

Oleuropein is a Powerful Sensitizer of Doxorubicin-mediated Killing of Prostate Cancer Cells and Exerts Its Action via Induction of Autophagy

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Abstract The phenolic component Oleuropein (OLEU), a bioactive natural product, has recently shown antiproliferative properties. Doxorubicin (DXR) is an anthracycline present in many chemotherapeutic schemes, although limited due to its cardio-toxic effects. Important research effort has been devoted therefore, to reducing DXR toxicity without compromising its antitumor efficacy. The anticancer actions of DXR and OLEU were assessed, on PC-3 prostate cancer cells, while cell cycle distribution and rate of apoptosis were assessed by flow cytometry. The autophagic process was determined via immunoblotting and immunofluorescent staining. Finally, cell extracts were analyzed by NMR spectroscopy. The present study showed that both DXR and OLEU inhibited PC-3 cells proliferation, while the co-treatment with DXR and OLEU resulted in an additive inhibition. Although the addition of OLEU to DXR did not alter significantly the cell cycle distribution, exhibited by each treatment alone, and produced a marginal increase on the rate of apoptosis, both compounds produced a remarkable induction of autophagy. In addition, treated cells exhibited significant metabolite alterations. This study demonstrates that OLEU, a basic component of the everyday diet, is capable of lowering significantly the cytotoxic dose of DXR, while obtaining an important anti-proliferative effect in prostate cancer cells.

Keywords: oleuropein, doxorubicin, autophagy, prostate, metabolic profiling

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1. Introduction

Prostate cancer is very common in Western populations and is the second leading cause for male death from cancer in North America. The incidence of cancer in Mediterranean countries is lower than in the rest of European countries and the United States. This is mostly described by the lower rate of the large bowel, breast, endometrial, and prostate cancers by a number of epidemiological studies, and the major reason for this, apart from possible genetic factors, is attributed to the dietary practices [1]. The traditional Mediterranean diet is characterized by high consumption of foods of plant origin, relatively low consumption of red meat, and high consumption of olive oil and its products [2].

Doxorubicin (DXR) is a potent anthracycline antibiotic that has been used in anticancer therapy for decades. However, DXR is also well known to exert toxic effects on normal tissues. Especially in the heart, DXR can induce an accumulative dose-dependent cardiomyopathy that ultimately leads to congestive heart failure. Despite its severe toxicity, DXR remains a major component of most chemotherapeutic regimens because of its efficacy and broad spectrum antitumor activity. As a result, sustained research effort has been devoted to identifying effective drugs or strategies that can reduce DXR toxicity without compromising its antitumor efficacy [3].

Since olive oil is one of the most valuable nutritional agents and is also the major energy source in the Mediterranean diet, several studies were recently conducted on the beneficial effects of olive oil (extracted mechanically from olives, the fruit of *Olea europaea* L) and on oleuropein, the most prominent phenolic

compound in olive fruit and leaves. Oleuropein (OLEU) has several pharmacological properties, including antioxidant, anti-inflammatory, anti-atherogenic and anti-cancer actions and for these reasons, it is commercially available as food supplement in Mediterranean countries [4,5]. Although Oleuropein has been shown to exhibit anti-proliferative activity *in vitro* against a number of cancer types, the underlying action mechanism remains largely unknown [6,7]. In addition, OLEU has been shown to be cardioprotective against acute doxorubicin cardiotoxicity and has been shown to exhibit anti-ischemic and hypolipidemic activities [8,9].

The healthy effects of plant polyphenols, some of which characterize the so-called Mediterranean diet, have been shown to arise from epigenetic and biological modifications resulting, among others, in autophagy stimulation. Autophagy is one of the major pathways for degradation of cellular components in animal cells that controls the turnover of long-lived proteins and organelles. Although initially identified as a process induced by cellular starvation, an autophagic pathway is now recognized as the cellular response to a variety of stimuli [10,11]. During autophagy, cytoplasmic components are engulfed by double membrane-bound structures (autophagosomes) and delivered to lysosomes for degradation. During autophagosome formation, a cytosolic microtubule-associated protein-light chain 3 (LC3) is processed and conjugated with phosphatidylethanolamine (PE). The PE-conjugated LC3, LC3-II, is inserted into autophagic vesicle membranes. Because the amount of LC3-II correlates with the number of autophagosomes, LC3-II is widely considered as an autophagosomal marker for monitoring autophagy [12,13].

