PRECLINICAL STUDIES

Growth inhibition induced by antiprogestins RU-38486, ORG-31710, and CDB-2914 in ovarian cancer cells involves inhibition of cyclin dependent kinase 2

Alicia A. Goyeneche · Erin E. Seidel · Carlos M. Telleria

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Summary Antiprogestins have been largely utilized in reproductive medicine, yet their repositioning for oncologic use is rapidly emerging. In this study we investigated the molecular mediators of the anti-ovarian cancer activity of the structurally related antiprogestins RU-38486, ORG-31710 and CDB-2914. We studied the responses of wt p53 OV2008 and p53 null SK-OV-3 cells to varying doses of RU-38486, ORG-31710 and CDB-2914. The steroids inhibited the growth of both cell lines with a potency of RU-38486 > ORG-31710 > CDB-2914, and were cytostatic at lower doses but lethal at higher concentrations. Antiprogestin-induced lethality associated with morphological features of apoptosis, hypodiploid DNA content, DNA fragmentation, and cleavage of executer caspase substrate PARP. Cell death ensued despite RU-38486 caused transient up-regulation of anti-apoptotic Bcl-2, ORG-31710 induced transient up-regulation of inhibitor of apoptosis XIAP, and CDB-2914 up-regulated both XIAP and Bcl-2. The antiprogestins induced accumulation of Cdk inhibitors p21cip1 and p27kip1 and increased association of p21^{cip1} and p27^{kip1} with Cdk-2. They also promoted nuclear localization of p21cip1 and p27kip1, reduced the nuclear abundances of Cdk-2 and cyclin E, and blocked the activity of Cdk-2 in both nucleus and cytoplasm. The cytotoxic potency of the antiprogestins correlated with the magnitude of the inhibition of Cdk-2 activity, ranging from G1 cell cycle arrest towards cell

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A. A. Goyeneche · E. E. Seidel · C. M. Telleria (☒)
Division of Basic Biomedical Sciences,
Sanford School of Medicine of The University of South Dakota,
Vermillion, SD, USA

e-mail: carlos.telleria@usd.edu

death. Our results suggest that, as a consequence of their cytostatic and lethal effects, antiprogestin steroids of well-known contraceptive properties emerge as attractive new agents to be repositioned for ovarian cancer therapeutics.

Keywords Cyclin dependent kinase $2 \cdot p21^{cip1} \cdot p27^{kip1}$. Antiprogestins · Ovarian cancer

Introduction

The first antiprogestin synthesized was RU-38486 (a.k.a. "RU-486"), now named mifepristone [1]. RU-38486 has been mainly used as blocker of progesterone receptors in the uterus, where it increases the sensitivity to myometrial contractions induced by prostaglandin analogues, leading to early termination of pregnancy [2]. Yet RU-38486 is useful for other reproductive indications, such as oral contraception, menstrual cycle regulation and emergency contraception [3]. More recently, RU-38486 emerged to treat endocrine-related diseases such as uterine leiomyoma and endometriosis [4].

The potential use of RU-38486 in oncology is promising. In non-reproductive tissues, RU-38486 inhibited the growth of gastric cancer cell lines [5] and of meningioma cells [6]. In reproductive tissues, RU-38486 blocked proliferation and killed benign and malignant endometrial cancer cells [7]. In prostate cancer, RU-38486 blocked growth of androgensensitive and androgen-insensitive LNCaP cells in vivo and in vitro [8]. In breast cancer, RU-38486 inhibited the growth of T-47D cells [9], and in MCF-7 cells it had an additive lethal effect when combined with antiestrogen tamoxifen [10]. In MCF-7 cells, RU-38486 had a synergistic lethal interaction with the Chk1 inhibitor 7-hydroxystaurosporine (UCN-01) [11] or with 4-hydroxystamoxifen [12]. Also, RU-



38486 blocked the growth of MCF-7 sublines resistant to 4-hydroxytamoxifen [13] and was lethal to progesterone receptor- and estrogen receptor-negative MDA-MB-231 cells [14]. In *p53/BRCA1*-deficient mice, RU-38486 prevented the formation of breast tumors [15]. RU-38486 also inhibited the growth of cervical adenocarcinoma cells in vitro and in vivo [16].

The action of antiprogestins in ovarian cancer has received limited attention. First in 1996 it was revealed that RU-38486 arrested OVCAR-3 and A2780 cells at the G1 phase of the cell cycle [17]. More recently we demonstrated the efficacy of RU-38486 as a single agent in an in vivo preclinical setting and found that its growth inhibitory effect was associated with inhibition of DNA synthesis, G1 cell cycle arrest, and down-regulation of transcription factor E2F1 needed for S phase progression [18]. We also demonstrated that RU-38486 inhibits the growth of ovarian cancer cells regardless of p53 genetic makeup and platinum sensitivity [19]. Finally, we have shown that cytostatic concentrations of RU-38486 added after courses of lethal platinum-based chemotherapy prevent repopulation of remnant cancer cells escaping and surviving the insult of the platinating agent [20].

ORG-31710 and CDB-2914 are two members of a family of selective progesterone receptor modulators with a similar structure to RU-38486, as they all contain a dimethylaminophenyl substitution at the 11\beta-position that confers antiprogestin activity [2, 4, 21] (Fig. 1). ORG-31710 and CDB-2914, however, were designed aiming to decrease the antagonistic effect of RU-38486 on the glucocorticoid receptor by substitutions made at the 17α side chain [4]. Limited information is available regarding the growth inhibition and oncologic value of these two antiprogestins. Studies in rats show that ORG-31710 and CDB-2914 were effective in reducing the growth of established DMBA-induced breast tumors by increasing apoptosis and blocking cell proliferation [22, 23]. In cultured human uterine leiomyoma cells, CDB-2914 inhibited cell proliferation down-regulating PCNA expression, and inducing apoptosis up-regulating PARP expression and reducing Bcl-2 abundance [24]. Further a recent randomized controlled clinical trial reported that CDB-2914 significantly reduced leiomyoma growth [25]. ORG-31710, on the other hand, increased apoptosis in human periovulatory granulosa cells [26].

Based on our previous findings on the in vitro and in vivo growth inhibitory effect of RU-38486 in ovarian cancer cells, the goal of this study was to investigate the molecular mediators of the anti-ovarian cancer activity of RU-38486 and of the two structurally related antiprogestins, ORG-31710 and CDB-2914 (Ulipristal). We report that RU-38486, ORG-31710 and CDB-2914 all are cytostatic at lower concentrations, up-regulating and promoting nuclear localization of the cyclin dependent kinase (Cdk) inhibitors p27kip1 and p21cip1, reducing nuclear abundances of Cdk-2 and cyclin E, and reducing the activity of Cdk-2. At higher concentrations the three antiprogestins blunted the activity of Cdk-2 leading to the death of the ovarian cancer cells, which was associated with morphological features of apoptosis, hypodiploid DNA content, fragmentation of the DNA, and cleavage of the executer caspase substrate PARP.

