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## **Research Article**

## Carboplatin and taxol resistance develops more rapidly in functional BRCA1 compared to dysfunctional BRCA1 <sup>(3)</sup> ovarian cancer cells

## **Q1** Steven Busschots<sup>*a,b*</sup>, Sharon O'Toole<sup>*b*</sup>, John J. O'Leary<sup>*a,b*</sup>, Britta Stordal<sup>*a,c,\**</sup>

<sup>a</sup>Department of Histopathology, Trinity College Dublin, Central Pathology Laboratory, St James's Hospital, Dublin 8, Ireland <sup>b</sup>Department of Obstetrics and Gynaecology, Trinity College Dublin, Trinity Centre, St James's Hospital, Dublin 8, Ireland <sup>c</sup>Department of Natural Sciences, Hendon Campus, Middlesex University, London NW4 4BT, United Kingdom

#### ARTICLE INFORMATION

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#### ABSTRACT

A major risk factor for ovarian cancer is germline mutations of BRCA1/2. It has been found that (80%) of cellular models with acquired platinum or taxane resistance display an inverse resistance relationship, that is collateral sensitivity to the other agent. We used a clinically relevant comparative selection strategy to develop novel chemoresistant cell lines which aim to investigate the mechanisms of resistance that arise from different exposures of carboplatin and taxol on cells having BRCA1 function (UPN251) or dysfunction (OVCAR8). Resistance to carboplatin and taxol developed quicker and more stably in UPN251 (BRCA1-wildtype) compared to OVCAR8 (BRCA1-methylated). Alternating carboplatin and taxol treatment delayed but did not prevent resistance development when compared to single-agent administration. Interestingly, the sequence of drug exposure influenced the resistance mechanism produced. UPN251-6CALT (carboplatin first) and UPN251-6TALT (taxol first) have different profiles of cross resistance. UPN251-6CALT displays significant resistance to  $CuSO_4$  (2.3-fold, p=0.004) while UPN251-6TALT shows significant sensitivity to oxaliplatin (0.6-fold, p=0.01). P-glycoprotein is the main mechanism of taxol resistance found in the UPN251 taxane-resistant sublines. UPN251 cells increase cellular glutathione levels (3.0-fold, p=0.02) in response to carboplatin treatment. However, increased glutathione is not maintained in the carboplatin-resistant sublines. UPN251-7C and UPN251-6CALT are low-level resistant to CuSO<sub>4</sub> suggesting alterations in copper metabolism. However, none of the UPN251 sublines have alterations in the protein expression of ATP7A or CTR1. The protein expression of BRCA1 and MRP2 is unchanged in the UPN251 sublines. The UPN251 sublines remain sensitive to parp inhibitors veliparib and CEP8983 suggesting that these agents are candidates for the treatment of platinum/taxane resistant ovarian cancer patients.

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\*Corresponding author.

E-mail addresses: busschos@tcd.ie (S. Busschots), shotoole@tcd.ie (S. O'Toole), olearyjj@tcd.ie (J.J. O'Leary), b.stordal@mdx.ac.uk (B. Stordal).

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## Introduction

103104Ovarian cancer is the 5th most prevalent cancer amongst European105women and is the leading cause of death from a gynaecological106malignancy. The majority of patients present with late stage disease107and have an approximately 30% 5-year survival rate [4]. The108standard treatment is surgical debulking followed by intravenous109platinum-taxane combination chemotherapy ([41,53]). This treat-110ment often fails and patients relapse with chemoresistant disease.

A strong family history of ovarian or breast cancer, which is often linked to BRCA1/2 germline mutations, is one of the greater risk factors associated with the disease. Deleterious germline mutations are found in 8.6-13.7% of ovarian cancer patients ([43,49,51]). These mutations cause BRCA1 dysfunction leading to reduced expression of functional BRCA1. A recent study, examining both somatic and germline mutations in ovarian cancer, has revealed that incidence for BRCA1/2 mutations might be even higher at 18.3% [20]. A woman with a BRCA1 mutation has a 39-46% chance of developing ovarian cancer [19]. BRCA1 function has not been fully elucidated but it has been shown to have roles in a number of cellular processes including DNA damage repair, apoptosis, cell cycle regulation, transcriptional control and ubiquitination ([28,38]). 

A systematic review of the literature by [58] revealed that the majority (80%) of cellular models with acquired platinum or taxane resistance displayed an inverse resistance relationship, that is collateral sensitivity to the other agent. A subsequent systematic review by [56] revealed that BRCA1 was the mostly likely genetic player in this relationship. Cells with BRCA1 defects have reduced efficiency in repairing DNA adducts and show increased apoptosis in response to platinums conferring sensitiv-ity [12,66]. The response to taxanes, in BRCA1 deficient cells is reduced apoptosis conferring resistance [29]. The opposite is true for cells with functional BRCA1 [47,61]. 

In this study, chemoresistant ovarian cancer cell lines were developed from established ovarian cancer cell lines using a novel comparative selection strategy. UPN251, which has functional BRCA1 (BRCA1 wild-type due to reversion mutation [59]) and OVCAR8, which has dysfunctional BRCA1 lending to reduced BRCA1 expres-sion (due to BRCA1 methylation) were used in order to investigate the development of chemoresistance in relation to BRCA1 status. This study highlights the effects of BRCA1 function and dysfunction on the development of resistance. In particular it focuses on its effects on the inverse resistance relationship between platinums and taxanes and its effect on alternating platinum and taxane doses. 