The objective of the present study was the evaluation of the cytostatic action of OLEU, DXR and, more interestingly, of the combined treatment with OLEU and DXR, on PC-3 prostate cancer cells. Furthermore, the scope of this study was the investigation of the mechanism of action of OLEU and DXR, in terms of the impact on the autophagic pathways and the identification of the metabolic profile of prostate cancer cells.

2. Materials and Methods

2.1. Oleuropein Isolation

The oleuropein used in this study was isolated from *Olea europaea* (*Oleaceae*) leaves as described before [8]. Briefly, air-dried and pulverized leaves were extracted with mechanical stirring for 12 h with acetone. The extract was evaporated to dryness and was washed with a mixture of dichloromethane/methanol (98/2). The insoluble material was separated, dried and submitted to medium pressure liquid chromatography with Si gel 60 Merck, using dichloromethane-methanol gradient as the eluent to afford pure oleuropein.

The structure elucidation of oleuropein was carried out using spectroscopic and spectrometric methods as well as comparison with literature data [14]. Specifically, 1D&2D NMR spectra (COSY, COSYLR, HSQC-DEPT, HMBC) were recorded in deuterated methanol (CD₃OD - Merck), on a Bruker Avance III spectrometer (Bruker Biospin GmbH, Reinsteten, Germany) operating at 600.11 MHz

for ¹H and at 150.11 MHz for ¹³C, with a 5 mm inverse detection probe. The residual ¹H (3.33 ppm) and ¹³C (49.50 ppm) signals of CD₃OD were used as internal standard. 1D & 2D NMR experiments were performed with standard pulse programs, at room temperature. HRMS & HRMS/MS data were obtained by direct infusion method using an hybrid LTQ-Orbitrap Discovery Mass Spectrometer (Thermo Scientific, Bremen, Germany) equipped with an ESI probe, in positive mode. Oleuropein's purity (95%) is the maximum that can be obtained from a natural compound.

2.2. Cell Line

The PC-3 cells, an androgen insensitive, p53-negative and K-Ras mutated human prostate cancer cell line, were obtained from the American Type Cell Culture (ATCC, Bethesda, Md). This cell line was grown in 75-cm² culture flasks at 37°C in 5% CO₂ using Dulbecco's modified Eagle's medium F/12 (DMEM/F-12, Gibco, USA) containing 10% fetal bovine serum (FBS, Gibco, USA) [15].

2.3. MTT Proliferation Assay

PC-3 cells were plated at a cell density of 500 cells/well in 96-well plates 24 hrs prior to treatment with the appropriate drug at various concentrations for 96 hrs. The anticancer actions of DXR (3-100 nM) and of OLEU (3-400 µg/mL) were assessed, alone and in co-treatment, on PC-3 cells. The cells were incubated with 10% MTT (Sigma M-5655, USA) added directly in the medium for 4 hrs at 37°C. The medium was then aspirated and the cells were solubilized with the organic solvent dimethylsulfoxide (DMSO). Absorbance was determined in a VERSA max microplate reader (Molecular Devices Corporation, USA) at 540 nm and results are presented as the percent of OD in the treated wells versus the controls [16].

2.4. Flow Cytometry

Cell-cycle analysis was performed following incubation of PC-3 cells with DXR (50 nM) and OLEU (200 µg/mL), alone and in co-treatment, for 24, 48 and 72 hours. Cells were then washed in PBS, fixed in 50% ethanol solution (v/v), and stained with an RNase-containing propidium iodide (PI) solution (50 µg/ml). DNA content was analyzed on a FACS Calibur (Becton Dickinson, San Jose, CA) flow cytometer using the ModFit software (Verity Software House, Topsham, ME). Apoptosis of PC-3 cells was assessed by analysing DNA content in Annexin V-FITC and propidium iodide (PI) stained cells (Annexin V-FITC Apoptosis Detection Kit, BD Systems) on a BD Accuri C6 flow cytometer.