Materials and methods

Cell culture and drugs

The human ovarian carcinoma cell line OV2008 was obtained in 2003 from Dr. Howell (University of California, San Diego) and was maintained in RPMI 1640 (Mediatech, Herndon, VA) supplemented with 5% heat inactivated fetal bovine serum (Atlanta Biologicals, Lawrencenville, GA), 10 mM HEPES (Mediatech), 4 mM L-glutamine (Mediatech), 1 mM sodium pyruvate (Mediatech), 1 x non-essential amino acids (Mediatech), 100 IU penicillin (Mediatech), and 100 μg/ml streptomycin (Mediatech). SK-OV-3 ovarian cancer cells were obtained in 2003 at passage 23 from the American Tissue Culture Collection (Manassas, VA) and were routinely maintained in RPMI 1640 (Mediatech) supplemented with 5% fetal bovine serum (Atlanta Biologicals), 10 mM HEPES (Mediatech),

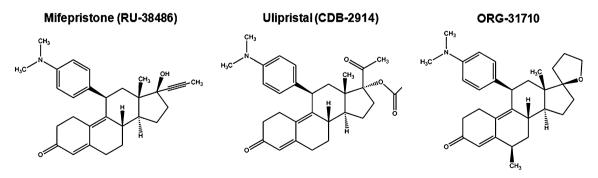


Fig. 1 Chemical structure of antiprogestins



4 mM L-glutamine (Mediatech), 0.45% D (+) glucose (Sigma Chemical Company, St. Louis, MO), 1 mM sodium pyruvate (Mediatech), 1 x non-essential amino acids (Mediatech), 100 IU penicillin (Mediatech), 100 μg/ml streptomycin (Mediatech), and 0.01 mg/ml human insulin (Roche, Indianapolis, IN). Both cell lines were cultured at 37°C in a humidified atmosphere in the presence of 5% CO₂. Treatment of the cells with RU-38486 (Sigma), ORG-31710 (generously provided by N.V. Organon, Oss, The Netherlands), or CDB-2914 (kindly provided by HRA Pharma, Paris, France) were done from 20 mM stock solutions in DMSO; the maximal concentration of DMSO in medium was 0.2% (v/v).

Cell proliferation and viability

Triplicate cultures were trypsinized, pelleted by centrifugation at 500 g for 5 min, and washed with PBS. The cells were resuspended in ViaCount reagent (Guava Technologies, Hayward, CA) and studied using the Guava ViaCount application in the Guava EasyCyte Mini microcapillary cytometer (Guava Technologies) as we previously reported [20]. When indicated, the proliferation IC50 values were determined using software designed to study drug interaction that calculates the median effective dose, Dm, which is analogous to the IC50 (Calcusyn, Biosoft, Cambridge, UK).

Cell cycle analysis

After treatment, cells were trypsinized, pelleted by centrifugation at 500*g* for 5 min, washed with PBS, and fixed with 4% paraformaldehyde. Cells were once again washed with PBS and pelleted by centrifugation at 500*g* for 5 min. Then approximately 100,000–200,000 cells were resuspended in 200 μl of cell cycle buffer [3.8 mM sodium citrate (Sigma), 7 U/ml RNase A (Sigma), 0.1% (v/v) Triton X-100 (Sigma), and 0.05 mg/ml propidium iodide (Sigma)] at a concentration of 500–1000 cells/μl. Cells were analyzed for the capacity of their DNA to bind propidium iodide utilizing the Guava EasyCyte Mini microcapillary cytometer and the cell cycle application of the CytoSoft 4.1 software (Guava Technologies).

Immunoblot analysis

Cells were scraped, pelleted, washed twice with PBS, and lysed by the addition of two volumes of NP-40 lysis buffer containing 50 mM Tris–HCl (pH 7.5), 150 mM NaCl, 0.5% NP-40 (Sigma), 50 mM sodium fluoride (Sigma), 1 mM PMSF (Sigma), 2 μ g/ml pepstatin (Sigma), 2 μ g/ml leupeptin (Sigma), 2 μ g/ml aprotinin (Sigma), and 1 mM orthovanadate (Sigma). Lysates were centrifuged at 16,000g for 15 min at 4°C, and the supernatant was considered the

whole cell extract, which was assayed for protein content using the bicinchoninic acid method (BCA; Pierce, Rockford, IL). The whole cell extracts were appropriately diluted in 3 x concentrated electrophoresis sample buffer, boiled for 10 min, and stored at -80°C until electrophoresed. Equivalent amounts of proteins (50 µg) per point were loaded in 12% (w/v) acrylamide gels, subjected to SDS-PAGE and transferred to PVDF membranes. The blots were blocked in 5% (v/v) non-fat milk in TBS containing 0.1% (v/v) Tween 20 (T). Blots were probed overnight with the appropriate dilution of each of the primary antibodies. The membranes were washed 3×5 min in TBS-T and incubated with 1: 10,000 dilution of peroxidaseconjugate secondary antibody (#111-035-003; Jackson ImmunoResearch Laboratories, West Grove, PA) for 30 min at room temperature. The blots were again washed, developed by chemiluminescence, and exposed to radiographic film. Blots were stripped and reprobed with an antibody directed against the ubiquitous proteins glyceraldehyde-3-phosphate dehydrogenase (GAPDH), or β-actin, to control for protein loading. Primary antibodies for the following proteins were used at the indicated dilutions. Cyclin E (clone HE12; $0.5 \mu g/ml$), and $p21^{cip1}$ (clone 6B6; $2 \mu g/ml$) were from BD Pharmigen (San Diego, CA); p27^{kip1} (clone 57; 1: 2,000) was from BD Transduction Laboratories (San Diego, CA); XIAP (#2042; 1:500), poly (ADP-ribose) polymerase (PARP; #9592; 1:1,000) and caspase-3 (#9662; 1:1,000) were from Cell Signaling Technology (Danvers, MA); Cdk-2 (M2; 1:1,000) and Bcl-2 (100; 1:1000) were from Santa Cruz Biotechnology (Santa Cruz, CA); GAPDH (ab9485; 1:10,000) was from Abcam Inc. (Cambridge, MA); β-actin (clone AC-15; 1:20,000) was from Sigma.

