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Targeting BRCA1/2 deficient ovarian cancer with CNDAC-based drug combinations

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Abstract

Purpose—The mechanism of action of CNDAC (2'-C-cyano-2'-deoxy-1-β-d-arabino-pentofuranosyl-cytosine) is unique among deoxycytidine analogs because upon incorporation into DNA it causes a single strand break which is converted to a double strand break after DNA replication. This lesion requires homologous recombination (HR) for repair. CNDAC, as the parent nucleoside, DFP10917, and as an oral prodrug, sapacitabine, are undergoing clinical trials for hematological malignancies and solid tumors. The purpose of this study is to investigate the potential of CNDAC for the therapy of ovarian cancer (OC).

Methods—Drug sensitivity was evaluated using a clonogenic survival assay. Drug combination effects were quantified by median effect analysis.

Results—OC cells lacking function of the key HR genes, *BRCA1* or *BRCA2*, were more sensitive to CNDAC than corresponding HR proficient cells. The sensitization was associated with greater levels of DNA damage in response to CNDAC at clinically achievable concentrations, manifested as chromosomal aberrations. Three classes of CNDAC-based drug combinations were

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Electronic supplementary material

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Author contributions

Conception and design: X. Liu, W. Plunkett. Development of methodology: X. Liu, Y. Jiang. Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): X. Liu, Y. Jiang, B. Nowak, B. Qiang, N. Cheng, Y. Chen. Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): X. Liu, Y. Jiang, W. Plunkett. Writing, review, and/or revision of the manuscript: X. Liu, Y. Jiang, W. Plunkett. Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): X. Liu, Y. Jiang, B. Nowak. Study supervision: W. Plunkett.

Compliance with ethical standards

Conflict of interest

The authors declare that they have no conflict of interest.

Ethical approval

This article does not contain any studies with human participants or animals performed by any of the authors.

Supplementary material

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Supplementary material 1 (PPT 256 KB)

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Supplementary material 2 (DOC 50 KB)

investigated. *First*, the PARP1 inhibitors, rucaparib and talazoparib, were selectively synergistic with CNDAC in BRCA1/2 deficient OC cells (combination index < 1) at a relatively low concentration range. *Second*, cisplatin and oxaliplatin had additive combination effects with CNDAC (combination index ~ 1). *Finally*, paclitaxel and docetaxel achieved additive cell-killing effects with CNDAC at concentration ranges of the taxanes similar for both BRCA1/2 deficient and proficient OC cells.

Conclusions—This study provides mechanistic rationales for combining CNDAC with PARP inhibitors, platinum compounds and taxanes in ovarian cancer lacking BRCA1/2 function.

Keywords

CNDAC; PARP inhibitor; Platinum compound; Taxane; Clonogenic survival; Drug combination

Introduction

Ovarian cancer (OC) ranks the fifth of all women's cancers with very high mortality (69%) relative to breast cancer (19%) [1]. Nearly 30 out of 100 women with a *BRCA1* or *BRCA2* gene mutation will develop OC by the age of 70 years, compared to fewer than 1 out of 100 women in the general US population [2]. Although a small population (5–13%) of OC patients carry germline mutations in *BRCA1* or *BRCA2*, somatic *BRCA1/2* mutations [3, 4] and hypermethylation in the *BRCA1* promoter region [5, 6, 7, 8] have been revealed in sporadic ovarian cancers. Loss of BRCA1/2 function causes a deficiency in homologous recombination repair of DNA double strand breaks.

Current therapies for OC include the first-line chemotherapeutics, platinum compounds and taxanes. Tumors with loss of BRCA1/2 function are sensitized to platinum compound therapy, whereas the taxanes are largely non-selective. In recent years, PARP inhibitors (PARPis) have emerged as a class of novel therapeutic agents preferentially targeting BRCA1/2-deficient cells due to synthetic lethality [9, 10, 11]. Three PARPis—olaparib, rucaparib and niraparib—have been approved by the FDA to treat patients with advanced ovarian cancer that is associated with deleterious BRCA mutations. However, *BRCA1/2*-mutated cancer cells are able to acquire resistance to cisplatin and PARPis by restoring BRCA1/2 functions through secondary *BRCA1/2* mutations [12, 13, 14, 15, 16]. Therefore, there is a need to develop novel therapeutic agents and combinations to target BRCA1/2-deficient OC and overcome such resistance.

CNDAC is a nucleoside analog with a unique action mechanism of inducing single strand breaks (SSBs) after incorporation into DNA. Incorporated CNDAC is transformed into a de facto chain terminator, CNddC, which is extremely hard to remove [17, 18, 19]. The SSBs are subsequently converted into double strand breaks (DSBs) during a second S-phase. This unique mechanism of action has distinguished CNDAC among deoxycytidine analogs. We demonstrated that CNDAC-induced DNA damage is primarily repaired through the ATM-dependent homologous recombination (HR) pathway, shown by sensitization of cells deficient in either ATM, RAD51D, XRCC3 or BRCA2 [20]. In contrast, cells lacking HR function are not sensitized to cytarabine or gemcitabine [21]. Repair of CNDAC-induced damage is independent of p53 [20]. Shapiro's group reported hypersensitivity of human

colon cancer cells lacking BRCA1 or BRCA2 to CNDAC [22], confirming our findings in hamster lines [20].

Sapacitabine, the orally bioavailable prodrug of CNDAC, is undergoing clinical trials in acute myeloid leukemia and MDS with encouraging outcomes [23, 24]. It is currently in Phase III registration trial for elderly AML patients (NCT01303796). CNDAC (as DFP-10,917) is in Phase I/II trial for AML and ALL (NCT01702155), and clinical responses in treatment for refractory or relapsed AML patients has been reported [25]. In this study, we investigated response of ovarian cancer (OC) cell lines to CNDAC and explored CNDAC-based combination strategies in BRCA1/2 deficient and proficient OC cells. A corollary is that matching the mechanism of action of sapacitabine or CNDAC to the Brca1/2 deficiency in OC may selectively sensitize the tumor.

Materials and methods

Materials

The nucleoside analogue CNDAC was synthesized as described [17]. CO-338 (PF-01367338-BW, rucaparib camsylate salt) was obtained from Clovis Oncology through MTA. Talazoparib (BMN 673), cisplatin, oxaliplatin, paclitaxel, and docetaxel were purchased from Selleck Chemicals (Houston, TX).

