2780cis cells in a concentration-dependent fashion, and unexpectedly, of also SKOV3 cells, with a similar efficacy as the ER β -selective agonist KB9520. This efficacious and unexpected cell growth inhibitory effect of E2 in the SKOV3 cells may, however, be explained by the reported loss-of-function mutation of ER α in these cells²³. Because of the mutated ER α in SKOV3 cells, we transfected A2780cis cells with a wild-type ER α expression vector (Fig. 2a). As shown in Fig. 2b, PPT resulted in a robust stimulation of A2780cis/ER α cell growth as compared to SKOV3 cells (Fig. 1b), with maximum stimulation at a concentration of 5–10 nM. E2 still caused inhibition of cell growth, similar to that of KB9520, despite the expression of wild-type ER α (Fig. 2b). In combination with a fixed concentration of PPT (5 nM), KB9520 inhibited the growth stimulatory effect of PPT in a concentration-dependent fashion (Fig. 2c). Combination

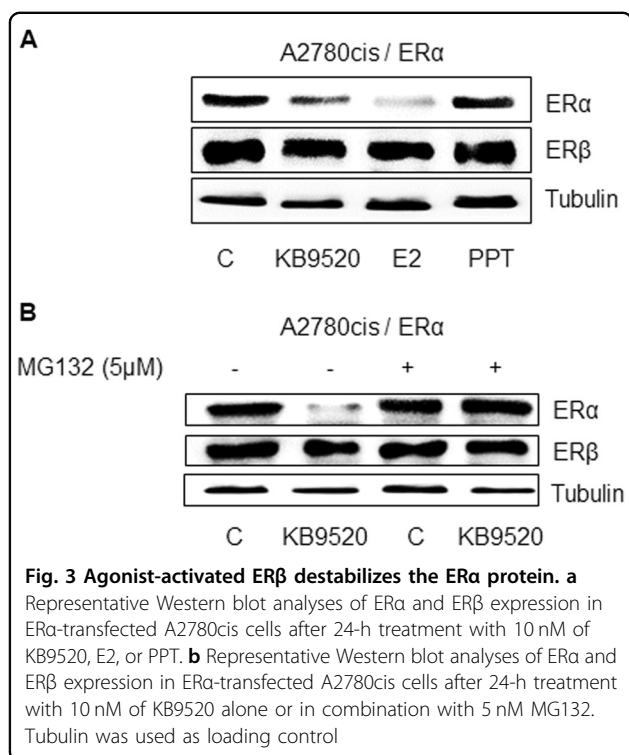


of a fixed concentration of E2 with increasing concentrations of KB9520 resulted in an additive effect on cell growth inhibition.



Agonist-activated ER β destabilizes the ER α protein

It has previously been reported that ER β has an inhibitory effect on ER α expression and activity in BG-1 epithelial ovarian cancer cells^{12, 13}. To explore the potential mechanism of the inhibitory effect of ER β on ER α activity in A2780cis cells, we treated the A2780cis/



ER α cells with E2 and KB9520 for 24 h. As shown in Fig. 3a, the ER β -selective agonist KB9520 and the non-ER subtype-selective agonist E2 resulted in decreased ER α protein levels as compared to PPT and untreated A2780cis/ER α cells. To further investigate this effect of ER β on ER α protein stability, the A2780cis/ER α cells were treated with KB9520 in the absence or presence of the proteasome inhibitor MG132 (Fig. 3b). Treatment with MG132 resulted in stabilization of the ER α protein at control levels in the presence of KB9520, suggesting that ER β plays a role in proteasome-mediated ER α protein degradation.

Additive effect of KB9520 on cisplatin and paclitaxel sensitivity in A2780cis and SKOV3 cells

The ER β -selective agonist KB9520 had a synergistic effect on cisplatin sensitivity in the malignant pleural mesothelioma REN cell line in vitro and in vivo²⁴. In this study, treatment of A2780cis and SKOV3 cells with a fixed concentration of KB9520, on top of increasing concentrations of cisplatin, resulted in an additive effect over the full range of cisplatin concentrations (Fig. 4a, S1A, B), whereas PPT was without effect. Treatment with E2 also resulted in an additive effect in combination with cisplatin but less pronounced than KB9520 (data not shown). KB9520 treatment also increased the level of cleaved PARP1 in combination with cisplatin (Fig. 4b, c).