Cdk-2 immunoprecipitation and histone H1 kinase assay

An aliquot (100 µg of protein) from each NP-40 cell lysate was incubated overnight at 4°C with constant rotation in 1 ml of NP-40 lysis buffer containing 1 µg polyclonal rabbit antibody to Cdk-2 (M2; Santa Cruz Biotechnology). Immunocomplexes associated with Cdk-2 were collected after incubation for 2 h with protein A/G PLUS-Agarose beads (Santa Cruz Biotechnology). The immune-complexes were washed twice with kinase buffer [50 mM HEPES (pH 7.2), 10 mM MgCl₂, 1 mM DTT, 10 mM βglycerophosphate, and 1 mM sodium fluoride]. Subsequently, the beads were resuspended in 30 µl of kinase buffer containing 2 µg of histone H1 (Upstate Cell Signaling Solutions, Lake Placid, NY), 5 µM ATP (Upstate), and 5 μ Ci of $[\gamma^{32}P]$ ATP (MP Biomedicals, Irvine, CA). The reaction mixtures were incubated at 30°C for 30 min, the reaction was terminated with 30 µl of 2 x electrophoresis sample buffer, boiled, and separated on 12% SDS/polyacrylamide gels. Gels were stained with



Coomassie Blue (Sigma) to visualize the histone H1 bands, dried, and autoradiographed.

Subcellular fractionation

Upon treatment with vehicle or antiprogestins, cells were washed in PBS, scrapped, pelleted, and resuspended in low salt lysis buffer [10 mM HEPES (pH 7.9), 10 mM KCl, 1.5 mM MgCl₂, 1 mM DTT, 1 mM PMSF, 1 mM sodium orthovanadate, 2 µg/ml aprotinin, 2 µg/ml leupeptin, and 2 µg/ml pepstatin], incubated on ice for 15 min and homogenized with a hand pestle. The lysates where then centrifuged for 10 min at 800 x g at 4°C. The pellet represents the nuclear fraction. The supernatant was further centrifuged for 10 min at 16,000 x g at 4°C to remove any contaminant nuclei. The new supernatant was considered the cytosolic fraction. The crude nuclear fraction was washed twice with low salt lysis buffer, pelleted down for 5 min at 800 x g, resuspended in NP-40 lysis buffer (described above), and rocked for 1 h at 4°C. The suspension was centrifuged at 16,000 x g for 20 min to remove debris, and the supernatant was considered the nuclear fraction. The protein concentrations of both factions were determined as described above. Fifty µg of each of the protein fractions were subjected to SDS-PAGE, transferred to PVDF membranes, and immunoblotted with antibodies against p27kip1, p21cip1, Cdk-2, cyclin E, and GAPDH or β-actin for loading control.

4', 6-Diamidino-2-phenylindole dihydochloride (DAPI) staining and phase contrast microscopy

OV2008 or SK-OV-3 cells cultured on 6-well plates were exposed to either vehicle (DMSO), or 20 or 40 µM antiprogestins for 96 h. After treatment, detached cells were collected, centrifuged at 500g for 5 min, fixed and resuspended in 100% methanol, adhered to a microscope slide, and stained for 10 min with DAPI (Molecular Probes, Eugene, OR). Nuclear morphology was observed and photographed using a Zeiss Axiovert M200 inverted fluorescence microscope (Carl Zeiss, Thornwood, NY). Cells that remained adherent to the original chamber slide were also fixed in 100% methanol, stained with DAPI and photographed. All cell preparations were simultaneously photographed using a phase contrast objective.

DNA fragmentation

Floating and adherent cells were pelleted and digested overnight at 50°C in a buffer composed of 100 mM NaCl, 10 mM Tris HCl (pH 8.0), 25 mM EDTA (pH 8.0), 0.5% SDS and 0.1 mg/ml proteinase K (Life Technologies, Rockville, MD). The genomic DNA was extracted from

the digested cells with phenol/chloroform/isoamyl alcohol (25:24:1, v/v/v), precipitated, and digested for 60 min at 37°C with 1 µg/ml ribonuclease (deoxyribonuclease-free; Roche). After extraction and precipitation, an equal amount of DNA for each sample (2 µg) was separated by electrophoresis on a 2.5% agarose gel, impregnated with SYBR Gold nucleic acid gel stain (Molecular Probes), examined using an ultraviolet transilluminator, and photographed with the Amersham Typhoon Fluorescence imaging system (Amersham Biosciences Corp., Piscataway, NJ). A 100 bp DNA ladder (Promega, Madison, WI) was utilized for determining the size of the fragments of DNA.

Results

Antiprogestins inhibit, in a dose-related manner, the growth of p53 wild type and p53 mutant ovarian cancer cells, eliciting concentration dependent cytostatic and lethal effects

To explore whether RU-38486, ORG-31710 or CDB-2914 can inhibit the growth of ovarian cancer cells of different genetic backgrounds, we studied the response to the antiprogestins in OV2008 cells that express wild type p53 [27], and SK-OV-3 cells that carry a deletion of a single nucleotide as a consequence of which no p53 mRNA transcripts are expressed [28]. The two cell lines were exposed to vehicle or increasing concentrations of the antiprogestins for 72 h. At the end of the experiment, the cells were evaluated and analyzed by microcapillary cytometry for cell number, cell viability, and cell cycle distribution. Results shown in Fig. 2a and d illustrate that both cell lines were growth inhibited by the three antiprogestins in a dose-related manner. In OV2008 cells, RU-38486 had a growth inhibition concentration 50% or IC50 lower than that of ORG-31710 or CDB-2914 (Fig. 1b). In SK-OV-3 cells, RU-38486 and ORG-31710 had similar growth inhibition potency which was, however, higher than that of CDB-2914 (Fig. 2e). Neither RU-38486 nor ORG-31710 or CDB-2914 showed lethality towards the cells at the 20 µM concentration. At the concentration of 40 µM, however, the three antiprogestins reduced the viability of both cell lines, with the effect being more prominent in OV2008 cells (Fig. 2c) than in SK-OV-3 cells (Fig. 2f).