2.5. Immunofluorescence Labeling

For the detection of the autophagic marker LC-3, PC-3 cells were plated at a cell density of 20,000 cells / well on Lab-Tek chamber slides were treated with DXR (50 nM) and OLEU (100 µg/mL, 200 µg/mL), alone and in co-treatment, for 24h. After a wash with cold phosphate-buffered saline (PBS), they were fixed in methanol for 10 min at -20°C, air-dried, washed again with PBS, incubated in 3 % H₂O₂ in methanol for 10 min, incubated in 50 mM ammonium chloride in PBS for 15 min at room

temperature and then washed with PBS. After blocking non-specific protein binding by incubation with blocking serum (Ultra Vision Detection System Anti-Polyvalent, Lab Vision Corporation, USA) for 20 min at room temperature and washing with PBS, the cells were incubated with primary antibody (LC3 rabbit polyclonal antibody; Novus, USA) for 60 min at room temperature. After another wash with cold PBS, the cells were treated with fluorescein isothiocyanate (FITC)-labeled secondary antibody (Santa Cruz Biotechnology, USA) for 60 min at room temperature, washed again with PBS, mounted with Aqua Poly Mount (Polysciences, Germany) and analyzed using a Nikon fluorescence microscope with a wide-band green filter.

2.6. Protein Isolation

After 48h treatment with the appropriate factors cells, grown in 25-cm² culture flasks, were washed twice with ice-cold PBS and scraped into chilled PBS with protease inhibitors. The cells were pelleted and the pellets were resuspended in cell lysis buffer containing Tris-HCl, NaCl, NaF, Na₄P₂O₇, Na₃VO₄, DTT, Triton X-100, protease inhibitors and phosphatase inhibitor cocktail I (Sigma, USA) 1/100. Cells were passed through a 21 gauge needle, followed by incubation on ice for 20 minutes. After centrifugation the supernatants were collected and assayed for protein concentration using the Bradford method (Bio-Rad Laboratories, USA).

2.6.1. Western Blot Analysis

20 µg of each protein sample were resolved on SDS 4-15% gradient PAGE and electrophoretically transferred to nitrocellulose (Bio-Rad Laboratories, USA) for overnight incubation, at 4°C, with LC3 rabbit polyclonal antibody (Novus, USA). Blots were then probed with peroxidase-conjugated species appropriate secondary antibodies (Santa Cruz Biotechnology, USA) and visualized by ECL (GE Healthcare, Germany) and exposed to an X-ray film (Kodak, Germany). Equal protein loading was verified using anti-GAPDH monoclonal antibody (Santa Cruz Biotechnology, USA). Intensity ratios of the LC3-II over LC3-I proteins were calculated and the control sample was used as reference value.

2.7. Cell Extraction and Sample Preparation for NMR Experiments

Cell samples (approximately 10⁷ cells/NMR sample) were extracted by the methanol-chloroform-water (M/C) protocol described by *Le Belle et al.* [17] (n=5/group). Methanol and chloroform (in reagent-grade) were added to the frozen cell pellets (4 °C) in a ratio of 2:1 (v/v; 250 µL/cell pellet). The tissue-solvent mixture was allowed to thaw before being transferred to centrifuge tubes. The cell pellet-solvent mixture was sonicated. After approximately 15 min in contact with the initially added solvents, chloroform and distilled water were added to the samples in a ratio of 1:1 (250 µL/cell pellet), in order to form an emulsion. The samples were then centrifuged at 13,000 rpm for 20 min. The upper phase (methanol and water) was separated from the lower (organic) phase and both fractions were dried at room temperature using a speed-vac. The protein pellets from the cells were re-extracted,

but the separated fractions were pooled with the original extracted fractions before drying the samples.

The aqueous cell extracts were reconstituted in 700 µL of 99.9% D₂O at pH 7.4 phosphate buffered (150 mM) containing 0.01% 1,1,2,2-tetradeutero-3-trimethylsilylpropionic acid (TSP) as internal standard and 0.2 mM NaN₃ for microbial growth inhibition. The corresponding lipid cell extracts were reconstituted in 700 µL of CDCl₃ using tetramethylsilane (TMS) as an internal standard for further NMR experiments.