Additive growth inhibition of A2780cis cells was also observed when cells were treated with KB9520 in combination with paclitaxel (Fig. S2).

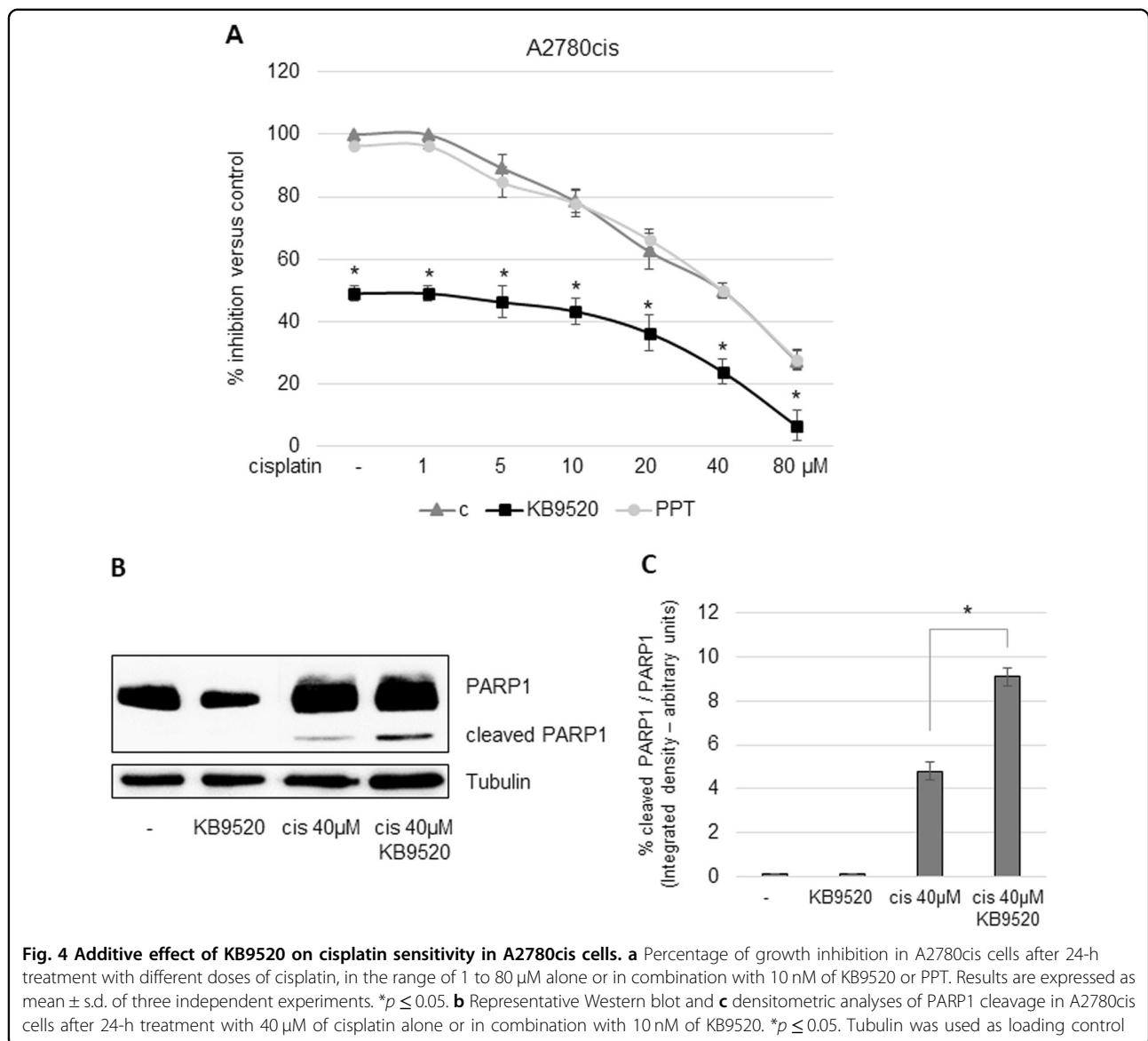
Role of KDM6B and SIRT1 for the expression and function of ER β and its response to KB9520 in A2780cis cells