The cytostatic effect of the 20 μ M dose of the three antiprogestins was reinforced by the fact that after 72 or 96 h treatment the cells remained attached to the culture plate, although in a reduced number when compared to vehicle-treated controls (Fig. S1). It is interesting to note that at cytostatic concentrations the antiprogestins not only



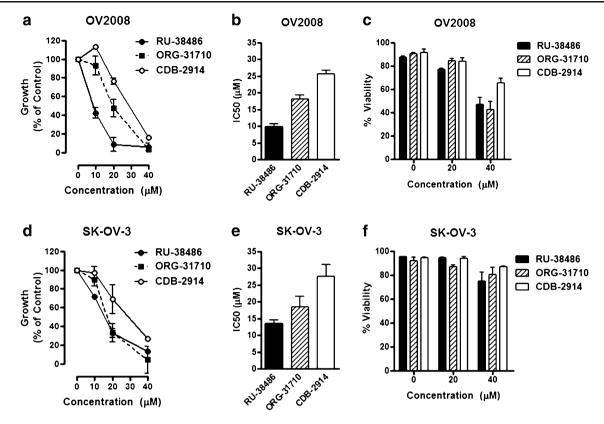


Fig. 2 Effect of antiprogestins on the growth of p53 wild type (OV2008) and p53 null (SK-OV-3) ovarian cancer cells. OV2008 a-c or SK-OV-3 d-f cells were cultured in the presence of the indicated concentrations of RU-38486, ORG-31710, or CDB-2914. The number of cells and their percent viability were recorded at the beginning of the experiment and after 3 days of treatment using exclusion fluorochromes in a microcapillary cytometer. The difference between

the number of viable cells in vehicle-treated controls at 0 h and after 3 days of culture was considered to be 100%. The growth of the treated groups is expressed as percentage of control **a** and **d**. For each dose–response curve, the amount of drug needed to achieve 50% growth inhibition (IC50) was calculated utilizing drug interaction software **b** and **e**. Cell viability is shown in separate graphs (**c** and **f**). Results are the average of three separate experiments

blocked cell proliferation but also altered the morphology of the cells. More evident in SK-OV-3 than in OV2008 cells, the cells acquired an elongated morphology. Conversely, more evident in OV2008 than in SK-OV-3 cells, scattered cells with multi-nucleation and multi-vacuolation were evident (Fig. S1).

When the cell cycle traverse of OV2008 was studied by microcytometric analysis of propidium iodide-labeled DNA particles, it was evident that 20 μ M antiprogestins had a cytostatic but not a lethal effect. This was indicated by the lack of cellular particles containing hypodiploid (Sub-G1) DNA content, the increased proportion of cells transiting the G0-G1 phase of the cell cycle, which was accompanied by a reduction in the proportion of cells allocated to the S phase, and no apparent changes in the proportion of cells transiting G2/M (Fig. 3a). When the antiprogestins were used at the 40 μ M concentration, there was an evident increase in the percentage of hypodiploid-DNA containing cellular particles that was more evident for RU-38486 and ORG-31710 when compared to CDB-1914. The lethality of high concentration antiprogestins was further evidenced for

RU-38486 and ORG-31710 by the reduction in the percentage of cells transiting G0-G1, which was not seen in CDB-2914 that still showed G0-G1 block (Fig. 3a).

To study the dynamics of the cell cycle upon antiprogestin challenge, cell cycle distribution was assessed in OV2008 cells after 12, 24, or 48 h treatment (Fig. 3b) or in SK-OV-3 cells after 24, 48, 72, or 96 h treatment (Fig. 4). To link the kinetics of the cell cycle with the growth inhibitory and lethal effects of the compounds, for these time-course studies we utilized the lethal 40 µM concentration. In OV2008 cells RU-38486 caused, after 24 h treatment, Sub-G1 DNA accumulation without any increase in G1, suggesting the cells are likely dying without arresting (Fig. 3b, upper panels, and lower left panel); however, the same concentration of ORG-31710 and CDB-2914 caused G1 arrest after 12 and 24 h treatment (Fig. 3b, upper panels). ORG-31710 showed Sub-G1 accumulation beginning at 48 h exposure with a parallel decline in G1, whereas CDB-2914 still showed G1 arrest 48 h after treatment (Fig. 3b, lower left panel). The three compounds abrogated growth as indicated by the stagnant cell number



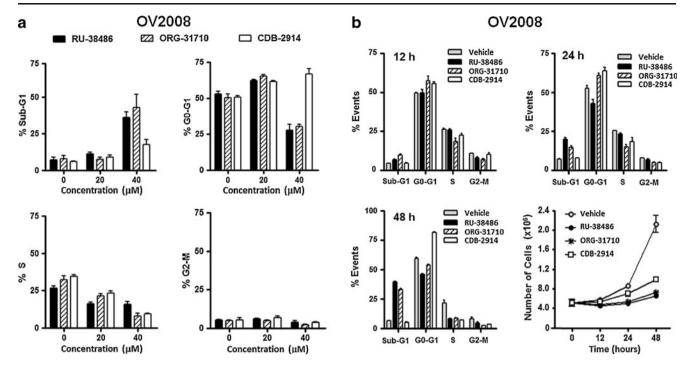


Fig. 3 Effect of antiprogestins on cell cycle distribution in OV2008 cells. **a** Logarithmically growing OV2008 ovarian cancer cells were exposed to 0, 20 or 40 µM RU-38486, ORG-31710 or CDB-2914 for 72 h. The percentage of cells with Sub-G1 DNA content (upper left), or with DNA content consistent with G0-G1 (upper right), S (lower left), and G2-M (lower right) phases of the cell cycle was determined by microcapillary cytometric analysis after propidium iodide staining of the DNA. **b** Logarithmically growing OV2008 cells were exposed

to DMSO (vehicle) or 40 μ M antiprogestins for 12 h (upper left), 24 h (upper right), or 48 h (lower left). The percentage of events consistent with Sub-G1, G0-G1, S, and G2-M DNA content was analyzed after propidium iodide staining utilizing microcapillary cytometry. The number of cells was recorded at the beginning of the experiment and after different periods of incubation with or without treatments (lower right)

in the treated groups along the study (Fig. 3b, lower right panel). In SK-OV-3 cells, the kinetics of the cell cycle was slightly different. Twenty-four, 48 or 72 h after treatment with any of the compounds, the cells accumulated in G1 with a reduced proportion of cells still in S phase when compared to vehicle-treated controls (Fig. 4a-c). After 96 h, however, the three antiprogestins triggered accumulation of Sub-G1 DNA particles when compared to the values at 72 h (Fig. 4d). The three compounds abrogated growth along the study when compared to vehicle-treated controls (Fig. 4e).

Lethal concentrations of antiprogestins induce morphological and biochemical changes consistent, at least in part, with apoptotic cell death

After culturing OV2008 or SK-OV-03 cells with 20 μ M antiprogestins, no lethality was observed; however, signs of loss of viability were found at the 40 μ M concentration (Figs. 2c and f, 3a and b, and 4). Morphological studies show that the number of OV2008 cells still adherent after 96 h treatment with 40 μ M antiprogestins is minimal, yet with differences among the treatments, with the effects being those of RU-38486 > ORG-31710 > CDB-2914. The

majority of the cells at this point in time appear detached and with morphological features similar to those shown by cisplatin-treated cells (Fig. 5a). SK-OV-3, although less sensitive to the lethal effects of the antiprogestins, show a remarkably reduced number of still adherent cells after 96 h treatment with a concomitant enhanced number of floating cells with similar morphological features to those shown by cisplatin-treated cells (Fig. 5b). Fragmentation of the DNA was clearly observed after 60 h exposure to 40 μ M RU-38486 and ORG-31710 in OV2008 cells, whereas the effect was less pronounced in CDB-2914-treated cells at the same point in time (Fig. 5c, left panel). For SK-OV-3 cells, treatment with 40 μ M of the three compounds showed DNA laddering upon 120 h incubation (Fig. 5c, right panel).