Cell lines

The BRCA1-null ovarian carcinoma cell line, UWB1.289 (designated as “– BRCA1”) and its complemented line, UWB1.289 + Brca1 (designated as “+BRCA1”) [26] were purchased from the American Type Culture Collection (ATCC). Both lines were grown in a mixture of two basic media – 50% ATCC-formulated RPMI-1640 and 50% mammary epithelial growth medium (<http://www.atcc.org/Products/All/CRL-2945.aspx#culturemethod> (<http://www.atcc.org/Products/All/CRL-2945.aspx#culturemethod>)) supplemented with 3% fetal bovine serum (FBS). G418 was added in the culture medium for the + BRCA1 line to a final concentration of 200 µg/mL. BRCA2-mutant ovarian adenocarcinoma cell line, PEO1 (designated as “– BRCA2”) and its revertant line, PEO1 C4-2 (designated as “BRCA2 rev.”) [16] were kindly provided by Dr. Tashiyasu Taniguchi (Fred Hutchinson Cancer Research Center, Seattle, WA). Both lines were cultured in DMEM supplemented with 15% FBS and Glutamax. All cells were free of mycoplasma, as certified by Characterized Cell Line Core Facility at M. D. Anderson Cancer Center (MDACC) using the MycoAlert kit from Lonza (Switzerland). Short tandem repeat DNA fingerprinting was used to authenticate human cell lines by the Characterized Cell Line Core. All lines were used in experiments for no more than 15 passages after recovering from frozen seed stocks that were validated prior to freezing.

Immunoblotting

Whole cell lysates were loaded on 4–12% Bis-Tris gradient gels (Bio-Rad) and separated proteins were transferred onto nitrocellulose membranes. Immunoblotting images were captured by Odyssey CLx Imaging System (Li-Cor, Lincoln, NE). The following antibodies were used: BRCA1 polyclonal C-20, Santa Cruz (sc-642); BRCA1 monoclonal MS110,

Calbiochem (OP92); BRCA2 polyclonal, BioVision (#3675); β -actin monoclonal AC-15, Sigma–Aldrich (A5441).

Cytogenetic analysis

Colcemid (100 ng/mL) was added to cell cultures before harvesting. Mitotic spreads were prepared as described [27]. A minimum of 50 metaphase spreads were analyzed for each sample by the T.C. Hsu Molecular Cytogenetics Core at MDACC.

Clonogenic cell survival assay

Exponentially growing cells were seeded to attach in 6-well plates 1 day before cells were exposed to a range of concentrations of drugs for 24 h. After washing into drug-free medium or medium containing a PARPi, cells were incubated for 7–8 days depending on the growth rate of the particular cell line, and colonies (containing ≥ 50 cells) in triplicate wells were stained with crystal violet after fixation. Then colonies were counted either manually under a dissecting microscope or electronically using GelCount (Oxford Optronix, Oxford, UK). IC₅₀ values of single agents were calculated using the GraphPad Prism6 software (GraphPad Software, Inc. San Diego, CA) based on clonogenic survival rates in triplicate.

Median-effect analysis of drug combinations

Cells were treated with two agents combined in fixed molar ratios over a range of concentrations for each agent that achieve 10–90% clonogenic inhibition alone. Clonogenic survival was analyzed by the median-effect method [28, 29] using the CalcuSyn software (Biosoft, Ferguson, MO) as described [30]. The calculated combination index (CI) < 1 , $= 1$ and > 1 indicate being synergistic, additive and antagonist, respectively.

Results

Ovarian cancer cells deficient in BRCA1 and BRCA2 are sensitive to CNDAC

To determine the impact of BRCA1 and BRCA2 on sensitivity to CNDAC, we used clonogenic assays to compare OC cells lacking BRCA1 or BRCA2 with the respective proficient cells. The BRCA1-null OC cell line, UWB1.289, was $>$ threefold more sensitive to CNDAC (IC₅₀ \sim 10 nM, mean of three independent experiments) than the BRCA1-complemented line, UWB1.289 + BRCA1 (mean IC₅₀ \sim 31 nM, Fig. 1a). Similarly, the BRCA2 mutant and defective cell line, PEO1, was $>$ fivefold more sensitive to CNDAC (IC₅₀ \sim 10 nM, mean of two independent experiments) compared to PEO1 C4-2 line (mean IC \sim 59 nM, Fig. 2a), which has restored BRCA2 function by a secondary 50 mutation [16]. Thus, both BRCA1 and BRCA2 are required for OC cells to repair CNDAC-induced DSBs.

To find the underlying cause for the sensitization of BRCA1/2 deficient OC cells to CNDAC, we measured DNA damage at the chromosomal level (Fig. 1 b). Before CNDAC treatment, 52% (26/50) of BRCA1-null UWB1.289 cells had aberrant metaphases, including 10% (5/50) metaphases which had too many aberrations to allow accurate counting (unscorable, Fig. 1 c). Among the scorable metaphases—the sum of metaphases without aberration and with scorable aberrations (90%, 45/50), each had average 0.24 chromosomal break and 0.29 fusion per metaphase (Fig. 1 d). The ratio of aberrant metaphases in BRCA1-

complemented cells was 32% (16/50), including 4% (2/50) with unscorable aberrations (Fig. 1 c). The scorable metaphases (96%, 48/50) each had average 0.17 break and 0.08 fusion (Fig. 1 d). Thus, lack of BRCA1 led to a greater basal level of chromosomal aberrations. When exposed to 15 nM CNDAC for 27 h (approx. one cell cycle), more BRCA1-null cells had aberrant mitosis (67%, 35/52), including 29% (15/52) metaphases with severe chromosomal damage (unscorable, Fig. 1 c). Among the scorable metaphases (71%, 37/52), the average break number was 0.54 and fusion number 0.3 per metaphase (Fig. 1 d). The ratios of aberrant metaphase (94%, 48/51) and unscorable aberrant metaphase (65%, 33/51) further increased dramatically with cells exposed to CNDAC after a second cell cycle at 54 h (Fig. 1 c), demonstrating that more DSBs are formed after a second S-phase. The average numbers of breaks and fusions were 1.56 and 0.83, respectively, among those scorable metaphases (35%, 18/51, Fig. 1 b, d). In contrast, when BRCA1-repleted cells were treated with 15 nM CNDAC, the ratio of aberrant mitosis increased to 42% (21/50) at 27 h and 48% (24/50) at 54 h. The ratio of unscorable aberrant mitosis remained constant (2–4%, Fig. 1 c). The average break and fusion numbers in those scorable metaphases did increase, but far less than the BRCA1-null cells (Fig. 1 d). These results demonstrate a clear relationship between the deficiency in BRCA1 and the greater level of chromosomal damage after CNDAC, indicating that BRCA1 is essential for the repair of CNDAC-induced DSBs.