## Methods

## Cell culture

The human ovarian cancer cell lines UPN251 and OVCAR8 were sourced from the MD Anderson Cancer Centre. Cells were grown in RPMI 1640 medium (Sigma R8758-500ML) supplemented with 10% foetal calf serum (Lonza DE14-801F), free of antibiotics. All cells were maintained in a humidified atmosphere of 5% CO<sub>2</sub> at 37 °C. Only cells at log phase of growth were used in experimentation. Cell lines were routinely checked for mycoplasma and were mycoplasma-free. The cell lines are both adherent, grow in a monolayer and are of

## Table 1 – Summary of BRCA1/2 and p53 mutation status in UPN251 and OVCAR8 ovarian cancer cells.

	BRCA1	BRCA2			
Cell line	Mutation status	Methylated	Mutation status		
UPN251	Wild type 1199del29ª,	No	Wild type		
	1246delA <sup>b</sup>				

Analysis of BRCA1/2 status was carried out by [59].

<sup>a</sup> Homozygous deleterious mutation compensated for by another.

<sup>b</sup> Reversion mutation.

epithelial serous histotype. UPN251 originated from a patient who had failed first line platinum/taxane chemotherapy and had relapsed after subsequent treatment of 8 rounds of single-agent taxol chemotherapy (personal communication, Hamilton). UPN251 is BRCA1 wild-type due to a secondary reversion mutation [59]. OVCAR8 was developed from a patient who had undergone treatment with high-dose carboplatin who exhibited progressive ovarian cancer [54]. OVCAR8 is BRCA1 wild-type but is methylated in the promoter region resulting in reduced gene expression of BRCA1 [59]. An overview of UPN251 and OVCAR8's BRCA1 and BRCA2 status is given in Table 1. The cell lines were short tandem repeats (STR) fingerprinted in order to confirm identity. Methylation status was examined and confirmed by Myriad Genetics.

#### Cytotoxicity assays

Acid phosphatase cytotoxicity assays [68] were used to determine cytotoxicity as per method used by [59]. Cells were allowed to attach overnight and then received 5-day exposures to drugs. See supplementary material, Table S1 for list of chemotherapy drugs used in this study including their molecular weight and conversion of 1 unit/ml to unit MW.

### Western blotting

Western blotting was performed as per the method used by [57]. Primary and secondary antibodies used are listed in supplementary material, Table S2. Drug treated cells received  $2 \mu g/ml$  carboplatin or 15 ng/ml taxol for 72 h.

#### Total cellular glutathione assay

Analysis of total cellular levels of glutathione (GSH) was carried out as per the method by [57] which was adapted from [60]. Plates were read and kinetics measured using the FLUOstar OPTIMA (BMG LABTECH) multifunctional microplate reader (405 nM at 30 °C).

## Cell selection strategy outline

Cell lines were treated with carboplatin or taxol as per Fig. 1. Sublines were named in the format of 'Parental cell line-Round and Treatment'. For example UPN251-4T refers to UPN251 treated with 4 rounds of single-agent taxol treatment'. The round parameter can take the values '1'-'7' and the treatment parameter can be 'C' (single-



carboplatin in round 1), 'T' (single-agent taxol) and 'TALT' (alternating treatment starting with taxol in round 1). Treatments were 4-5 weeks apart allowing for all cells to recover before subsequent Q4 drugging. For each round of selection, below steps are followed.

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268 Cells were plated into a T25 flask at a cell density of  $2.6 \times 10^4$ 269 cells per flask and drugged on day 2 as per the selection strategy 270 outline (2 and  $4 \mu g/ml$  carboplatin and 60 and 12 ng/ml for 271 UPN251 and OVCAR8, respectively). On day 5 drugged media was removed and replaced with fresh drug-free media. Over 272 273 subsequent days all T25 flasks were examined for confluence 274 using a novel method to calculate an area fraction output [2]. 275 Upon reaching confluence, cells were re-seeded into a T75 flask. 276 Leftover cells were used to freeze stocks. Cytotoxicity assays were 277 performed at 1 week intervals for 3 weeks and were compared to 278 the parental lines in order to calculate fold resistance. Once all 279 cells had recovered, the next round of drugging commenced 280 following the same format as above (provided the cells were 4 this time).

### **Statistics**

All experiments were repeated at a minimum in biological triplicate excluding the cell selection strategy. Statistical significance analysis was performed by Student's t-test in Microsoft Excel using a two tailed analysis and two samples of equal variance settings.

#### Results

#### Parental BRCA1 protein expression

The BRCA1 protein expression of OVCAR8 (methylated) and UPN251 (un-methylated) was examined by Western blotting.

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n=1

UPN251

n=2 n=3 n=3

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OVCAR8 UPN251 OVCAR8 ARTICLE IN PRESS

OVCAR8 has  $26\% \pm 7\%$  of the expression of UPN251 ( $p = 3.3 \times 10^{-4}$ , Fig. 2). This correlates with the BRCA1 methylation status of the cell lines.

#### Cell selection strategy

The baseline IC<sub>50</sub> values of OVCAR8 were carboplatin  $1.3 \pm 0.2 \ \mu$ g/ml (n=11) and taxol  $1.2 \pm 0.2 \ n$ g/ml (n=10). The baseline IC<sub>50</sub> values of UPN251 were carboplatin  $0.8 \pm 0.1 \ \mu$ g/ml (n=7) and taxol  $17.9 \pm 5.6 \ n$ g/ml (n=9). OVCAR8 has a slightly higher baseline IC<sub>50</sub> to carboplatin and a much lower baseline IC<sub>50</sub> to taxol when compared to UPN251.