We have previously reported an important role for the Jumonji domain containing 3 histone demethylase KDM6B for the expression of ER β in human epithelioid and biphasic pleural mesothelioma cell lines²⁵. In this report we observed that KB9520-activated ER β increased the levels of *KDM6B* almost 20-fold, whereas the expression of the methyltransferase *EZH2* was unaffected (Fig. S3). Knock-down of *KDM6B* affected the function of ER β as a ligand-dependent transcription factor. siRNA inhibited expression of *KDM6B* resulted in increased expression of SIRT1, at both the mRNA and protein levels, and decreased transcripts of the ER β gene (*ESR2*), with a subsequent consequence on the ER β protein level (Fig. 5a, b). Whereas acetylation of ER β and complex formation with p300 was increased in a KB9520-dependent fashion in control cells (Fig. 5c), knock-down of KDM6B led to decreased acetylation of the ER β protein and its complex formation with p300, irrespective of the presence or absence of KB9520 (Fig. 5d). In addition, the cellular sensitivity to cisplatin and the additive effect of KB9520 was lost (Fig. 5e, f, S4). Also the KB9520-mediated inhibition of *SIRT1* expression was significantly reduced following knock-down of *KDM6B* (Fig. 5g, h). KB9520 did not induce ER α acetylation and, in addition, led to reduced p300:ER α association (Fig. S5).

Inhibition of *SIRT1* expression led, on the other hand, to increased *KDM6B* expression and increased levels of ER β protein, without increased transcription of the *ESR2* gene (Fig. 6a, b). Absence of SIRT1 expression and activity (by gene silencing or inhibition of protein catalytic activity by EX527) resulted in increased ligand-dependent acetylation of ER β and re-association with acetylated p300 (Fig. 6c, S6A, B). Moreover, SIRT1 depletion restored sensitivity to KB9520 in combination with cisplatin (Fig. 6d, e). To strengthen our suspicion that SIRT1 has a more direct role in the deacetylation of ER β , A2780cis cells were treated with and without KB9520 for a shorter time point than 2 h. Cell extracts immunoprecipitated with a p300 antibody showed that SIRT1 exists in a complex with p300 and ER β and that this association is ligand dependent (Fig. 6f).

Discussion

The standard of care for the management of advanced ovarian cancer has been unchanged for many years and includes maximum cytoreductive surgery followed by platinum-based chemotherapy (carboplatin or cisplatin) in combination with paclitaxel^{2, 8}. Although the response



rate for first-line carboplatin and paclitaxel is 70–80%, the majority of women with advanced ovarian cancer will relapse or progress and eventually develop chemotherapy-resistant disease^{2, 9}. Clinical trials in the search for innovative, targeted therapies for treatment of ovarian cancer is ongoing but so far no standard second-line treatment stands out as superior with regard to efficacy or safety.

It has previously been reported that ER β mediates ovarian cancer cell growth repression by decreasing the cellular content of, among others, retinoblastoma, phospho-AKT, cyclin D1, and A2 as well as upregulating the cyclin-dependent kinase inhibitor p21^{13,14,26, 27}. In this report we have focused on other aspects of ER β function and activity as a ligand-activated transcription factor and growth inhibitor of tumor cells.

We initiated the studies on the human ovarian cancer cell lines, SKOV3 and A2780cis, by determining their expression of ER α and ER β , respectively, and by exploring the effect of the non-subtype-selective, natural hormone E2, which binds to ER α and ER β with similar affinity, and the two subtype-selective agonists KB9520 (ER β -selective) and PPT (ER α -selective), on the growth of these cells (Fig. 1). In the presence of both ER subtypes, we could show that KB9520 had a dominant inhibitory effect on cell growth, antagonizing the growth stimulatory effect of PPT (Fig. 2), which is in agreement with published results on DPN and PPT on the mouse mammary cell line HC11²⁸. Perhaps less expected was that E2 also exerted a dominant cell growth inhibition, with the same efficacy as KB9520, despite the presence of also ER α (Fig. 2). This result is in disagreement with published results on the mouse