The basal level of chromosomal aberrations in BRCA2 deficient PEO1 cells (67% with aberrations) was much greater than that in BRCA2 restored PEO1 C4–2 cells (14% with aberrations, Fig. 2c). This is similar to the BRCA1 deficient situation, suggesting innate genetic instability in the absence of BRCA1/2. Remarkably, after a 72-hour incubation with 25 nM CNDAC, all metaphases in BRCA2 mutant cells presented severe chromosomal damage which could not warrant accurate counting (100% unscorable, Fig. 2b–d), whereas only 30% (15/50) metaphases in BRCA2 reverted cells had aberrations and all of them were scorable (Fig. 2c, d). Therefore, BRCA2 is also required for cells to repair DNA damage caused by CNDAC. These cytogenetic studies provide direct evidence for the sensitization of OC cells to CNDAC in the scenario of BRCA1/2 deficiency.

To evaluate the possibility that distinct levels of chromosomal lesion after CNDAC exposure is attributed to differential accumulation of the nucleoside analog in cells, we measured cellular levels of the triphosphate of CNDAC, CNDACTP by anion-exchange HPLC. Both BRCA2-mutant and -revertant cell lines showed a slight dependency of CNDACTP accumulation on drug concentration (1 and 3 μ M) and incubation time (2 and 4.5 h). However, there was no significant difference between the mutant and restored cells comparing CNDACTP concentrations detected from whole cell extracts under all conditions (Fig. S1).

Synergistic interaction between CNDAC and PARP inhibitors is selective in BRCA1 and BRCA2 defective ovarian cancer cells

The combination of CNDAC with a PARPi is synergistic in rodent cells with defective homologous recombination (HR) function, such as those lacking XRCC3, RAD51C or RAD51D [30]. We hypothesized that CNDAC potentiates the synthetic lethal condition in HR deficient cells exposed to a PARPi, because CNDAC-induced DNA damage largely

depends on the HR pathways for repair [20], whereas PARPi affects both the base excision repair (BER) and HR pathways [31, 32]. To test this combination strategy in human ovarian cancer cells. BRCA1-null OC cells were ~ 80 fold more sensitive to rucaparib ($IC_{50} \sim 13$ nM) than were BRCA1 complemented cells ($IC_{50} \sim 1040$ nM, Fig. S2A). BRCA2-revertant OC cells were 18-fold more resistant to rucaparib ($IC_{50} \sim 1800$ nM) than were BRCA2-mutated cells ($IC_{50} \sim 100$ nM, Fig. S2B). These results are consistent with the well-accepted reports of synthetic lethal killing of BRCA-defective cells by PARPis [9, 33].

Combination of rucaparib and CNDAC at a ratio of 10:1 achieved $CI < 1$ in UWB1.289 cells, indicating synergy in a situation of BRCA1 deficiency (Fig. 3 a). The rucaparib-CNDAC combination in BRCA1-complemented cells (ratio 50:1) also achieved $CI < 1$ (Fig. 3 b), but in much greater concentration ranges of both agents compared to that used for the deficient cells. When rucaparib and CNDAC were combined at a ratio of 10:1 in BRCA2 deficient PEO1 cells, the synergy was more obvious ($CI < 1$, Fig. 3 c). Similarly, this combination at a ratio of 50:1 (rucaparib : CNDAC) was synergistic ($CI < 1$) at greater concentrations of each agent in BRCA2-revertant cells relative to that in the mutated cells (Fig. 3 d). We also tested another potent PARPi, talazoparib in the BRCA2 paired cell lines. Combination of talazoparib with CNDAC (ratio 1:12) resulted in $CI < 1$ in PEO1 cells (four experiments, same below), showing synergistic interaction between the two agents when lacking BRCA2 (Fig. 3 e). Similar to the rucaparib-CNDAC combination, the talazoparib-CNDAC combination achieved $CI < 1$ in BRCA2-restored cells at much higher concentrations of both agents than that in the mutated cells (Fig. 3 f). Thus, combinations of CNDAC with rucaparib or talazoparib clearly exerted a synergistic interaction between PARPi and CNDAC. For BRCA1/2-deficient cells, the concentrations of both agents needed were too low to kill the proficient cells, demonstrating selectivity for the BRCA1/2 mutants. However, this selectivity of synergistic cell killing was lost at greater concentrations in BRCA1/2-repleted or -restored cells.

Combination of CNDAC with platinum compounds

Platinum agents are frequently used in OC chemotherapy regimen. They act by forming intra- and inter-strand crosslinks in DNA. We have reported additive combination effect of cisplatin or oxaliplatin with CNDAC in both HR-proficient and deficient cells [30]. In the context of OC, we sought to investigate the combination effect of CNDAC with cisplatin and oxaliplatin in BRCA1/2 paired OC cell models. In BRCA1-deficient UWB1.289 cells, combination of cisplatin and CNDAC at a ratio of 1.8:1 achieved $CI \leq 1$, while CI for the oxaliplatin-CNDAC combination (6:1) was < 1 (Fig. 4a). Similarly in BRCA1-complemented cells, the cisplatin-CNDAC combination (1.5:1) was additive ($CI \sim 1$), whereas that of oxaliplatin - CNDAC (5:1) was slightly synergistic ($CI < 1$, Fig. 4 b). Since the IC_{50} values of cisplatin and oxaliplatin (single agent) were approximately twice as much in BRCA1-repleted as in BRCA1-defective cells, the combination effect was specific for the defective cells at lower concentrations.