To further explore the signaling molecules involved in the lethal action of antiprogestins in ovarian cancer cells, we treated OV2008 cells with 40 μ M antiprogestins and collected whole cell extracts after 24, 48 or 72 h incubation. Results in Fig. 5d show that RU-38486 caused cleavage of the full-length caspase-3, 116 kDa PARP substrate into the 86 kDa cleaved and inactive form 24 h after treatment, an effect that was more evident at 48 h, in association with down-regulation of cell cycle



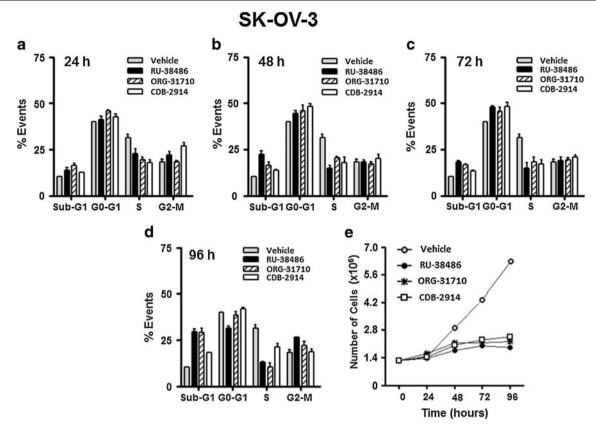


Fig. 4 Effect of antiprogestins on cell cycle distribution in SK-OV-3 cells. Logarithmically growing SK-OV-3 cells were exposed to DMSO (vehicle) or 40 μ M RU-38486, ORG-31710 or CDB-2914 for 24 h a, 48 h b, 72 h c or 96 h d. The percentage of events consistent with Sub-G1, G0-G1, S, and G2-M DNA content was

analyzed after propidium iodide staining utilizing microcapillary cytometry. The number of cells was recorded at the beginning of the experiment and after different periods of incubation with or without treatments ${\bf e}$

related proteins Cdk-2 and p21cip1, and cleavage of p27kip1. RU-38486-induced lethality was associated with down-regulation of the inhibitor of apoptosis protein XIAP but, curiously, it occurred in the presence of high levels of anti-apoptotic protein Bcl-2 when compared to vehicle-treated controls. ORG-31710-induced cell death associated with PARP cleavage and down-regulation of Cdk-2, in the presence of high levels of Cdk inhibitors p21cip1 and p27kip1. In contrast to RU-38486, ORG-31710-induced cytotoxicity occurred with higher levels of XIAP and no changes in Bcl-2 when compared to vehicle-treated controls. Finally, lethality associated to CDB-2914 occurred with up-regulation and cleavage of PARP, up-regulation of XIAP, and up-regulation and cleavage of Bcl-2, in the presence of high levels of p21cip1 and p27kip1. Altogether results in Fig. 5 suggest that concentrations of antiprogestins higher than 20 µM are lethal to p53-wild type OV2008 and p53-null SK-OV-3; the molecular mediators of the cell death mechanisms, as well as their interrelationship with cell cycle regulatory proteins, differ among the antiprogestins and warrant further investigations.

Cytostatic concentrations of antiprogestins increase association of p21^{cip1} and p27^{kip1} with Cdk-2, promote nuclear localization of p21^{cip1} and p27^{kip1}, reduce nuclear abundances of Cdk-2 and cyclin E, and inhibit the activity of Cdk-2

In association with cyclin E, Cdk-2 is essential to drive the cell cycle through the G1-S transition and into the S phase whereas p21cip1 and p27kip1 are usually involved in inhibiting cyclin E/Cdk-2 activity [29]. We questioned whether the activity of Cdk-2 was related to the growth inhibitory effect of the antiprogestins. OV2008 cells were cultured for 12, 24, or 48 h in the presence of 20 µM RU-38486, ORG-31710, or CDB-2914 (Fig. 6a). Twelve hours after treatment with RU-38486, p21cip1 levels increased with respect to vehicle-treated controls. Such increase was even higher 24 h later, and persisted 48 h following drug exposure. Treatment with ORG-31710 also led to an increase in p21^{cip1} levels 12 h after exposure, a further increase at 24 h, but it was followed by a decline at 48 h. Finally, CDB-2914 induced a mild increase in p21cip1 levels at 12 h, followed by higher levels at 24 h and a further



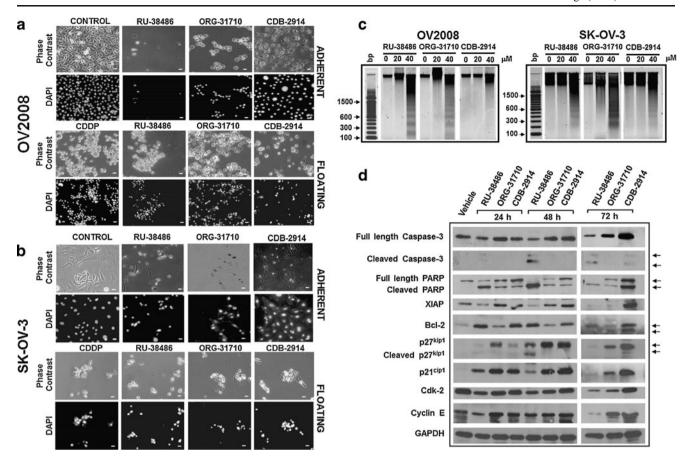


Fig. 5 Lethality of antiprogestins towards ovarian cancer cells. OV2008 or SK-OV-3 cells were cultured in the presence of 40 μM RU-38486, ORG-31710 or CDB-2914. After 60 h (OV2008) or 120 h (SK-OV-3), floating cells were collected, adhered to a microscope slide, stained with DAPI, and microscopic phase contrast and fluorescence images were obtained from the same fields. As positive control of apoptotic cell death, floating cells from cisplatin (CDDP) treated cells were included in the morphological studies **a** and **b**. Adherent untreated cells are also shown (CONTROL). Scale bar, 20 μm. **c** A similar experiment was done in which all floating and