2.7.1. NMR Experiments

All NMR experiments were acquired on a 600 MHz Avance III spectrometer (Bruker GmbH, Germany) equipped with a B-ACS-60 sample changer at temperature probe 27°C. For each sample 1D 1H NMR (noesypr1d pulse sequence) and ultra-fast *J*-resolved spectra were acquired with water signal suppression during relaxation and mixing time. For the 1D spectra 32 free induction decays (FIDs) were collected into 64k data points at a spectral width of 12335.5 Hz and with acquisition time of 2.65 s, relaxation delay of 4 s and mixing time 0.010 s. The NMR spectra were phased corrected and referenced to the chemical shift of TSP at 0.00 ppm using the TOPSPIN v.3.1 software (Bruker BioSpin). The *J*-resolved experiments were acquired with 64k points for a spectral width of 10000 Hz. Prior to Fourier transformation an exponential weighting factor corresponding to a line broadening of 0.3 Hz was applied. For selected samples 2D HSQC heteronuclear and 2D COSY and TOCSY homonuclear spectra were recorded to aid resonance assignment according to spectral details described elsewhere [18].

2.7.2. Data Reduction of the NMR Spectra and Univariate Statistical Analysis

NMR spectra (spectral width from 9.49 to 0.71 ppm) were segmented into 431 chemical shift regions of 0.02 ppm width using the software package AMIX Statistics v.3.9.14 (Analysis of MIXtures, Bruker BioSpin) and the integrals were calculated. Regions of a) 7.67-7.70 ppm (unknown peak), b) 4.85-4.70 ppm (residual water suppression trace), b) 3.38-3.34 ppm (methanol) and c) 0.71-0.00 ppm (region with no significant information) were removed from the data set before the application of AMIX package. Data were normalized according to the total intensity of the spectra.

2.8. Statistical Analysis

All the variables are expressed as mean ± standard error of the mean and compared among groups using One-way Analysis of Variance model (ANOVA) with Bonferroni correction and Tukey post-hoc analysis via the software package IBM SPSS Statistics v. 21. A p value <0.05 was considered to be statistically significant. For the NMR analysis, the z-scores were also calculated from normalized bucket integrals and averaging for each group according to the formula $z\text{-score} = (\text{mean}_{\text{Group}i} - \text{mean}_{\text{Control}}) / \text{SD}_{\text{Control}}$, in order to highlight the metabolites which are responsible for the differentiation of the metabolic profiles between the four groups (doxorubicin, oleuropein, doxorubicin-oleuropein, control).

3. Results

3.1. Cytotoxicity / MTT Assay

Increasing concentrations of OLEU (3-400 $\mu\text{g/mL}$) produced a dose-dependent inhibition in the proliferation of PC-3 cells (Figure 1A). Continuous exposure to DXR (3-100 nM) produced a maximal inhibition of 66 % on the growth of PC-3 cells at the high doses tested (Figure 1B). The co-treatment of PC-3 cells with DXR (3-100 nM) and 100 $\mu\text{g/mL}$ of OLEU did not alter the cell proliferation inhibition obtained by DXR alone, whereas the co-treatment of PC-3 cells with DXR (3-100 nM) and 200 $\mu\text{g/mL}$ of OLEU resulted in an additive inhibition of cell proliferation even at very low doses of DXR (3-12,5 nM) (Figure 1B).