We also compared these combinations in BRCA2 mutated and revertant OC cells. The combination of cisplatin-CNDAC (25:1) in BRCA2 deficient cells was additive ($CI \leq 1$), while that of oxaliplatin-CNDAC (75:1) was synergistic ($CI < 1$, Fig. 4 c). The trend

reversed in BRCA2 restored cells though. The cisplatin–CNDAC combination (12:1) was synergistic ($CI < 1$), whereas that of oxaliplatin–CNDAC (22:1) was additive ($CI \leq 1$, Fig. 4 d). Since the IC values of cisplatin and oxaliplatin (single agent) were 50 approximately 4–5 fold greater in BRCA2 restored cells than that in BRCA2 defective cells, the combination effect with CNDAC was specific for defective cells at lower concentrations.

Taxane agents and CNDAC have additive effect in cells lacking HR function

Taxanes are another class of agents commonly used in chemotherapy for OC. We have shown additive effect of CNDAC in combination with either of the two well-established tubulin-directed agents, paclitaxel or docetaxel, in HR defective cells and proficient cells as well [30]. Here BRCA1 or BRCA2 deficient OC cells were not sensitized to paclitaxel or docetaxel (Fig. 5 a–d, single agent curves), probably because the taxanes specifically impact on mitotic spindle by suppressing the depolymerization of microtubules. Combination of paclitaxel and CNDAC had additive effect ($CI \sim 1$) in both BRCA1-null cells (ratio 2:25, Fig. 5 a left panel) and BRCA1-complemented cells (ratio 1:25, Fig. 5 b left panel). Similarly, the docetaxel–CNDAC combination was additive in BRCA1 paired line (ratios 1:40 and 3:250, respectively, Fig. 5 a, b right panels). In the BRCA2 paired cell lines, the combinations of paclitaxel–CNDAC (ratio 3:20 for BRCA2 mutant, 2:75 for BRCA2 revertant) and docetaxel–CNDAC (ratio 3:50 for BRCA2 mutant, 7:500 for BRCA2 revertant) had additive effect with $CI \sim 1$ (Fig. 5 c, d). This effect is selective in BRCA1/2-deficient cells at lower concentrations of CNDAC that allow survival of proficient cells.

Discussion

Clinical trials on CNDAC and sapacitabine have been conducted largely in hematological malignancies. The clinical trials of sapacitabine for the treatment of solid tumors have not passed Phase II yet [34, 35]. Our prior and current studies have indicated activity of CNDAC in solid tumor cell lines. CNDAC has preferential killing effect in OC cells lacking BRCA1 or BRCA2. These deficient cells also manifested more remarkable chromosomal damage before and after exposure to CNDAC, compared with their wild-type counterparts. The BRCA1/2-associated genetic instability is the cause of the sensitization of BRCA1/2 deficient OC cells. Reciprocally, since BRCA1/2 plays a critical role in the repair of CNDAC-induced DNA damage, the absence of BRCA1/2 results in greater levels of chromosomal aberrations in cells treated with CNDAC.

We have demonstrated synergistic interaction between CNDAC and PARPis, as well as additive effects of CNDAC with platinum compounds and taxane agents, in hamster cell model systems [30]. In the current study, we applied these combination strategies in human OC cells and found the effect of combinations largely consistent with that in hamster cells.

Olaparib is the first PARPi which received accelerated approval from FDA in 2014 for ovarian cancer patients with deleterious germline and/or somatic mutations in BRCA1/2 [36]. A priority review has been granted to the new drug application for olaparib as a maintenance therapy in repleted patients with platinum-sensitive OC [37]. Another PARPi, rucaparib was approved for OC with deleterious BRCA1/2 mutations in 2016 [38]. A third PARPi, niraparib (MK-4827) was recently approved by FDA for the maintenance treatment

of recurrent OC, which does not require testing for BRCA mutation [39]. The synergy between a PARPi and CNDAC in BRCA1/2 deficient cells could be attributed to synthetic lethality in which both the HR and base excision repair (BER) pathway are not functional. PARPi blocks BER by competing the NAD⁺ binding site in PARP, while repair of CNDAC-induced damage requires HR function. We have demonstrated that BER has little influence on cellular response to CNDAC [40]. In other words, inhibition of BER pathway does not enhance the cytotoxicity of CNDAC. Therefore, inhibition of PARP catalytic activity is unlikely the mechanism for synergistic combination of PARPi and CNDAC in BRCA1/2 proficient cells. Talazoparib potentiates the cell-killing effect of CNDAC more efficiently than rucaparib in both HR defective and proficient cells (Fig. 3). Since talazoparib has greater ability in trapping PARP compared with rucaparib and olaparib [41], PARP trapping [42, 43, 44] might outweigh other synergy mechanisms in cells with normal HR function, which is exhausted by significant amount of DNA DSBs after exposure to high concentration of CNDAC (and PARPi as well).

As with other targeted therapies, resistance has arisen in the clinical use of PARP inhibitors. There are multiple mechanisms for the resistance to PARP inhibitors, including restoration of BRCA1/2 function through secondary mutations, upregulation of NHEJ capacity, reduction in PARP1 activity or level, and decreased intracellular concentration of PARPi due to efflux by *p*-glycoproteins [45, 46]. While complete remission was achieved for years in some high-grade serous OC (HGSOC) patients with somatic biallelic deletion of BRCA1/2 after olaparib therap, resistance also arose in other patients with one copy of intact (WT) BRCA1/2 due to copy number gain and/or upregulation of the remaining functional allele [47]. Therefore, PARPi resistance poses a substantial challenge in OC targeted therapy. We found that at concentrations which allow survival of the majority of wild type cells, the synergistic killing effect of the PARPi-CNDAC combination is selective in BRCA1/2 deficient OC cells. This directs to testing of the combination in animal studies such as using patient-derived xenograft models of OC with BRCA1/2 mutations. In the circumstances without BRCAness, selectivity is lost because higher concentrations are needed for both agents to exert synergy to a similar extent as in the defective cells. However, the concentrations of each agent used in combination are still less compared with those used as a single agent. These findings suggest that combining a PARPi and CNDAC might benefit OC patients broadly, with or without functional BRCA1/2, in overcoming or delaying potential resistance to PARP inhibitors.