attached cells were pelleted, total DNA was isolated, subjected to agarose electrophoresis, stained with SYBR Gold nucleic acid stain, and photographed with the Amersham Typhoon fluorescence imaging system. A 100 base pair (bp) marker was run in parallel. (d) OV2008 cells were exposed to DMSO (Vehicle) or 40 μ M RU-38486, ORG-31710 or CDB-2914 for 72 h. Whole protein extracts were obtained and separated by electrophoresis, and immunoblots were probed with the indicated antibodies to recognize cell cycle and cell death related proteins. The housekeeping gene GAPDH was used as protein loading control

increase at 48 h. A similar pattern of induction was observed for p27^{cip1} with small differences. RU-38486 induced a slight increase in p27^{kip1} levels at 12 h but a larger increase at 24 and 48 h, whereas ORG-31710 needed 24 h exposure to increase p27^{cip1}, and CDB-2914 only slightly increased p27^{cip1} levels even after 48 h exposure. No major modification in the total protein levels of Cdk-2 and cyclin E was observed in these experiments. When a similar experiment was conducted in SK-OV-3 cells p21^{cip1} and p27^{kip1} increased in response to the antiprogestins as well, but there was a pronounced decline in the abundance of Cdk-2 not observed in OV2008 cells (Fig. S2).

Because nuclear localization of Cdk-2 is required for its full activation and cell cycle progression [30, 31], we analyzed whether the antiprogestins affect the nucleocytoplasmic trafficking of the Cdk inhibitors p21^{cip1} and p27^{kip1}, cyclin E and Cdk-2. OV2008 cells treated with

vehicle or 20 μM antiprogestins for 48 h were subjected to subcellular fractionation and Western blot analysis. Results in Fig. 6b reveal that the antiprogestins induced an increase in nuclear localization of p21cip1 and p27kip1, which correlated with decreased Cdk-2 and cyclin E nuclear levels. As a consequence of the observed decrease in Cdk-2 and cyclin E nuclear abundance, together with the increase in the abundances of Cdk inhibitors, we asked whether such changes were reflected in the modulation of the activity of Cdk-2. We immunoprecipitated Cdk-2 from vehicle-treated OV2008 cells or from OV2008 cells treated with 20 µM antiprogestins for 48 h and did an in vitro kinase assay using histone H1 as substrate. The results in Fig. 6c show that the activity of Cdk-2 was inhibited in both nucleus and cytoplasm, and such inhibition was stronger when the cells were exposed to RU-38486 or ORG-31710, whereas CDB-2914 caused the weakest



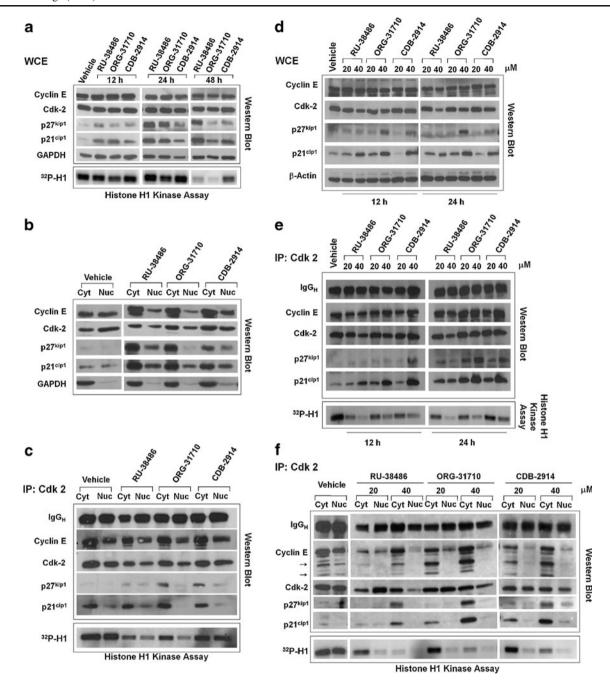


Fig. 6 Effect of antiprogestins on cell cycle regulatory proteins in ovarian cancer cells. OV2008 cells were exposed to DMSO (Vehicle), 20 **a-e** or 40 μM **d** and **e** RU-38486, ORG-31710, or CDB-2914 for the indicated times **a**, **d**, and **e** or 48 h **b** and **c**. **a** Whole protein extracts (WCE) were obtained and separated by electrophoresis, and immunoblots were probed with the indicated cell cycle related antibodies. Whole protein extracts were also immunoprecipitated with anti-Cdk-2 antibody and assayed for their capacity to phosphorylate histone H1 in vitro in the presence of ³²P ATP. **b** Isolation of nuclear and cytosolic fractions was achieved, proteins from each fraction were obtained and separated by electrophoresis, and immunoblots were probed with the indicated antibodies. **c** Nuclear and cytosolic extracts were imunoprecipitated with anti-Cdk-2 antibody, electrophoresed,

and probed with the indicated antibodies. The immunoprecipitates were also assayed for their capacity to phosphorylate histone H1 in vitro in the presence of ^{32}P ATP. d Time-course experiment on the effect of 20 or 40 μM antiprogestins on the expression of cell cycle-related proteins. e Whole cell extracts from previous experiment were imunoprecipitated with anti-Cdk-2 antibody, electrophoresed, probed with the indicated antibodies, and also assayed for their capacity to phosphorylate histone H1 in vitro. f A similar experiment was performed as in d but instead of WCE, nuclear and cytoplasmic protein extracts were isolated and immunoprecipitated with Cdk-2 antibody upon treatment of the cells with 20 or 40 μM antiprogestins for 24 h



inhibition. There was an increase in the amounts of p21^{cip1} and p27^{kip1} that co-immunoprecipitated with Cdk-2, which was more pronounced in the cytoplasm than in the nucleus, suggesting that the inhibition in Cdk-2 activity is at least in part likely due to an increase in the binding of the inhibitors p21^{cip1} and p27^{kip1}. The activity of Cdk-2 was also remarkably inhibited in both nucleus and cytoplasm of SK-OV-3 cells as shown when treated with RU-38486 (Fig. S3).

The magnitude of the increase in $p21^{cip1}$ and $p27^{kip1}$ and the decline in Cdk-2 levels induced by the antiprogestins was dose-dependent, with the particularity that the increase in $p21^{cip1}$ occurred earlier than that of $p27^{kip1}$ (Fig. 6d). When Cdk-2 was immunoprecipitated, there was a dose-dependent increase in the amounts of $p21^{cip1}$ and $p27^{kip1}$ that co-immunoprecipitated with Cdk-2, which was associated with a parallel decline in the activity of Cdk-2 (Fig. 6e). A dose- and time-dependent inhibition of Cdk-2 activity was also observed in SK-OV-3 cells when exposed to 20 or 40 μ M ORG-31710 (Fig. S4).