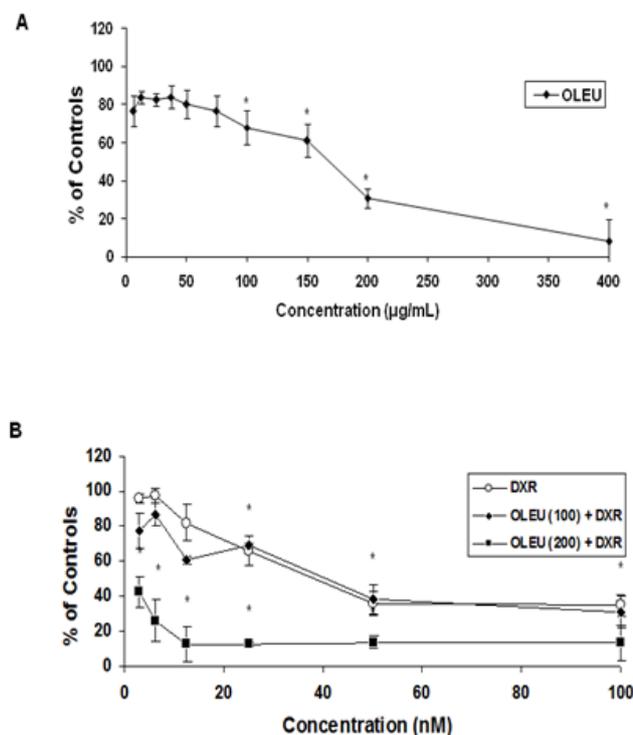


Figure 1. Dose-response curves for the continuous administration (96 hr) of a) OLEU alone (3-400 $\mu\text{g/mL}$), b) DXR alone (3-100 nM) and DXR in co-treatment with either 100 $\mu\text{g/mL}$ OLEU or 200 $\mu\text{g/mL}$ OLEU on PC-3 cells, as assessed by the MTT assay. Results are expressed as percentage of controls. Statistical analysis was performed in triplicate determination at $p < 0.05$.

3.2. Flow Cytometry

The cell cycle distribution showed that treatment of the PC-3 cells with 200 $\mu\text{g/mL}$ of OLEU for 24, 48 and 72 hours produced a marginal increase in the cell distribution in G0/G1 phase, while treatment with 50 nM of DXR showed the expected G2/M phase blockade and suppression of the S phase. The addition of OLEU to the treatment with DXR did not alter the effect exhibited by DXR alone (Figure 2A, Figure 2B, Figure 2C). Additionally, the combination of Annexin V-FITC and PI staining allows differentiation between early apoptotic cells, late apoptotic cells, necrotic cells and viable cells. The addition of OLEU to DXR, after 48 hours exposure, produced evidence of late apoptosis (ranging from 10-15 %) above that noted by each treatment alone, under these experimental conditions (data not shown).

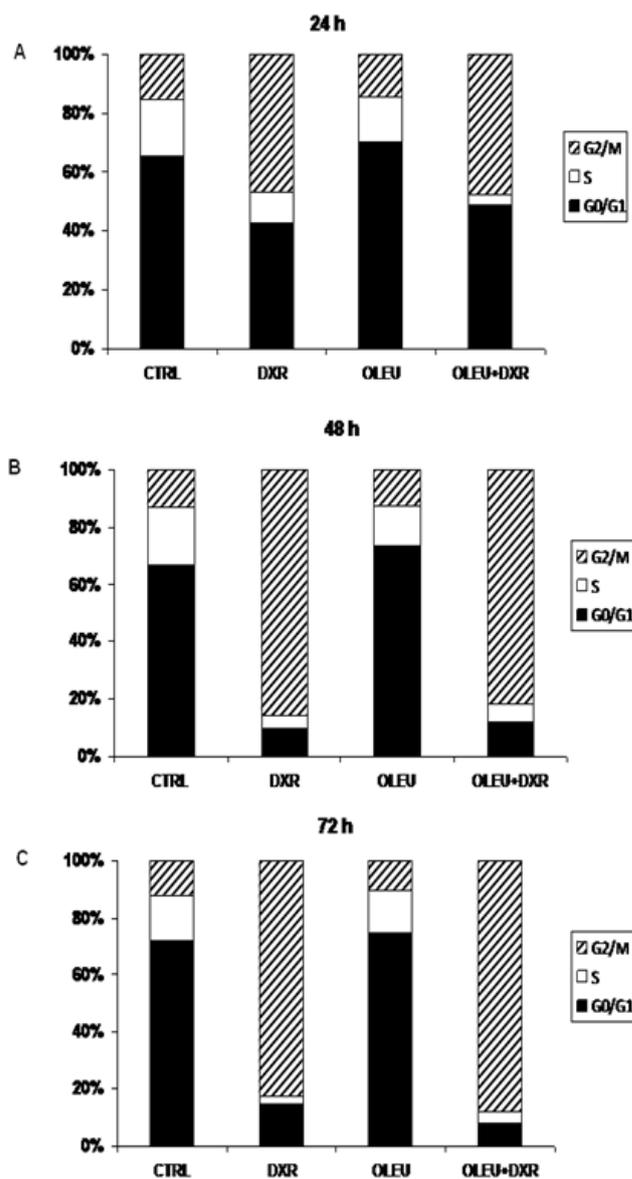


Figure 2. Flow cytometry graphs showing DNA content in PC-3 cells, after a) 24 hours b) 48 hours and c) 72 hours of exposure to DXR (50 nM) and OLEU (200 $\mu\text{g/mL}$), alone and in co-treatment in comparison to control cells (CTRL). Quantification of G0/G1, S and G2/M phases was based on staining by the DNA binding dye propidium iodide.