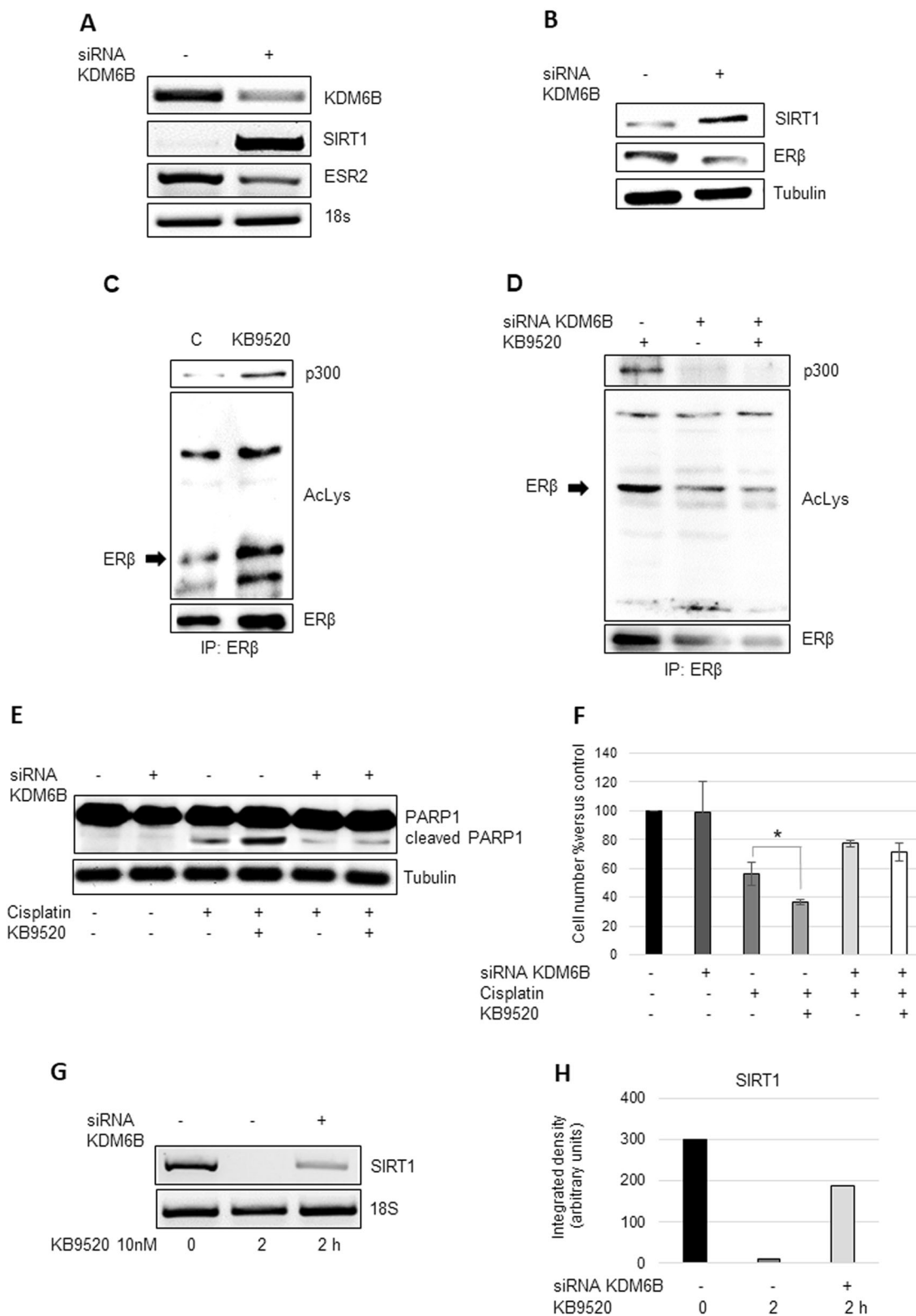


Fig. 5 (See legend on next page.)

Fig. 5 Role of KDM6B for the expression and function of ER β and its response to KB9520 in A2780cis cells. **a** Representative RT-PCR and **b** Western blot analyses of *KDM6B*, *SIRT1*, and *ESR2* expression in A2780cis cells transfected 48 h with non-specific or *KDM6B*-specific siRNAs. 18S rRNA and Tubulin were used as controls. **c** Immunoprecipitation of ER β , from lysates of A2780cis cells treated or not with 10 nM of KB9520, 2 h; lysine acetylation and co-immunoprecipitated proteins were detected by Western blot analyses using the respective antibodies (AcLys, ER β , and p300). **d** Immunoprecipitation of ER β , from lysates of A2780cis cells transfected 48 h with non-specific or *KDM6B*-specific siRNAs and treated or not with 10 nM of KB9520, 2 h; lysine acetylation and co-immunoprecipitated proteins were detected by Western blot analyses using the respective antibodies (AcLys, ER β , and p300). **e** Representative Western blot analysis of PARP1 cleavage in A2780cis cells transfected with non-specific or *KDM6B*-specific siRNAs after 24 h treatment with 40 μ M of cisplatin alone or in combination with 10 nM of KB9520. Tubulin was used as loading control. **f** Percentage of viable A2780cis cells transfected with non-specific or *KDM6B*-specific siRNAs after 24 h treatment with 40 μ M of cisplatin alone or in combination with 10 nM of KB9520. Results are expressed as mean \pm s.d. of three independent experiments. * $p \leq 0.05$. **g** Representative RT-PCR, and **h** respective densitometry, of *SIRT1* expression in A2780cis cells transfected with non-specific or *KDM6B*-specific siRNAs for 48 h and then treated or not with 10 nM of KB9520 for 2 h. 18S rRNA was used as housekeeping gene