In addition to the increased association of Cdk inhibitors p21^{cip1} and p27^{kip1} to Cdk-2, another contributing factor for the reduced Cdk-2 activity in response to antiprogestins appears to be a decline in the nuclear levels of cyclin E (Fig. 6b). To further confirm this assumption we immunoprecipitated Cdk-2 from cytosolic and nuclear fractions of OV2008 cells, which had been exposed for 24 h to 20 or 40 µM antiprogestins. Fig. 6f shows an evident dosedependent decline in Cdk-2 activity in both cellular compartments, together with a major decline in cyclin E nuclear levels, and cyclin E redistribution to the cytoplasm, where lower molecular weight cyclin E fragments were also observed. The association of p21cip1 and p27kip1 to Cdk-2 in the cytoplasmic fraction correlated with the decline in the activity of Cdk-2 in this cellular compartment even in the presence of cyclin E, which however may be undergoing accelerated proteasomal degradation. In the nuclear fraction, however, the correlation among binding of Cdk inhibitors to Cdk-2 and decline in Cdk-2 activity is not apparent, suggesting that it may be the lack of cyclin E rather than the increase in Cdk inhibitors the main cause for the blunted activity of Cdk-2 in the nucleus.

The dose-dependent decline in Cdk-2 activity observed in OV2008 cells (Fig. 6e and f), and SK-OV-3 (Fig. S4), correlated with a dose-dependent growth inhibitory effect elicited by the antiprogestins (Fig. 2 and Fig. S5). Altogether these results suggest that the dose-dependent antiprogestin-mediated inhibition of growth in ovarian cancer cells involves increased nuclear abundance of the Cdk inhibitors p21^{cip1} and p27^{kip2}, decreased Cdk-2 and cyclin E nuclear levels, redistribution of cyclin E to the cytoplasm, and a remarkable decline in the activity of the cell cycle regulatory protein Cdk-2 in both nuclear and cytoplasmic compartments.



Ovarian cancer is known as a silent killer due to its late detection and high mortality. Despite countless efforts to develop early diagnostic tools and new treatment approaches, the 5-year survival for these patients has only improved from 37% to 45% in the past 30 years [32]. To cure this disease efforts are geared to chemoprevention and assessment of risk factors, early detection biomarkers, identification of early disease symptoms, and development of targeted drugs to accompany standard therapy [33]. However, since screening strategies for early diagnosis have so far failed and most patients still die from the disease, new therapeutic options are desperately needed. The results presented in this work clearly show that three different antiprogestin compounds are cytotoxic to ovarian cancer cells displaying two main effects: (i) a cytostatic effect at lower concentrations blocking cell growth at the G1 phase of the cell cycle; and (ii) a lethal effect at higher doses associated with morphological features of apoptosis and fragmentation of the genomic DNA. The overall toxicity of antiprogestins involved a dose-dependent decline in the activity of the cell cycle regulatory protein Cdk-2 (Fig. 7).

Cdk-2 has been shown to be critical in promoting the transition of cells in the cell cycle from G1 to S phase [29]. For instance, cyclin E/Cdk-2 is needed for the stimulation of histone gene transcription [34], which is one of the major events that mark the entry into the S phase. To drive cell cycle progression, Cdk-2 should be free of p21cip1 and p27^{kip1} binding [29], bound to cyclin E, and allocated to the nucleus to phosphorylate cell cycle regulatory proteins [31, 35]. We show that antiprogestins affect the nucleocytoplasmic trafficking of Cdk inhibitors p21^{cip1} and p27^{kip1}, Cdk-2 and its co-factor cyclin E. We demonstrate that antiprogestins increase p21cip1 and p27kip1 abundances in both cytoplasm and nuclear compartments, which correlate with decreased Cdk-2 and cyclin E nuclear levels, increased cytoplasmic cyclin E and a remarkable decline in the activity of Cdk-2 in both subcellular compartments. The magnitude of inhibition of Cdk-2 activity was related to the growth inhibition potency of the compounds with RU-38486 > ORG-31710 > CDB-2914. Supporting our results, a decline in cyclin E-associated kinase activity (presumably Cdk-2) has been previously reported for T-47D breast cancer cells in response to ORG-31710 in the absence of significant changes in cyclin E and Cdk levels, but in the presence of elevated concentrations of p21cip1, suggesting that p21^{cip1} contributes to the reduction in Cdk-2 activity after antiprogestin treatment [9]. In ovarian cancer cells we show that not only the increased association of p21^{cip1} and p27kip1 to Cdk-2 may account for the reduced Cdk-2 activity in the nucleus in response to antiprogestins, but



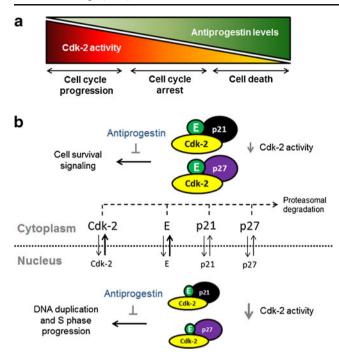


Fig. 7 Proposed model for the cytotoxic effect of antiprogestins in ovarian cancer cells. a In normal growing conditions or in the presence of very low doses of antiprogestins, the activity of Cdk-2 is elevated and the cell cycle progresses. At certain threshold of antiprogestin levels, the activity of Cdk-2 is reduced in association with cytostasis. At higher doses of antiprogestins, Cdk-2 activity is blunted leading to cell death. b Antiprogestins reduce nuclear and cytoplasmic activity of Cdk-2 in association with up-regulation of Cdk-2 inhibitors p21^{cip1} (p21) and p27^{kip1} (p27), down-regulation of Cdk-2 and cyclin E in the nucleus, and cyclin E redistribution to the cytoplasm where it may undergo accelerated proteasome degradation. The inhibition of nuclear activity of Cdk-2 leads to G1 cell cycle arrest due to abrogation of Cdk-2 driven pathways needed for DNA duplication and S phase progression. The inhibition of cytoplasmic Cdk-2 activity, when bypassing certain minimal activity threshold, ultimately leads to cell death likely due to abrogation of a Cdk-2 dependent cell survival pathway

also a reduction in Cdk-2 and cyclin E nuclear levels and redistribution of cyclin E to the cytoplasm, are related variables leading to blunting Cdk-2 nuclear activity needed for G1 to S transition. A recent study using LNCaP prostate cancer cells revealed that targeting Cdk-2 to the nucleus is sufficient to prevent growth inhibition triggered by 1,25 (OH)₂ D3 [36], suggesting that antiprogestin-mediated growth inhibition and growth arrest triggered by metabolites of vitamin D may share common molecular intermediaries.