#### Dose optimisation

Doses of drug for carboplatin and taxol used in the selection 356 strategy were selected from the following ranges respectively:-357 UPN251 (0.7-2 µg/ml, 10-100 ng/ml) and OVCAR8 (2.3-18.5 µg/ml, 358 2.3-14 ng/ml). Ranges were selected from the results of 3-day 359 cytotoxicity assays on parental cell lines (Table S3) initially encom-360 passing inhibitory concentration (IC) values ranging from 20 to 80. 361 Clinical relevance was validated by investigating clinical trial 362 publications and using pharmacokinetic studies to translate doses 363 364 from the clinic into usable doses in the laboratory.

For carboplatin and taxol a dose range of up to  $20 \ \mu g/ml$  and 120 ng/ml respectively was deemed clinically relevant following pharmacokinetic studies for a dose of carboplatin at AUC 5 and taxol at 175 mg/m<sup>2</sup> which are often administered to patients in clinical trials as single agents ([8,16,23,24,35,37,39,42,45,50,65]).

Cells were subjected to 3-day drug exposures and the time taken for cells to recover was recorded and compared to a drug free control. Desired criteria for the selected doses were that cells would display an initially large amount of cell death (>95%) followed by growth to confluence after drug exposure. Carboplatin

μ

**OVCAR8** 

**UPN251** 

BRCA1 - 220kDa

β-Actin 57kDa

doses of  $4 \mu g/ml$  and  $2 \mu g/ml$  and taxol doses of 12 ng/ml and 60 ng/ml were chosen for OVCAR8 and UPN251, respectively. From the recovery plots (Fig. 3(A) and (B)) we can see that recovery from taxol differed from carboplatin. Taxol treated cells saw a sharp decline in cell number over the first number of days followed by a quick return to confluence thereafter. With carboplatin a more prolonged decline and recovery was noted.

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#### Recovery

In general, all cells recovered quicker after drugging as the rounds of selection progressed. Fig. 4(A) shows recovery plots for each cell line grouped per ascending rounds of selection. In round 7 single-agent treatments received twice the usual dose and consequently recovery time increased. UPN251 cells recovered quicker than OVCAR8 cells and in both cell lines it took longer to recover from carboplatin treatments than taxol. Fig. 4(B) shows the sublines that were treated with carboplatin in each round grouped together for comparison purposes. Single-agent carboplatin treatments (solid bars) were compared with alternating treatments (dashed bars) that received carboplatin in each round. There was little difference in recovery between cell lines receiving treatment with the alternating agents compared to cell lines receiving single-agent carboplatin. Round 2 is the only exception to this. Fig. 4(C) shows the same as above but for taxol treatments. In this case alternating treatments always took longer to recover than single-agent treatments when receiving taxol.

#### Fold resistance

The fold resistance of each subline at weekly intervals for 3-weeks in each round of selection for carboplatin is shown in Fig. 4(D) and (E) and for taxol in Fig. 4(F) and (G) for UPN251 and OVCAR8, respectively. By round 6, UPN251-6T treated solely with taxol displayed the highest level of resistance (7-fold,  $p=0.1 \times 10^{-5}$ ). The sublines developed from UPN251 showed higher levels of resistance compared to those developed from OVCAR8. UPN251-6CALT, UPN251-6T and UPN251-6TALT all had significant resistance to taxol (4–8 fold,  $p = 0.4 \times 10^{-6} - 0.6 \times 10^{-6}$ ) while OVCAR8-6CALT, OVCAR8-6T and OVCAR8-6TALT all had significant resistance to taxol but to a lower extent (1.5–2.5 fold,  $p = 0.02 - 0.2 \times 10^{-6}$ ). All UPN251 sublines after their final round of selection (including UPN251-7T treated only with taxol) had significant resistance to carboplatin (1.6–3.5 fold,  $p = 0.3 \times 10^{-3} - 0.5 \times 10^{-6}$ ). Only OVCAR8-7C and OVCAR8-6TALT had significant resistance to carboplatin in the OVCAR8 sublines (1.3–2.6 fold,  $p = 0.04 - 0.3 \times 10^{-3}$ ). But again this was lower than in UPN251 sublines. In as early as the first round of selection UPN251-1C and UPN251-1T was significantly



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512resistant to carboplatin (1.5-fold,  $p=0.3 \times 10^{-2}$ ) and taxol (1.7-fold,513 $p=0.8 \times 10^{-2}$ ), respectively. These sublines retained significant514resistance with fold resistance increasing from round to round. All515UPN251 sublines receiving the opposite selecting agent in round 2516retained some degree of significant resistance to carboplatin, except517UPN251-2TALT. However UPN251-2TALT regained a significant level518of resistance by round 3.

The cell lines were stable in culture for up to 6 weeks after defrost at which point resistance began to decline (Supplementary

material (Fig. S1)). Experiments were therefore performed in 6 week blocks.

Fig. 5(A) shows the extent of resistance development after 6 rounds of selection for single-agent treatments in OVCAR8 and UPN251 sublines. This was examined to investigate whether cells with BRCA1 defects (OVCAR8) would develop resistance to platinums slower than taxanes with the opposite being true for cells with functional BRCA1 (UPN251). We found that taxol resistance developed quicker in both models irrespective of BRCA1 status.