Chemotherapy based on platinum and/or taxane compounds plays a crucial role as first-line treatment for OC. Our study showed largely additive effects in the combination of CNDAC with either of the two platinum compounds, cisplatin and oxaliplatin, selective in BRCA1/2 deficient OC cells at relatively lower concentrations. Thus, the cisplatin-CNDAC and oxaliplatin-CNDAC combinations are useful to attenuate the cytotoxicity of platinum. The current study also showed an additive combination effect of CNDAC with paclitaxel and docetaxel used at similar concentrations in BRCA1/2 proficient and deficient OC cells. Such combinations may also be useful for reducing toxicity of the respective taxane agents.

Similar to the case of PARP inhibitors, emergence of resistance to platinum compounds has become a major hurdle in related therapeutic regimens. Progress has been made in nearly

50% of carboplatin-resistant HGSOc by co-therapy with carboplatin and birinapant, a potent cIAP inhibitor, which re-sensitizes cells to carboplatin [48]. As a cisplatin derivative, carboplatin shares a similar mechanism of action. Cisplatin resistant tumors are often not cross-resistant to oxaliplatin, suggesting a distinct action mechanism [49]. Besides cross-resistance among PARP inhibitors, PARPi-resistant cells are resistant to cross-linking agents such as cisplatin and carboplatin, but not to microtubule poisons such as taxanes [50]. Thus, the combination of CNDAC with taxanes could provide a novel treatment modality to deal with the above cross-resistance in OC therapy.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Abbreviations

BER	Base excision repair
CNDAC	2'-C-cyano-2'-deoxy-1-β- d- <i>arabino</i> -pentofuranosyl-cytosine
CNDACTP	CNDAC triphosphate
CNddC	2'-C-cyano-2',3'-didehydro-2',3'-dideoxycytidine
CI	Combination index
DSB	Double strand break
FBS	Fetal bovine serum
HGSOc	High-grade serous ovarian cancer
HR	Homologous recombination
OC	Ovarian cancer
NHEJ	Non-homologous end joining
PARPi	PARP inhibitor
PCC	Premature chromosome condensation
SSB	Single strand break

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%20Ohnuma&author=K.%20Tugawa&author=T.%20Ohta&author=M.%20Roche-
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Personalised recommendations

1. Successful use of Bruton's kinase inhibitor, ibrutinib, to control paraneoplastic pemphigus in a patient with paraneoplastic autoimmune multiorgan syndrome and Lee, Andrew... Shumack, Stephen *Australasian Journal of Dermatology* (2017)
2. Ibrutinib as a Bruton Kinase Inhibitor in the Management of Chronic Lymphocytic Leukemia: A New Agent With Great Promise Foluso, Ogunleye... Jaiyesimi, Ishmael *Clinical Lymphoma Myeloma and Leukemia* (2015)
3. Bruton's tyrosine kinase inhibitors: first and second generation agents for patients with Chronic Lymphocytic Leukemia (CLL)

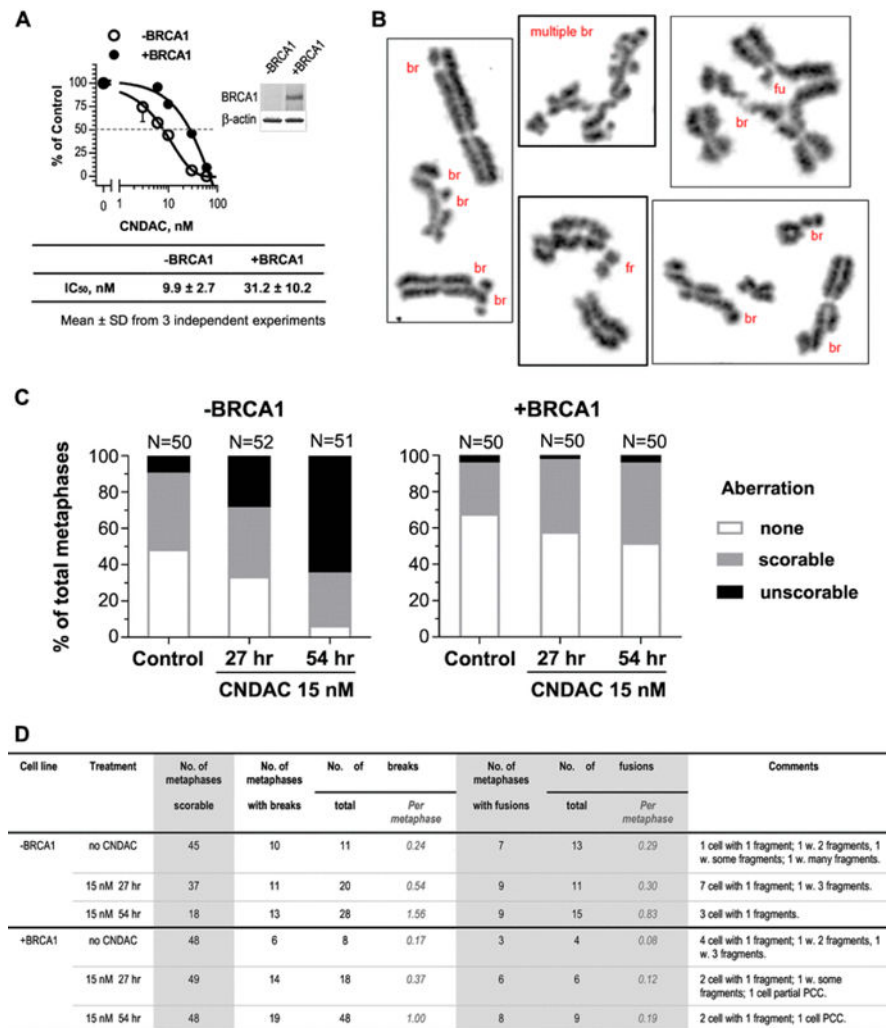


Fig. 1. BRCA1 deficiency confers sensitivity to CNDAC in ovarian cancer cells, which is associated with greater level of chromosomal aberrations. **a** Clonogenic survival of UWB1.289 (BRCA1 deficient, - BRCA1) and UWB1.289 + BRCA1 (complemented, + BRCA1) cell lines in response to CNDAC (3–60 nM, 24-hour exposure). Percentage of cell survival in both cell lines is plotted as a function of drug concentration, and a representative plot is shown. IC₅₀ values of CNDAC (Mean ± SD from three independent experiments) are listed. The insert shows the immunoblot of BRCA1 and β -actin (loading control) proteins. **b–d** CNDAC induces chromosomal aberrations in cells deficient and proficient in Brca1. **b** Representative images of chromosome abnormalities in - BRCA1 cells with many aberrations are shown. *br* Break, *fu* fusion, *fr* fragment. **c** Percentage of metaphases with chromosomal structural aberrations (including breaks, fusions and fragments) detected in - BRCA1 and + BRCA1 cell lines in response to 15 nM CNDAC for 27 and 54 h. **d** More details of chromosomal aberrations detected in metaphases that are scorable (with or without aberration) as in **c**. The number of scorable metaphases is calculated by subtracting the number of unscorable metaphases from the examined total. PCC stands for premature chromosome condensation