Because Cdk-2 is frequently up-regulated in ovarian tumors [37], the potent inhibition of Cdk-2 elicited by antiprogestins may be critically important from a translational therapeutics viewpoint. Moreover, because cytoplasmic localization of Cdk inhibitor p27^{kip1} in ovarian cancer patients has been associated with poor prognosis [38], by promoting an increase in p27^{kip1} in the nucleus, antipro-

gestins may be able to rescue the tight inhibitory control of Cdk inhibitors on Cdk-2 activity which is frequently lost in ovarian cancer. One question that deserves an answer is whether p21cip1 and/or p27kip1 that accumulate after antiprogestin exposure are required for antiprogestinmediated Cdk-2 inhibition and/or cell cycle arrest, or whether it is the decline in nuclear cyclin E levels itself sufficient to cause the reduction in nuclear Cdk-2 activity. In support of the latter hypothesis overexpression of cyclin E in LNCaP prostate cancer cells blocked 1, 25 (OH) 2 D3mediated growth inhibition, Cdk-2 relocalization to the cytoplasm, and inhibition of Cdk-2 activity [36], suggesting that a similar mechanism may be taking place in ovarian cancer cells upon antiprogestin treatment. Because in mammalian cells cyclin E is degraded in an ubiquitin- and proteasome-dependent pathway [39, 40], it is possible that by causing cyclin E redistribution to the cytoplasm antiprogestins promote cyclin E proteasomal degradation. This pharmacologic engagement of the proteasome system degrading G1 cyclins such as D1 and E has been previously proposed as a molecular target for cancer therapy [41].

A potential target of antiprogestin action is the ubiquitinproteasome system (UPS). This idea is based on the following facts: (i) to transition from G1 to S phase and to commit to DNA synthesis, the cells must degrade the Cdk-2 inhibitors p27^{kip1} and p21^{cip} via the Skp1-Cullin-Fbox protein/Skp2 (SCFSkp2) E3 ubiquitin ligase complex [42, 43]. This requires the Cdk-2-dependent phosphorylation of p27^{kip1} on Thr187 [42] and p21^{cip1} on Ser130 [43]; (ii) antiprogestins have a dual effect blocking Cdk-2 activity and triggering the accumulation of p21cip1 and p27^{kip1}, and these Cdk-2 inhibitors rely on the UPS for their disappearance to enforce the orderly progression of the cell cycle from G1 to the S phase; (iii) finally, there are remarkable similarities in the behavior of antiprogestins and proteasome inhibitors in inducing p21cip1 and p27kip1 accumulation before triggering caspase-associated lethality [18, 19, 44]. It is therefore possible that antiprogestins induce G1 growth arrest by interfering with the proteasome-mediated degradation of p27^{kip1}/p21^{cip1}, leading to Cdk-2 inhibition. It is also feasible that the sustained levels of p27kip and p21cip1 in response to cytostatic doses of antiprogestins are the consequence of a reduced recognition of the Cdk inhibitors by the UPS. Because ovarian cancer cells function with high activity of the UPS [44], this proteolytic machinery may be degrading Cdk inhibitors at a high rate, causing the reduced basal levels we found in ovarian cancer cells, thus favoring their proliferation. Antiprogestins may mitigate this process.

In addition to regulating cell cycle progression, Cdk-2 is involved in cell survival after DNA damage and in DNA repair pathways [45, 46]. As a survival factor, for instance, Cdk-2 phosphorylates the FOXO1 transcription activator of



pro-apoptotic genes, keeping them in the cytoplasm [45, 47] . If the activity of Cdk-2 is abolished by an antiprogestin, then FOXO1 may not be retained in the cytoplasm, consequently migrating to the nucleus where it promotes the expression of pro-apoptotic genes [45, 47]. The lethality of high concentration antiprogestins associated with features of apoptosis as compared to platinuminduced lethality in the same cell lines in terms of nuclear and DNA fragmentation. However, the molecular mediators of antiprogestin-mediated cell death varied among the steroids. Although cleavage of the caspase-3 substrate PARP was a commonality among RU-38486, ORG-31710 and CDB-2914, the latter also caused an upregulation of PARP which was also previously observed in cultured human uterine leiomyoma cells [24]. In addition, CDB-2914 caused up-regulation of the antiapoptotic proteins XIAP and Bcl-2, yet cell death still ensued but with less effectiveness than that observed after exposure to high concentrations of RU-38486 or ORG-31710 in which both XIAP and Bcl-2 are down-regulated after 3 days of treatment. Thus, the extended up-regulation of XIAP and Bcl-2 upon CDB-2914 treatment but not after RU-38486 or ORG-31710 may account for the reduced cytotoxic potency of CDB-2914. Although with different potencies, high concentrations of antiprogestins were able to lead the cells to crossing a cell death threshold or point of no return in which the pro-apoptotic load of the cell surpasses its anti-apoptotic buffering capacity.

Apoptosis induced by antiprogestins has also been reported in cultured human periovulatory granulosa cells in which RU-38486 and ORG-31710 caused an increase in the activity of executer caspase-3 and fragmentation of the DNA [26]. RU-38486 was also shown to cause apoptosis in breast, cervical, endometrial, and prostate cancer cells in association with activation of caspase-3, down-regulation of Bcl-2 and secretion of TGF β 1 [14, 48, 49]. Using ovarian cancer cell lines, our work expanded to CDB-2914 the previously reported cytotoxicity of RU-38486 and ORG-31710.

It is unknown whether antiprogestins inhibit cell growth involving progesterone receptors (PR), glucocorticoid receptors (GR) or neither one of them. In fact it has been previously suggested a dissociation between the antihormone and anti-proliferative activity of antiprogestins [50]. This is further supported by studies in MDA-MB-231 breast cancer cells lacking both ER and PR, in which RU-38486 retained its antiproliferative activity [14]. On the other hand, our laboratory using SK-OV-3 cells [18] and others using SV-40 transformed ovarian cystadenoma cells [51] have shown that RU-38486 elicits progesterone-like effects in terms of growth inhibition although with greater potency than progesterone, whereas others [52] using HOC-7 ovarian carcinoma cells demonstrated that high concen-

tration (30 μ M) progesterone stimulates p21^{cip1} and p27^{kip1} expression and inhibit Cdk-2 activity mimicking our observations with antiprogestins. Whether progesterone and antiprogestins share similar mechanism of action when acting as anti-proliferative agents, and which are their downstream targets, need to be investigated to identify the genetic background of ovarian cancers required for susceptibility to growth inhibition by antiprogestins.

In summary, the results from the present work provide evidence suggesting that antiprogestins originally utilized for reproductive medicine can be repurposed or repositioned for another modality-of-use, their anti-ovarian cancer property, which involves their capacity to ablate the activity of the cell cycle regulatory protein Cdk2.

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