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Fig. 5(B (i) and (ii)) shows the results of fold resistance after 6 rounds of selection for carboplatin and taxol, respectively. This was investigated as we hypothesised that cells receiving alternating treatments of carboplatin and taxol should develop resistance slower or not at all when compared to single-agent treatments. We can see from these graphs that single-agent treatments have higher fold resistance than all of the alternating treatments.

Fig. 5(C (i) and (ii)) shows the results of fold resistance compare the point in time when each subline had received 3 doses of carboplatin or taxol, respectively. In this case resistance has developed at the same rate in alternating treatments compared to single agent administration in OVCAR8 and resistance has developed quicker in alternating treatments compared to single agent administration in UPN251.

### Mechanisms of drug resistance in UPN251 sublines

Investigation of drug resistance mechanisms were carried out on UPN251 sublines only. OVCAR8 sublines were not examined any further as they developed only low levels of unstable resistance (Fig. 4(E and G)).

#### Drug screen

A drug screen was performed in order to evaluate cross resistance to other drugs and to help elucidate resistance mechanisms that have developed in the cells. A total of 11 drugs and 2 inhibitors were used. Inhibitors include, buthionine sulphoximine (BSO) an inhibitor of glutathione (GSH) (Drew, Miners [7]) and elacridar an inhibitor of *P*-glycoprotein (*P*-gp) a member of the ATP binding cassette (ABC) transporter family [22]. Table 2 gives a summary of all cytotoxicity data collected.

633All of the sublines of UPN251 were significantly resistant to634carboplatin (fold change=1.5-3.2,  $p=0.2 \times 10^{-2}-0.5 \times 10^{-8})$ ,635with single-agent carboplatin developed UPN251-7C being the636highest. The addition of 12.5 µg/ml BSO had the effect of lowering637IC<sub>50</sub> values across all UPN251 sublines. UPN251 and all sublines638(except UPN251-7T) showed significant decreases (p=0.0003-6390.008). Fold resistance however, stayed at a similar level. Sig-640nificant cross resistance to cisplatin and copper sulphate (CuSO<sub>4</sub>)

was also seen in sublines developed with carboplatin treatments. One exception to this is UPN251-6TALT, which is not resistant to CuSO<sub>4</sub>. Oxaliplatin showed significant cross resistance for sublines developed as single-agent treatments while alternating treatments showed no significant cross resistance. UPN251-6TALT showed collateral sensitivity to oxaliplatin (fold change=0.6, p=0.01). The UPN251 sublines developed with taxol all have significant taxol resistance (fold change =4.3–9.0,  $p=0.5 \times 10^{-3}-0.2 \times 10^{-7}$ ), with UPN251-7T having the highest fold resistance. UPN251-7C developed with carboplatin was not resistant to taxol. Comparing taxol, with and without 0.25 µg/ml elacridar, across all cell lines reveals significant drops in IC<sub>50</sub> values ( $p=0.1 \times 10^{-3}$  to  $0.4 \times 10^{-7}$ ). An almost identical trend is seen with vinblastine, and olaparib±elacridar which are all *P*-gp substrates ([3,30]). Doxorubicin and docetaxel also displayed crossresistance in sublines developed with taxol. A significant degree of collateral sensitivity to docetaxel was observed in UPN251-7C (fold change=0.4, p=0.002).

Parp inhibitors veliparib and CEP8983 both showed no significant change in  $IC_{50}$  when compared to the parental cell lines and could both be candidates for treating platinum/taxane-resistant ovarian cancer.

#### Total cellular glutathione assay

Using a total cellular glutathione (GSH) assay (Fig. 6) no significant difference in GSH levels were seen when UPN251 sublines were treated with carboplatin. However UPN251 parental cells saw a significant 3-fold increase in total cellular GSH levels with the addition of 2 µg/ml carboplatin (p=0.02) for a 3-day exposure. Treatment with 12.5 µmol BSO for a 3-day exposure gave significantly reduced levels of GSH for UPN251 and its sublines, when compared to treatment free control cells (fold reduction=9.3–27.8, p=0.5 × 10<sup>-2</sup>–0.02). This was the same dose of BSO which was used in our drug screen.

#### Post selection Western blots

*P*-gp protein expression for UPN251-6CALT and UPN251-7T (control and taxol treated) and UPN251-6TALT (taxol treated) are significantly up-regulated when compared to UPN251 control (Fold Change =  $2.3 \pm 0.9 - 7.3 \pm 2.8$ , p = 0.04 - 0.003, Fig. 7(A)). *P*-gp is significantly decreased in UPN251-7C (carboplatin treated). There are no significant changes in protein expression for both ATP7A (Fig. 7(B)) and CTR1 (Fig. 7(C)) when compared to UPN251 control. MRP2 was not expressed in UPN251 and resistant sublines (supplementary material Fig. S2(A)). There was no change in BRCA1 protein expression between UPN251 parental cells and UPN251-7C and UPN251-7T resistant sublines (supplementary material Fig. S2(C)).