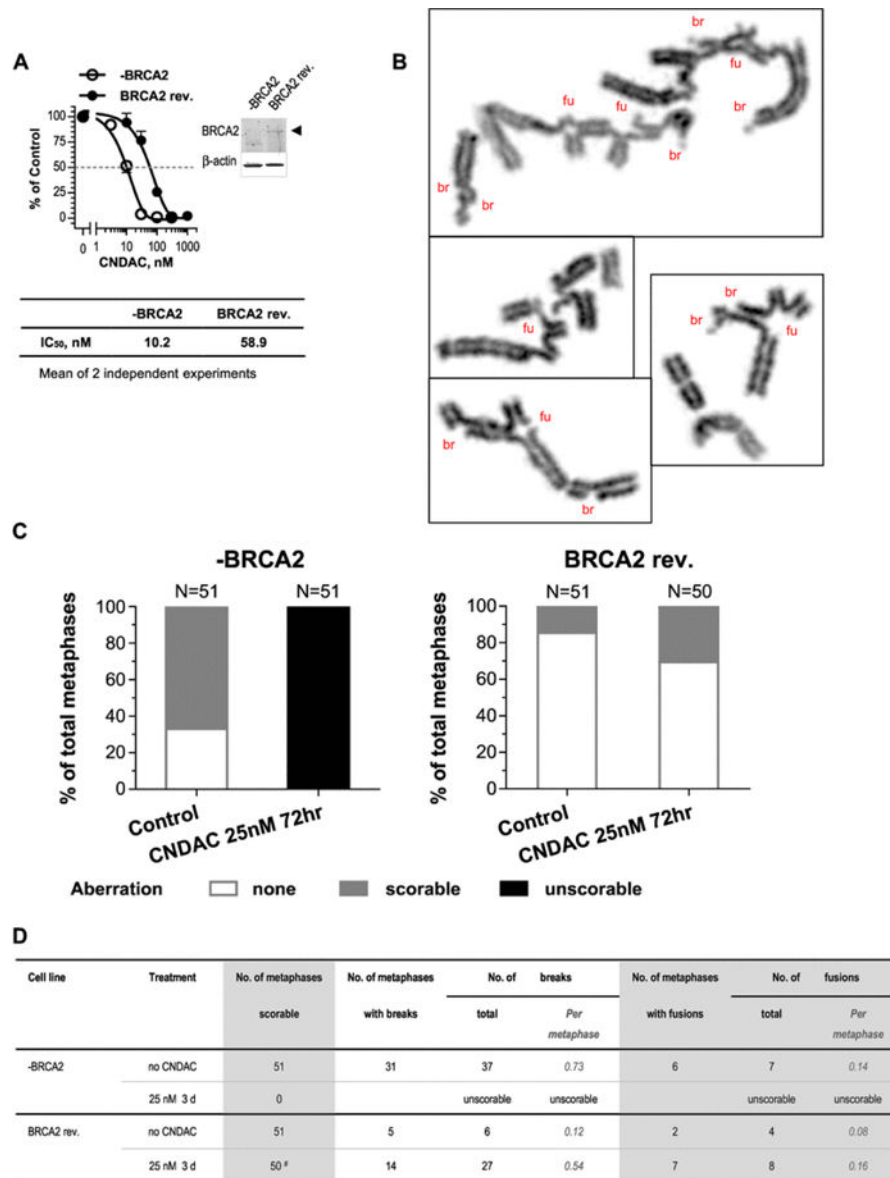


Fig. 2. BRCA2 deficiency also confers sensitivity to CNDAC in ovarian cancer cells, which is associated with greater level of chromosomal aberrations. **a** Clonogenic survival of PEO1 (BRCA2 deficient, -BRCA2) and PEO1 C4-2 (BRCA2 revertant, BRCA2 rev.) cell lines after a 24-hour incubation with CNDAC (3–300 nM and 10–1000 nM, respectively). Percentage of cell survival in both cell lines is plotted as a function of drug concentration, and a representative plot is shown. IC₅₀ values of CNDAC (mean of two independent experiments) are presented. The insert shows the immunoblot of BRCA2 and β -actin proteins. **b–d** CNDAC induces chromosomal aberrations in Brca2 deficient and repleted cells. **b** Representative images of chromosome abnormalities in - BRCA2 cells with many aberrations are shown. *br* Break, *fu* fusion. **c** Percentage of metaphases with chromosomal aberrations determined in - BRCA2 and BRCA2 rev. cell lines in response to 25 nM CNDAC for 72 h. **d** More details of chromosomal aberrations detected in metaphases that

are scorable. The number of scorable metaphases is calculated by subtracting the number of unscorable metaphases from the examined total. # one metaphase with too many aberrations excluded. Those numbers are not added in the table. That particular metaphase has over 25 breaks and about 11 fusions