mammary cell line HC11²⁸. This discrepancy may, however, be due to the difference in origin of the cell lines since our E2 results are in agreement with published data in ER β overexpressed, human ovarian, and breast cancer cell lines that express endogenous ER α ; ER β has a dominant, negative influence on ER α activity^{12,13,29, 30}.

That the proteasome inhibitor MG132 prevented loss of the ER α protein, following activation of ER β with KB9520, suggests that ER β is responsible for targeting ER α for proteasome pathway degradation (Fig. 3a, b).

Due to the ER β -mediated loss of the ER α protein, presented data strongly support our notion that the growth inhibition by E2, in the presence of both ER subtypes, is mediated by ER β . However, also other mechanisms may explain the dominant growth inhibitory effect of ER β such as: formation of ER α :ER β heterodimer in the presence of E2 and that ER β dictates the activity of the heterodimer, competition with ER α for binding to sites on DNA within target genes or competition for interaction with coregulatory proteins (Fig. S5)^{29–31}.

Although we received a similar increase in cleaved PARP1 by KB9520 in combination with cisplatin in the A2780cis cells as in the human malignant pleural REN cell line, we did not observe a synergistic effect of cell growth inhibition (Fig. 4, S1)²⁴. The mechanistic difference in response, additive vs. synergism, is unknown to us and needs further and deeper investigation.

Posttranslational acetylation of ER α was recently reported to play an important role for its activity as a ligand-activated transcription factor³² and for its stability³³. In this study we can for the first time present data showing that also acetylation of ER β is important for its function as a ligand-dependent regulator of cellular events (Figs. 5, 6).

That KDM6B is important for ER β expression was previously reported by us²⁵. In this report, we show that KDM6B is also important for ER β function. Ligand-dependent acetylation of ER β and its interaction with p300 was lost following depletion of KDM6B (Fig. 5c compared to 5d). Moreover, the ligand-dependent,

additive effect of ER β on PARP1 cleavage in combination with cisplatin vanished (Fig. 5e, f) and the ability of ER β to downregulate SIRT1 expression in a ligand-dependent fashion was lost (Fig. 5g, h). Whether the transcriptional regulation of the SIRT1 gene by ligand-activated ER β is direct or indirect through KDM6B needs further studies; KB9520-activated ER β upregulated *KDM6B* expression approximately 20-fold (Fig. S3).

SIRT1 was found to exist in a complex with ER β and p300 at the 1 h time point but not at 2 h or later, implying that deacetylation of ER β is a sequential event (Fig. 6e compared to 6c and S6). We speculate that loss of ER β acetylation follows SIRT1-mediated deacetylation of lysine residues 1020/1024 within the cell cycle regulatory domain 1 of p300, which thereafter results in reduced p300-mediated ER β acetylation^{32–35}.

Loss of ER β acetylation and thereby ligand-dependent ER β function may be due to decreased DNA-binding activity, inability to interact with coregulators for negative or positive transcriptional gene regulation, changed sub-cellular location or decreased binding affinity for KB9520³².

Knock-down of SIRT1 caused increased levels of ER β protein without increased transcription of the *ESR2* gene. The mechanism for this effect is unclear but may be a result of a stabilized ER β protein when acetylated.