#### Discussion

#### **Resistance models**

The mechanism of resistance that develops in a drug-resistance model can differ depending on the method of selection used. The most common methods of selection used to model resistance are increasing continuous administration ([33,55,64]) and low-dose intermittent incremental inducement ([17,27,62]) where cells are

Drug (Units)	Parent UPN251 IC <sub>50</sub>		Carboplatin single agent UPN251-7C IC <sub>50</sub>			Alternating carboplatin first UPN251-6CALT IC <sub>50</sub>		Taxol single agent UPN251-7T IC <sub>50</sub>			Alternatingtaxol first UPN251-6TALT IC <sub>50</sub>			
	Mean±SD	n	Mean ± SD	n	F	Mean ± SD	n	F	Mean ± SD	n	F	Mean ± SD	n	
Platinums and hear	vy metals													
Carboplatin (µg/	$1.0\pm0.01$	3	3.5±0.07 8***	3	3.3	1.8±0.02****	3	1.7	1.6±0.14 ***	3	1.5	2.0±0.07 ***	3	
ml)														
+BSO (12.5 μg/	$0.8 \pm 0.03$ # # #	3	$2.9 \pm 0.16$ # #	3	3.4	1.6 $\pm$ 0.06 <sup>#</sup> <sup>#</sup>	3	1.8	$1.3\pm0.23$	3	1.5	$1.7 \pm 0.07$ # #	3	
ml)														
Cisplatin (µg/ml)	$0.15\pm0.01$	3	0.48 ± 0.08 **	3	3.3	0.21 ± 0.02 **	3	1.6	$0.15\pm0.01$	3	1.0	0.27±0.04 **	3	
Oxaliplatin (µg/	$0.08 \pm 0.01$	3	0.2±0.03 **	3	2.1	$0.08 \pm 0.02$	3	1.0	0.12 ± 0.003 *	3	1.3	$0.04 \pm 0.008$ *	3	
ml)														
CuSO <sub>4</sub> (ng/ml)	$15.3\pm1.7$	5	28.5±11.4 *	5	1.9	34.7 ± 10.6 **	5	2.3	$17.0 \pm 2.9$	5	1.1	$15.0 \pm 1.3$	5	
Taxanes														
Taxol (ng/ml)	$14.9 \pm 1.9$	3	$14.9 \pm 1.7$	3	1.0	63.6±4.4 ***	3	4.3	133.3 ± 4.7 ***	3	8.9	84.4 ± 11.6 ***	3	
+Elacridar	1.8±0.34 <sup># # #</sup>	3	1.8±0.26 # # #	3	1.0	$1.9 \pm 0.17$ # # #	3	1.1	$1.9 \pm 0.17$ # # #	3	1.0	$2.9 \pm 0.51$ # # #	3	
(0.14 µg/ml)														
Docetaxel (ng/ml)	$4.1\pm0.3$	3	1.5 ± 0.6 **	3	0.4	8.5±1.5 **	3	2.1	20.3±4.0 ***	3	4.9	9.0 ± 2.6 *	3	
Parp Inhibitors														
Olaparib (µg/ml)	$1.7\pm0.36$	4	3.3±0.85 *	4	2.0	3.9±0.73 **	4	2.3	5.4±0.98 ***	4	3.2	3.7±0.4 ***	4	
+Elacridar	$1.3\pm0.14$	4	$2.5\pm0.6$	4	1.8	1.7±0.36 <sup>#</sup> <sup>#</sup>	4	1.2	$1.3 \pm 0.2$ # # #	4	1.0	$1.4 \pm 0.14$ # # #	4	
(0.14 µg/ml)														
Veliparib (µg/ml)	$13.1 \pm 2.91$	3	$14.5\pm0.7$	3	1.1	$14.6 \pm 1.29$	3	1.1	$10.4 \pm 1.58$	3	0.8	$14.7 \pm 1.6$	3	
+Elacridar	$13.7 \pm 2.83$	3	$15.2 \pm 1.79$	3	1.1	$14.0\pm1.9$	3	1.0	$10.7 \pm 1.81$	3	0.8	$14.3\pm1.6$	3	
(0.14 µg/ml)														
CEP-8983 (µg/ml)	$1.4 \pm 0.12$	3	$1.7\pm0.3$	3	1.2	$1.3 \pm 0.24$	3	0.9	$1.1 \pm 0.25$	3	0.8	$1.3\pm0.19$	3	
Vinca Alkaloids														
Vinblastine	$9.9 \pm 1.1$	4	$11.9 \pm 2.84$	4	1.2	29.2±4.54 ***	4	3.0	62.1±3.8 ****	4	6.3	31.2 ± 11.11 **	4	
(ng/ml)														
+Elacridar	3.2±0.57 <sup># # #</sup>	4	5.4±0.65 <sup>#</sup> <sup>#</sup>	4	1.7	3.3±0.73 <sup># # #</sup>	4	1.1	4.6±1.22 <sup># # #</sup>	4	1.5	3.6±0.73 <sup>#</sup> <sup>#</sup>	4	
(0.14 µg/ml)														
Anthracyclines														
Doxorubicin	$39.1 \pm 10.15$	5	$46.8 \pm 2.96$	5	1.2	62.8±3.02 **	5	1.6	117.9±20.2 ***	5	3.0	59.6±6.44 **	5	
(ng/ml)														
Inhibitors														
BSO (µg/ml)	$5.1 \pm 1.87$	4	14.4 ± 2.07 ***	4	2.8	$7.6 \pm 0.22$	4	1.5	$7.9 \pm 2.8$	4	1.5	29.2 ± 11.12**	4	
	$25 \pm 04$	5	1.5+0.3 **	5	0.6	0.8+0.2 ***	5	0.3	1.3±0.5 ***	5	0.5	0.8±0.2 ***	5	



representation of the clinical setting in ovarian cancer, where patients receive drug infusion every 3-4 weeks [41]. Ref. [67] compares the differences in using the pulse versus intermittent incremental strategy in the same ovarian cancer cell lines. They found great differences in the resistance mechanisms that appeared from both strategies. The consensus was that although the intermittent incremental strategy produced higher levels of fold resistance, the mechanisms evolved using the pulse strategy were closer to the mechanisms seen in the clinic and serves as a



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**JPN251** Control

JPN251 Carboplatin UPN251-7C Control

Fig. 6 - (A)-(D).