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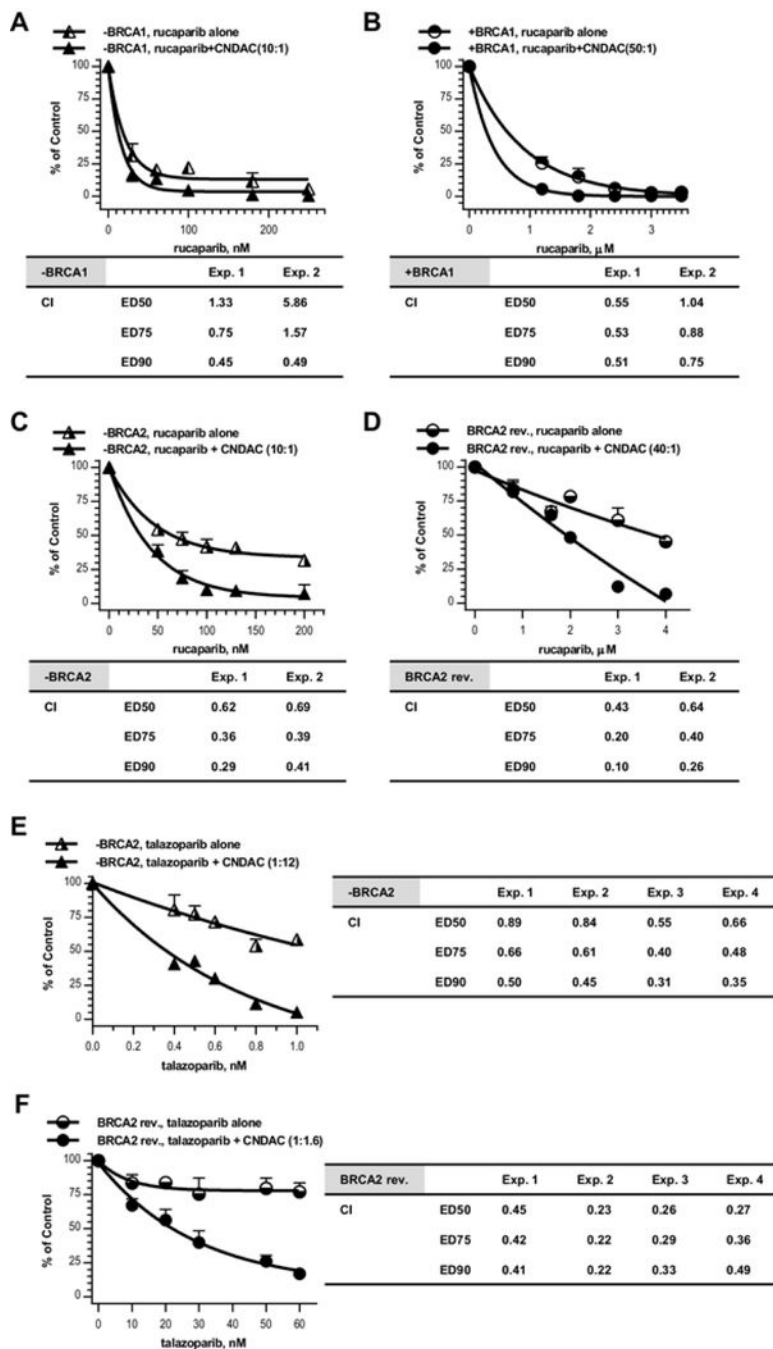
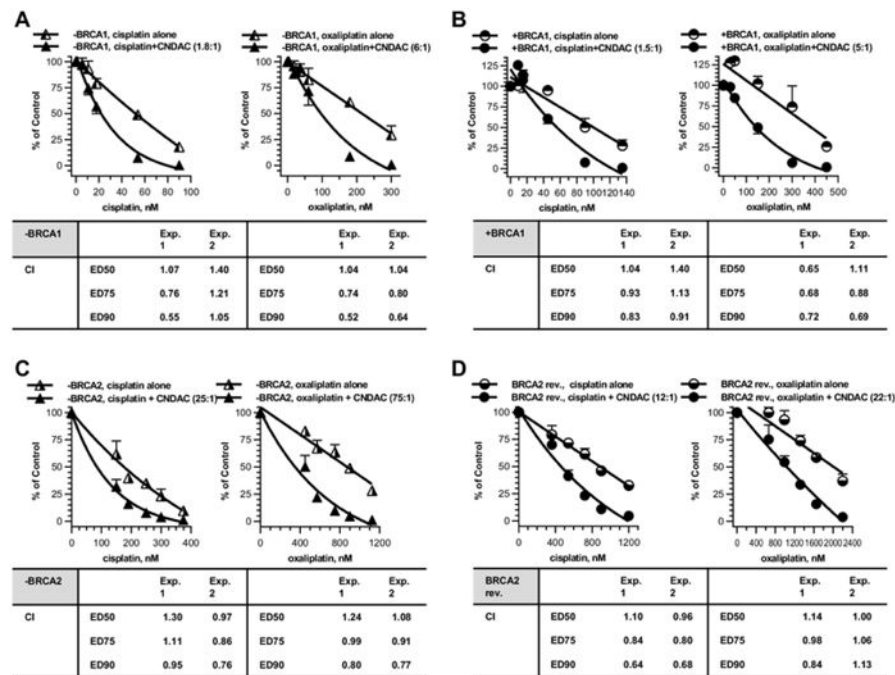


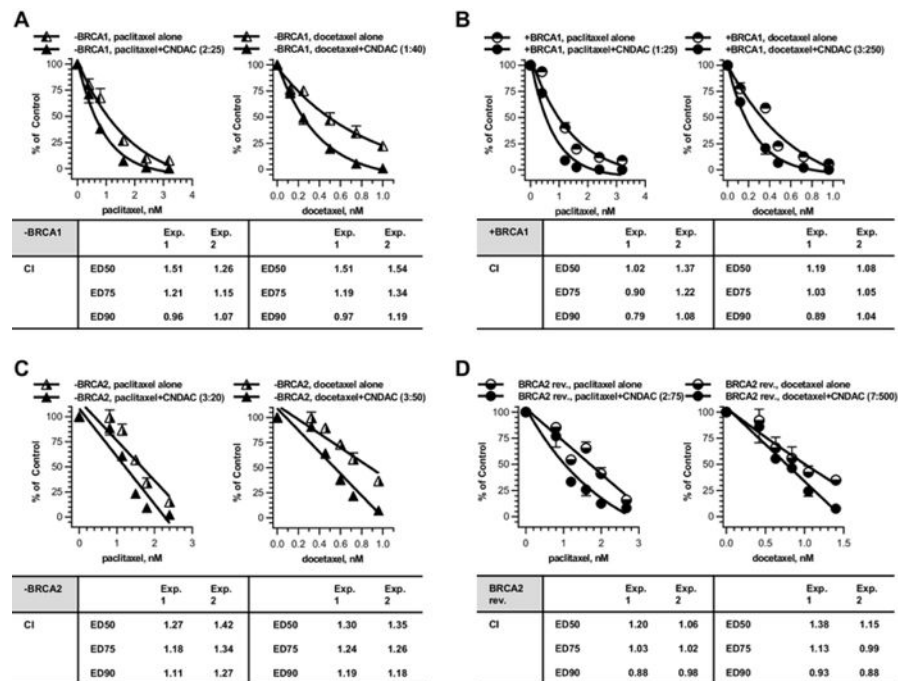
Fig. 3.