Which lysine residues in ER β that are targets for acetylation need further studies. In conclusion, in this report, we have provided data showing that the function and ligand-dependent activity of ER β , as an ovarian tumor cell growth inhibitor, is dependent of posttranslational acetylation and that the level of KDM6B and the NAD⁺-dependent deacetylase SIRT1 play important roles in this process. We have also demonstrated that selective activation of ER β is additive to cisplatin and paclitaxel in the inhibition of ovarian tumor cell growth and that ER β has a dominant cell regulatory effect over ER α , even in the presence of E2. Moreover, we have shown that KB9520-activated ER β has a strong stimulatory effect on *KDM6B* expression, that most likely add to the ER β -mediated

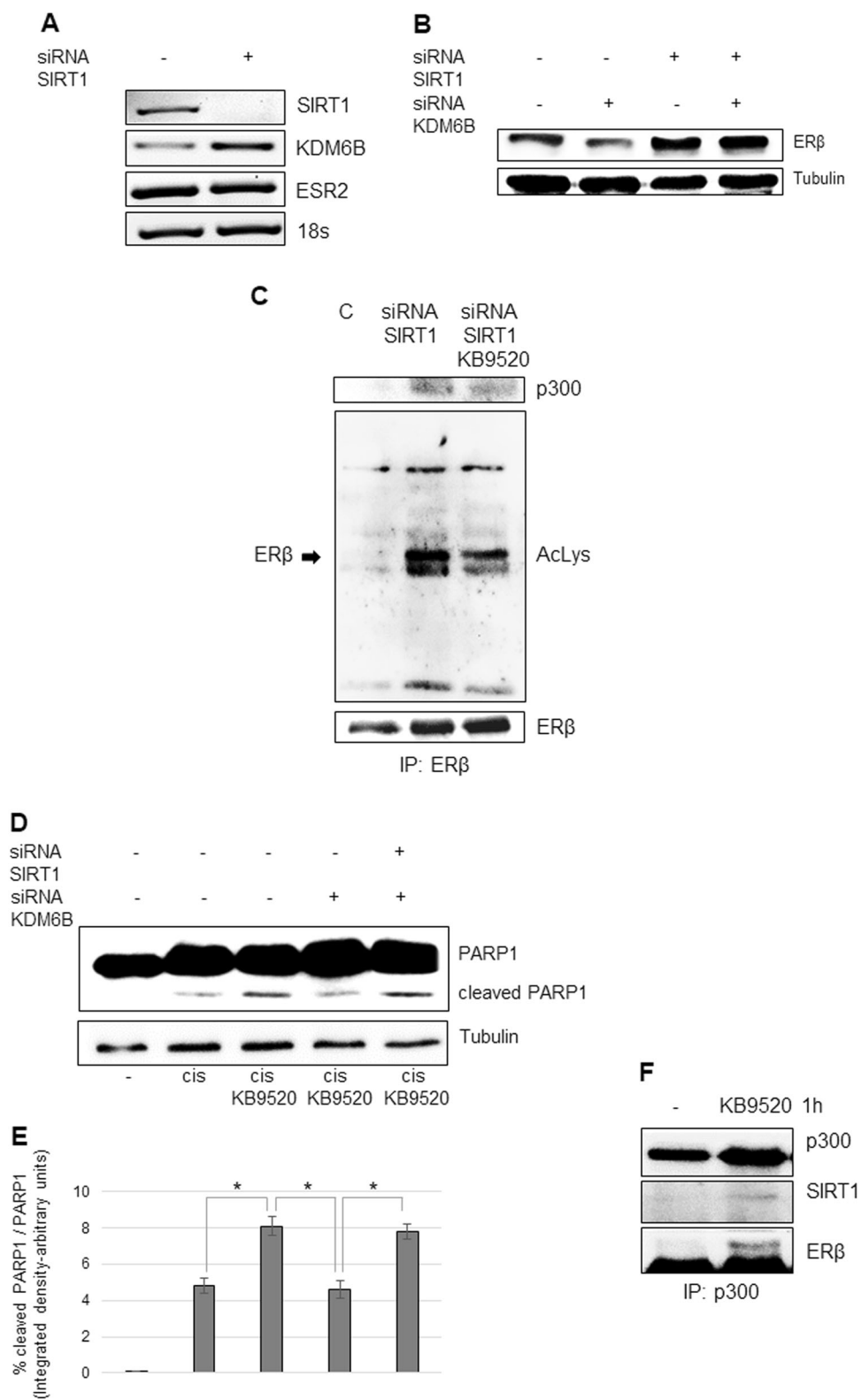


Fig. 6 (See legend on next page.)