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941 more 'appropriate' model in studying drug resistance in ovarian
942 cancer. Therefore mechanisms produced from this study have the
943 potential to closely mirror the clinical mechanisms of resistance
944 for ovarian cancer.

#### 946 **Resistance development**

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948 We hypothesised that ovarian cancer cells with BRCA1 defects 949 (OVCAR8) would develop resistance to platinums slower than taxanes with the opposite being true for cells with functional 950 951 BRCA1 (UPN251). Fig. 5(A) shows the extent of resistance devel-952 opment after 6 rounds of selection for single-agent treatments. 953 This hypothesis holds true for OVCAR8 sublines, but not for 954 UPN251 sublines, as taxol resistance developed quicker in both 955 models irrespective of BRCA1 status. Possible reasons for this occurrence may be that taxol treated cells recover quicker than 956 957 carboplatin-treated cells, and therefore resistance can develop 958 faster in taxol-treated cells.

959 We hypothesised that cells receiving alternating treatments of 960 carboplatin and taxol should develop resistance slower or not at 961 all when compared to single-agent treatments. On first inspection this seems to hold true. Fig. 5(B) shows the results of fold 962 963 resistance after 6 rounds of selection for taxol and carboplatin. 964 We can see from these graphs that single-agent treatments have 965 higher fold resistance than all of the alternating treatments. 966 Alternatively, however, if we compare the point in time when 967 each subline had received three doses of taxol (Fig. 5(C) (i)) or three doses of carboplatin (Fig. 5(C) (ii)) during the course of the 968 969 treatment strategy and compared the extent of resistance devel-970 opment, the opposite conclusion could be reached. In UPN251, 971 despite the fact that alternating treatments received the same 972 amount of taxol or carboplatin over a longer period of time when 973 compared to single-agent treatments (five or six rounds in 974 alternating versus three rounds for the single-agent), resistance 975 development was higher in alternating treatments when com-976 pared to single-agent treatments. In OVCAR8, alternating and 977 single-agent treatments are at a similar resistance level. This 978 ambiguity in our results may stem from an inability to directly 979 compare the results of the two drugs due to their different 980 mechanisms of action and speed of recovery from drug treatment.

981 An interesting finding from the selection strategy is that cells 982 treated with taxol, having received carboplatin in the previous 983 round, show large increases in taxol resistance, larger than the 984 increase seen when cells were treated with taxol in the previous 985 round (Fig. 4(E)). Cells with carboplatin pre-treatment also take 986 longer to recover compared to cells which have only received 987 taxol as seen from our area fraction data (Fig. 4(C)). Carboplatin 988 therefore seems to enhance a cell's capacity to become taxol 989 resistant. Further to this, alternating treatments generally dis-990 played notable jumps in taxol resistance in a round in which they 991 received taxol. This is usually greater than the increase in 992 resistance seen per round from single-agent taxol treatments. 993 This is the opposite of what we would have predicted given the 994 inverse resistance relationship between platinum and taxanes 995 Q2 (Stordal et al. 2009). We hypothesised that pre-treatment with 996 one agent would sensitise to the other. We saw no evidence of an 997 opposite effect. Taxol pre-treatment did not affect the amount of 998 carboplatin resistance that developed.

In the literature an in vitro study showed that when cisplatin preceded taxol treatment, lessened antitumor activity was seen

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when compared to taxol before cisplatin [26]. In ovarian cancer cell lines the sequence of cisplatin before taxol reduces taxol induced apoptosis. This was found using DNA fragmentation assays, fluorescence microscopy and flow cytometry [25]. An in vivo mouse study showed that this sequence (cisplatin then taxol) had significant increases in morbidity and mortality associated with it when compared with taxol before cisplatin [36]. In the clinic taxol is given 3 h before carboplatin in order to circumvent carboplatin's myelosupressive affects [42]. Taxol reduces the proportion of bone marrow precursors circulating at the time when carboplatin is given which reduces toxicity when compared to the opposite administration. In non-small cell lung cancer clinical studies, with chemotherapy naive patients, the sequence of carboplatin then taxol administration in combination treatments, showed no sequence-dependant toxicities or pharmacokinetic interactions. However it is not clear whether the different sequences affected response data ([14,21]).

Some evidence which supports the inverse resistance relationship hypothesis is that increased *P*-gp expression was seen in sublines which had taxol treatment during selection while decreased expression was seen in UPN251-7C (carboplatin treated) (Fig. 7(A)). Long term monitoring of UPN251-7C's resistance to taxol showed significant sensitivity to taxol (data not shown). Down regulation of *P*-gp in this cell line may explain this occurrence. As UPN251-7T is resistant to taxol and has high *P*-gp expression it may imply that *P*-gp is involved in the mechanism of the inverse resistance phenotype.