The PARP inhibitors rucaparib and talazoparib synergize with CNDAC in both BRCA1/2 proficient and deficient ovarian cancer cell lines. **a** BRCA1 deficient (– BRCA1) cells were exposed to rucaparib (30–250 nmol/L) 1 h before addition of CNDAC (3–25 nmol/L). **b** BRCA1 complemented (+ BRCA1) cells were exposed to rucaparib (1.2–3.5 μmol/L in Exp. 1, or 1–3 μmol/L in Exp. 2) 1 h before addition of CNDAC (24–70 nmol/L in Exp.1, or 25–75 nmol/L in Exp.2). **c** BRCA2 deficient (– BRCA2) cells were exposed to rucaparib (5–200 nmol/L) 1 h before addition of CNDAC (5–20 nmol/L). **d** BRCA2 revertant (BRCA2 rev.)

cells were exposed to rucaparib (0.8–4 $\mu\text{mol/L}$) 1 h before addition of CNDAC (20–100 nmol/L). **a–d** 24 h after CNDAC addition, drugs were washed out and cells were incubated in medium with rucaparib at the corresponding concentrations until colonies were fixed. Top, representative of 2 independent experiments, comparison of clonogenicity of – BRCA1 cells after treatment with rucaparib alone versus rucaparib–CNDAC combination at a fixed ratio as indicated. Bottom, CIs from median-effect analysis of the 2 experiments are presented in the accompanying inserts. **e** BRCA2 deficient (– BRCA2) cells were exposed to talazoparib (0.4–1 nmol/L in Exp. 1 and 2, or 0.4–1.6 nmol/L in Exp. 3 and 4) 1 h before addition of CNDAC (4.8–12 nmol/L in Exp. 1 and 2, or 3.6–14.4 nmol/L in Exp. 3 and 4). **f** BRCA2 revertant (BRCA2 rev.) cells were exposed to talazoparib (10–60 nmol/L in Exp. 1 and 2, or 15–100 nmol/L in Exp. 3 and 4) 1 h before addition of CNDAC (16–96 nmol/L in Exp. 1 and 2, or 15–100 nmol/L in Exp. 3 and 4). **e–f** Drugs were washed out 24 h after CNDAC co-incubation, and cells were incubated in medium with talazoparib at the corresponding concentrations until colonies were fixed. Top, representative of four independent experiments, comparison of clonogenicity of BRCA2 deficient cells after treatment with talazoparib alone versus talazoparib–CNDAC combination at a fixed ratio as indicated. Bottom, CIs from median-effect analysis of the four experiments are presented in the accompanying inserts

**Fig. 4.**

Combinations of platinum compounds cisplatin and oxaliplatin with CNDAC in BRCA1/2 proficient and deficient cells. **a** BRCA1 deficient OC cells (– BRCA1) were treated concomitantly with cisplatin (5.4– 90 nmol/L) or oxaliplatin (18–300 nmol/L) and then CNDAC (3– 50 nmol/L), and incubated for 24 h before washout. **b** BRCA1 complemented OC cells (+ BRCA1) were exposed to CNDAC (6– 90 nmol/L) subsequent to cisplatin (9– 135 nmol/L) or oxaliplatin (30– 450 nmol/L) for 24 h as in **a**. **c** BRCA2 deficient OC cells (– BRCA2) were treated concomitantly with cisplatin (150–375 nmol/L) or oxaliplatin (450–1125 nmol/L) and then CNDAC (6–15 nmol/L), and incubated for 24 h before washout. **d** BRCA2 revertant OC cells (BRCA2 rev.) were exposed to CNDAC (30–100 nmol/L) subsequent to cisplatin (360– 1200 nmol/L) or oxaliplatin (660–2200 nmol/L) for 24 h as in **C**. **a–d** Cells were incubated in drug-free medium until colonies were fixed and quantitated. Top, representative of 2 independent experiments, clonogenicity was compared for treatment with cisplatin alone versus cisplatin–CNDAC combination at a fixed ratio (left graph) or oxaliplatin versus oxaliplatin–CNDAC combination at a fixed ratio (right graph). Bottom, CIs from median-effect analysis of the two experiments are presented in the accompanying inserts

**Fig. 5.**

Combinations of taxane compounds paclitaxel and docetaxel with CNDAC in BRCA1/2 proficient and deficient cells. **a** BRCA1 deficient OC cells (-BRCA1) were treated concomitantly with paclitaxel (0.4–3.2 nmol/L) or docetaxel (0.125–1 nmol/L) and CNDAC (5–40 nmol/L), and incubated for 24 h before washout. **b** BRCA1 complemented OC cells (+BRCA1) were exposed concomitantly to paclitaxel (0.4–3.2 nmol/L) or docetaxel (0.12–0.96 nmol/L) and CNDAC (10–80 nmol/L) for 24 h as in **a**. **c** BRCA2 deficient OC cells (-BRCA2) were treated concomitantly with paclitaxel (0.81–2.4 nmol/L) or docetaxel (0.324–0.96 nmol/L) and CNDAC (5.4–16 nmol/L), and incubated for 24 h before washout. **d** BRCA2 revertant OC cells (BRCA2 rev.) were exposed to CNDAC (30–100 nmol/L) subsequent to paclitaxel (0.8–2.67 nmol/L) or docetaxel (0.42–1.4 nmol/L) for 24 h as in **c**. **a–d** Cells were incubated in drug-free medium until colonies were fixed and quantitated. Top, representative of 2 independent experiments, clonogenicity was compared for treatment with paclitaxel alone versus paclitaxel–CNDAC combination at a fixed ratio (left) or docetaxel versus docetaxel–CNDAC combination at a fixed ratio (right). Bottom, CIs from median-effect analysis of the two experiments are presented in the accompanying inserts