Fig. 6 Role of SIRT1 for the expression and function of ER β and its response to KB9520 in A2780cis cells. **a** Representative RT-PCR of *SIRT1*, *KDM6B*, and *ESR2* expression and **b** Western blot analyses of ER β expression in A2780cis cells transfected with non-specific or *SIRT1* and *KDM6B*-specific siRNAs. 18S rRNA and Tubulin were used as controls. **c** Immunoprecipitation of ER β , from lysates of A2780cis cells transfected for 48 h with non-specific or *SIRT1*-specific siRNAs and then treated or not with 10 nM of KB9520, 2 h; lysine acetylation and co-immunoprecipitated proteins were detected by Western blot analyses using the respective antibodies (AcLys, ER β , and p300). **d** Representative Western blot and **e** densitometric analysis of PARP1 cleavage in A2780cis cells transfected with non-specific or *SIRT1* and *KDM6B*-specific siRNAs after 24 h treatment with 40 μ M of cisplatin alone or in combination with 10 nM of KB9520. Tubulin was used as loading control. * $p \leq 0.05$. **f** Immunoprecipitation of p300, from lysates of A2780cis treated or not 1 h with 10 nM of KB9520; co-immunoprecipitated proteins were detected by Western blot analyses using the respective antibodies (p300, SIRT1, and ER β)

ovarian tumor inhibitory activity. In summary, presented data suggest that selective targeting of ER β with an agonist potentiate chemotherapy efficacy for the treatment of ovarian cancer and that downregulation or inhibition of SIRT1 may further enhance its therapeutic effect.

Materials and methods

Reagents and antibodies

The polyclonal antibodies specific for ER α (H-184), ER β (H-150), SIRT1, p300 and the monoclonal antibodies specific for PARP1, acetylated-lysine, and α -tubulin were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Anti-mouse and anti-rabbit IgG peroxidase or FITC-conjugated antibodies, the proteasome inhibitor MG132, the selective SIRT1 inhibitor EX527 and chemical reagents were from Sigma-Aldrich (St Louis, MO, USA). ECL, nitrocellulose membranes, and protein assay kit were from Bio-Rad (Hercules, CA, USA). TRIzol[®], culture media, sera, and LipofectAMINE transfection reagent were from Thermo Fisher (Waltham, MA, USA). The ER α -selective agonist PPT and 17 β -estradiol were from Tocris Bioscience (Bristol, UK). The ER β -selective agonist KB9520 was originally provided by Karo Bio AB (Huddinge, SE)²⁴. Today this ligand is owned by Oasmia Pharmaceuticals AB (Uppsala, SE).

Cell cultures and transfection

The ovarian cancer SKOV3 and A2780cis cell lines were purchased from Sigma-Aldrich (St Louis, MO, USA). Cells were grown in standard conditions in RPMI medium supplemented with 10% FBS, 100 μ g/ml streptomycin, and 10 μ g/ml penicillin at 37 $^{\circ}$ C in a humidified environment containing 5% CO₂. Mycoplasma infection was excluded by the use of Mycoplasma PlusTM PCR Primer Set kit from Stratagene (La Jolla, CA, USA). Cells grown to 80% confluence in tissue culture dishes were transiently transfected with the pcDNA3.1-ER α plasmid or with specific siRNAs from Qiagen (Hilden, Germany), using the LipofectAMINE reagent as described by the manufacturer.

Proliferation assays

Cells were seeded at a density of 10×10^4 cells/well in 6-well plates in RPMI medium supplemented with 10% FBS, 100 μ g/ml streptomycin, and 10 μ g/ml penicillin and incubated overnight at 37 $^{\circ}$ C in a humidified environment containing 5% CO₂ to allow adherence. Following treatment cells were trypsinized and stained with Trypan blue. The number of cells considered viable (unstained cells) was counted in a Bürker haemocytometer within 5 min after staining.

Cell cycle analysis

For cell cycle/apoptosis analysis, 5×10^5 cells were seeded in tissue culture plates and treated with 10 nM KB9520, 40 μ M cisplatin, or the combination of the two drugs for 24 h at 37 $^{\circ}$ C in a 5% CO₂ atmosphere. After incubation, detached and suspended cells were harvested in complete RPMI and centrifuged at $500 \times g$ for 10 min. Pellets were washed with PBS, fixed in ice-cold 75% ethanol at 4 $^{\circ}$ C, treated with 100 mg/ml RNase A for 1 h at 37 $^{\circ}$ C, stained with 25 μ g/ml propidium iodide and finally analyzed by using a flow cytometer FACS (Becton Dickinson, San Jose, CA, USA) and Modfit software (Verity Software House, Topsham, ME, USA).