Having received an equivalent cytotoxic drug treatment as UPN251 (BRCA1-wildtype), OVCAR8 (BRCA1-methylated) developed much less resistance to carboplatin or taxol over the same time period. All OVCAR8 sublines were less than 2-fold resistant to carboplatin and less than 2.5-fold resistant to taxol after 6 rounds of selection. This may be due to the cells BRCA1 methylation status. Cells deficient in BRCA1 have reduced efficiency in repairing DNA damage caused by cytotoxic agents. It has been shown that hypermethylation of the BRCA1 promoter region causes increased sensitivity to platinum drugs ([63]). Also in two ovarian cancer cell lines decreasing BRCA1 mRNA using inhibition assays correlated to increased sensitivity to platinums [46]. They also show that patients with low/intermediate levels of BRCA1 mRNA have a significantly improved overall survival following platinum-based chemotherapy compared to patients with high levels of BRCA1 mRNA. Ref. [71] showed that the ovarian cancer cell line SNU251, having a mutation in BRCA1 inhibiting its sub-nuclear assembly, increased its sensitivity to taxol. Also UPN251 and OVCAR8 have had different baseline sensitivities to the drugs used as they originated from patients who had had different levels of exposures to carboplatin and taxol which may have affected the development of resistance.

One caveat to our ability to directly compare resistance developed with carboplatin and taxol is that these drugs may not be directly comparable to each other, due to their different rates of recovery after drugging. This was seen in our dose finding experiment for the selection strategy (Fig. 3) and in the selection strategy itself (Fig. 4). Cells treated with taxol show high initial cell death followed by fast recovery, while carboplatin showed much slower recovery with slight elevations in cell number after drugging, followed by cell death and slow recovery. This difference could be due to the differences in platinum and taxane mechanisms of action. Platinums act mainly by forming nuclear

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1061 platinum adducts on DNA strands ([13,16]), while taxanes act by 1062 stabilising microtubules within the cell ([34,48]). Another caveat 1063 is that the cell lines used in this study were of different genetic backgrounds having been obtained from different patients. A 1064 1065 number of different elements may be at play that has the 1066 potential to affect our results. Future studies could be carried 1067 out in a BRCA1 mutant cell model and a transfected model where 1068 BRCA1 functionality is restored such as UWB1.289 and UWB1.289-1069 BRCA1 [6].

#### 1071 Mechanisms of taxol resistance

1072 Taxol resistance cell models are very common and many have 1073 been developed for ovarian cancer cell lines ([10,11,18,44,70]). 1074 Most of these models use different variations on the above 1075 mentioned intermittent incremental inducement and increasing 1076 continuous administration strategies. Our work is novel as a 1077 pulsed strategy which closely mirrors the clinic has not been 1078 used before in ovarian cancer and a model of taxane-resistance 1079 has not been previously developed in UPN251 to our knowledge. 1080 Over-expression of P-gp often arises as a mechanism of taxol 1081 resistance in cell models. P-gp is the main mechanisms of taxol resistance in our models. Cytotoxicity assays for P-gp substrates 1082 1083 taxol, vinblastine and olaparib + elacridar (Table 2) all show highly 1084 significant drops in IC<sub>50</sub> when P-gp is blocked with elacridar. 1085 Western data showed increased P-gp expression for sublines 1086 which had taxol treatment during selection (Fig. 7(A)). Therefore 1087 it is likely that taxol is being actively pumped out of the cell by 1088 *P*-gp causing taxol resistance.

#### 1090 Mechanisms of carboplatin resistance

Carboplatin resistant ovarian cancer cell lines are rare in the 1091 1092 literature. This is most likely because a combination of cisplatin 1093 and taxol was the standard chemotherapy treatment for advanced 1094 ovarian cancer before 2003, until carboplatin and taxol was 1095 deemed more favourable due to reduced toxicities associated 1096 with carboplatin ([9,42]). A publication by [31] reports on the 1097 development of 5 resistance cell models for ovarian cancer cell 1098 lines (2 carboplatin, 2 cisplatin and 1 taxol). They found a number of genes which were differentially expressed compared to par-1099 ental cells across all resistant models. Another study has devel-1100 1101 oped carboplatin resistant sublines from human larynx carcinoma 1102 cell line Hep 2 by continuous 5-day exposure of increasing doses 1103 of carboplatin. All of the 3 sublines developed had elevated levels 1104 of GSH, but only one of these had significant elevations [40].

1105 From the results of our GSH assays (Fig. 6) we can see that only 1106 UPN251 showed a significant increase in total cellular GSH levels 1107 in response to carboplatin treatment. The developed UPN251 1108 sublines had no significant increases compared to UPN251 with a 1109 carboplatin treatment of 2 µg/ml. This suggests that elevated GSH 1110 plays a role in the parental cells initial response to carboplatin 1111 and that UPN251 resistance sublines utilise other mechanisms. 1112 Treatment with a 12.5  $\mu$ M dose of BSO significantly decreases GSH 1113 in UPN251 and all sublines. This was the same dose of BSO used in our post selection drug screen with carboplatin (Table 2). Small 1114 1115 but significant drops in IC<sub>50</sub> were noted in UPN251 sublines, but 1116 no difference in fold change was noted. This indicates that 1117 increased total cellular GSH may not be a major mechanism of 1118 carboplatin resistance in our developed models.

1119 CuSO<sub>4</sub> (Table 2) had significantly higher IC<sub>50</sub>'s for UPN251-7C
1120 and UN251-6CALT. This indicates the possible involvement of

copper transporters ATP7A, ATP7B and CTR1 in carboplatin 1121 1122 resistance (Safaei, Howell [52]). However Western blots for ATP7A and CTR1 showed little difference in protein expression. These 1123 proteins may instead be relocated to different parts of the cell 1124 causing a resistance phenotype. An increase of ATP7A and ATP7B 1125 in the cellular membrane or a relocation of CTR1 to the golgi 1126 apparatus may lead to platinum resistance without a change in 1127 protein expression [57]. 1128

#### Combined resistance to platinums and taxanes

Models of taxane-platinum resistance are rare in the literature. One study developed a taxane-platinum resistant model for nonsmall cell lung cancer by exposing the cells to cycles of taxol and carboplatin, two cytotoxic agents with different mechanisms of action [5]. Another study developed a dual carboplatin and docetaxel resistant subline from A2780 ovarian cancer cells which are cross resistant to both agents as well as two singularly resistant sublines resistant to each agent but not cross resistant to the other. All of the sublines were selected for in parallel [1]. Gene profiling revealed that the dual model contains genetic changes not present in the singularly resistant models demonstrating that combined drug resistance may not be a simple combination of changes present in single-agent resistant cell lines but can contain novel changes.

Our model presents this novel aspect of subline development for ovarian cancer where sublines were exposed to alternating sequences of taxol and carboplatin. As a result UPN251-6CALT and UPN251-6TALT show significant cross resistance to both carboplatin and taxol used in their development. They show a carboplatin fold resistance of 1.7 and 2 and a taxol fold resistance of 4.3 and 5.7, respectively (Table 2). This is less than the UPN251 sublines selected with single agents but these have no significant cross resistance to carboplatin or taxol except UPN251-7T which is 1.5 fold resistant to carboplatin.

The cytotoxic agent which our sublines were exposed to first influenced the mechanisms of resistance that arose. UPN251-6CALT and UPN251-6TALT both received 3 rounds of drugging with carboplatin and taxol. The only difference is that UPN251-6CALT received carboplatin in the first round whereas UPN251-6TALT received taxol. As a result UPN251-6CALT displays significant fold resistance to  $CuSO_4$  (2.3 fold, p=0.004) while UPN251-6TALT shows no significant fold resistance. Also UPN251-6TALT shows significant sensitivity to oxaliplatin (p=0.01) while UPN251-6CALT has no significant fold change. This would indicate different resistance mechanisms being selected in these cells depending on initial drug exposure. Also, as neither UPN251-7C nor UPN251-7T had significant sensitivity to oxaliplatin while UPN251-6TALT did, this may indicate a novel mechanism of resistance being generated between this dual carboplatin/taxol resistant model and our singularly non-cross-resistant models. This evidence is supported by the finding of [1] discussed above.

# Treatment options for platinum/taxane resistant ovarian cancers

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The baseline  $IC_{50}$  values of parp inhibitors CEP8983 and veliparib in OVCAR8 was  $2.02\pm0.3 \ \mu\text{g/ml}$  (n=6) and  $5.8\pm1.1 \ \mu\text{g/ml}$  (n=4) respectively and in UPN251 was  $1.63\pm0.1$  (n=5) and  $17.5\pm6.8$  (n=4), respectively. UPN251 cells were intrinsically more resistant to CEP8983 than OVCAR8 but had similar baseline sensitivity to

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veliparib. It would be expected that UPN251 cells would be more
resistant to both PARP inhibitors as they have functional BRCA1
when compared with OVCAR8 that has non-functional BRCA1.

1184 These two parp inhibitors, CEP8983 and veliparib, were not 1185 affected by the multiple mechanisms of resistance that arose in 1186 our UPN251 sublines. They did not show any significant resistance 1187 development and may be candidates in treating platinum/taxane 1188 resistant ovarian cancers. This data and the results of our recent 1189 study on a panel of 41 ovarian cancer cell lines [59] suggests a broader activity of parp inhibitors in BRCA1 wild-type ovarian 1190 1191 cancer. This activity is likely due to a variety of mechanisms 1192 causing dysfunction in homologous recombination repair.

## Conclusions

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The development of taxane resistance was not slower that the 1197 development of platinum resistance in cells with functional BRCA1 1198 as was expected per the inverse resistance relationship. Taxol 1199 resistance developed quicker in BRCA1-wildtype and BRCA1-1200 methylated cells. Both resistance to carboplatin and taxol devel-1201 oped quicker and more stably in UPN251 (BRC1-wildtype) com-1202 pared to OVCAR8 (BRCA1-methylated). Also alternating carboplatin 1203 and taxol treatment delays but does not prevent resistance 1204 development when compared to single agent administration. This 1205 was expected from the inverse resistance relationship. However, 1206 interestingly, the sequence of drug exposure influenced the resis-1207 tance mechanism that developed in resultant sublines. UPN251-1208 6CALT and UPN251-6TALT have different profiles of cross resistance 1209 to drugs, one having received carboplatin and one having received 1210 taxol in round one of development being their only difference. 1211 Finally over expression of *P*-gp is the dominant mechanism of taxol 1212 resistance present in our UPN251 resistant sublines whereas 1213 multiple mechanisms of carboplatin resistance are postulated to 1214 be present in our cell models. 1215

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## Appendix A. Supporting information

Supplementary data associated with this article can be found in the online version at http://dx.doi.org/10.1016/j.yexcr.2014.12.001.

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