Cell lysis, immunoprecipitation, and immunoblot

Cells were extracted with 1% NP-40 lysis buffer (1% NP-40, 150 mM NaCl, 50 mM Tris-HCl pH 8.5 mM EDTA, 10 mM NaF, 10 mM Na₄P₂O₇, 0.4 mM Na₃VO₄) with freshly added protease inhibitors (10 μ g/ml leupeptin, 4 μ g/ml pepstatin, and 0.1 Unit/ml aprotinin). Lysates were centrifuged at $13,000 \times g$ for 10 min at 4 $^{\circ}$ C and the supernatants were collected and assayed for protein concentration with the Bio-Rad protein assay method. For immunoprecipitation experiments, 2 mg of extracted protein for each treatment were incubated with specific antibodies for 1 h at 4 $^{\circ}$ C and 50 μ l protein A-Sepharose beads. Proteins were separated by SDS-PAGE under reducing conditions. Following SDS-PAGE, proteins were transferred to nitrocellulose, reacted with specific antibodies and then detected with peroxidase-conjugate secondary antibodies and chemiluminescent ECL reagent. Digital images were taken with the Bio-Rad

ChemiDoc™ Touch Imaging System and quantified using Bio-Rad Image Lab 5.2.1.

RNA isolation and RT-PCR

Total RNA was extracted using TRIzol[®] reagent. Starting from equal amounts of RNA, cDNA used as template for amplification in the RT-PCR (5 µg), was synthesized by the reverse transcription reaction using RevertAid Minus First Strand cDNA Synthesis Kit from Fermentas-Thermo Scientific (Burlington, ON, Canada), using random hexamers as primers, according to the manufacturer's instructions. 20 ng of cDNA were used to perform RT-PCR amplification.

The primers sequences were: *ESR1*, Fw 5'-AACAAAGGCATGGAGCATCTGT-3' and Rev 5'-TGATGTAATACTTTTGCAAGG-3'; *ESR2*, Fw 5'-GTCAGGCATGCGAGTAACAA-3' and Rev 5'-GGGAGCCCTCTTTGCTTTTA-3'; *KDM6B*, Fw 5'-CCTCGAAATCCCATCACAGT-3' and Rev 5'-GTGCCTGTCAGATCCCAGTT-3'; *SIRT1*, Fw 5'-CTGGACAATTCCAGCCATCT-3' and Rev 5'-GGGTGGCAACTCTGACAAAT-3'. 18S RNA was simultaneously amplified using the primers: Fw 5'-AAACGGCTACCACATCCAAG-3' and Rev 5'-CCTCCAATGGATCCTCGTTA-3'.

The real-time RT-PCR was performed using the double-stranded DNA binding dye SYBR Green PCR Master Mix (Fermentas-Thermo Scientific) on an ABI GeneAmp 7000 Sequence Detection System machine, as described by the manufacturer. The instrument, for each gene tested, obtained graphical cycle threshold values automatically. Triplicate reactions were performed for each marker and the melting curves were constructed using Dissociation Curves Software (Applied Biosystems, CA, USA), to ensure that only a single product was amplified.

The primers sequences were: 18S, Fw 5'-CCCCTCGGCACCTTACG-3' and Rev 5'-TTTCAGCCTTGCAGCCATACT-3'; *KDM6B*, Fw 5'-CCTCGAAATCCCATCACAGT-3' and Rev 5'-GTGCCTGTCAGATCCCAGTT-3'; *EZH2*, Fw 5'-GCCAGACTGGGAAGAAATCTG-3' and Rev 5'-TGTGTTGGAAAATCCAAGTCA-3'.

Statistical analysis

Statistical evaluation of the differential analysis was performed by one-way ANOVA and Student's *t*-test.

Acknowledgements

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Authors' contributions

G.P. carried out cellular and molecular studies and made substantial contributions to analysis and interpretation of data. S.N. and L.M. participated

in the design and coordination of the study and drafted the manuscript. All authors have read and approved the final manuscript.

Conflict of interest

The authors declare that they have no conflict of interest.

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