3.3. Autophagy Imaging

The immunofluorescence experiments revealed the induction of autophagy, in PC-3 cells that received OLEU, DXR and the co-treatment scheme. It was noticed a more pronounced signal in cells that received the combination of 50 nM of DXR with 200 $\mu\text{g/mL}$ of OLEU (Figure 3).

3.4. Immunoblot Analysis for LC3

The unconjugated (LC3-I) and conjugated forms (LC3-II) of LC3 in PC-3 cells are resolved by protein electrophoresis as two distinct bands. In fact, treatment with 50 nM DXR, 200 $\mu\text{g/mL}$ OLEU and the co-treatment scheme (DXR + OLEU) resulted in the induction of LC3 membrane translocation, the disappearance of LC3-I protein and the consequent increase in the LC3-II/LC3-I ratio. The enhancement of autophagy is even more evident following the co-treatment of 50 nM DXR and 200 $\mu\text{g/mL}$ OLEU (Figure 4).

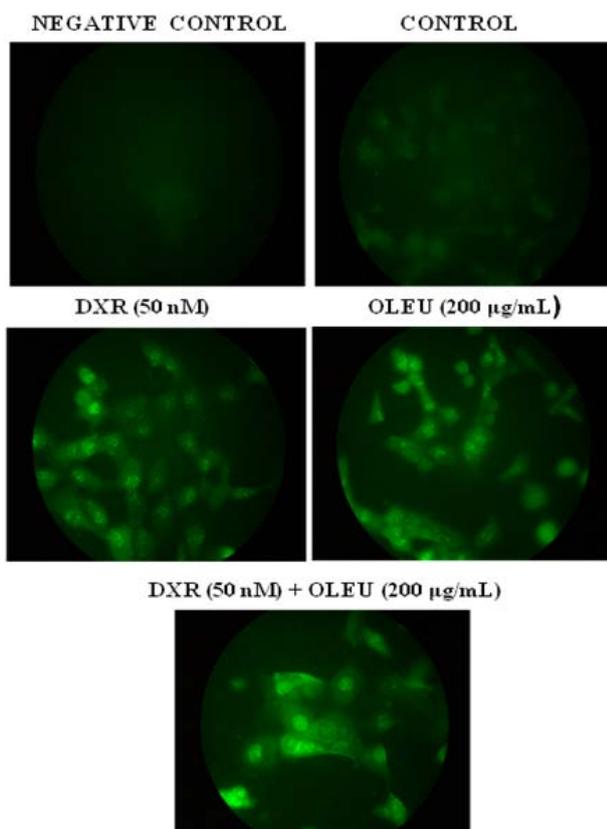


Figure 3. An example of the expression of LC3 (autophagic marker) in PC-3 cells in response to DXR (50 nM) and OLEU (200 µg/mL), alone and in co-treatment, by immunofluorescence, as described in the Materials and Methods section. All treated samples enhanced the expression of LC3 as compared to controls. Note the presence of autophagosomes in treated cells (Magnification: x60)

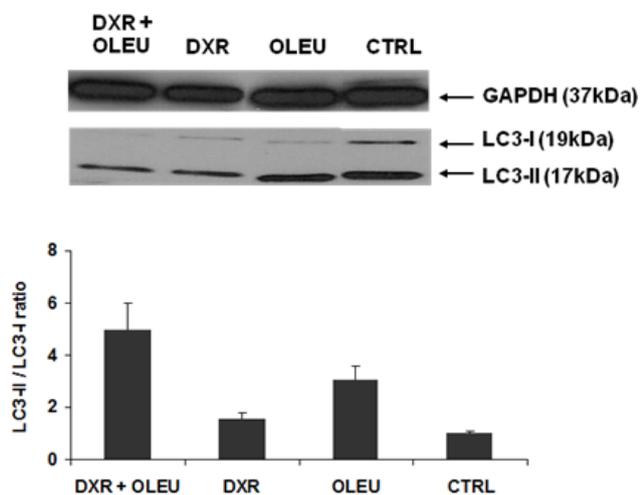


Figure 4. Representative Western blot and densitometry of the expression of the unconjugated (LC3-I) and conjugated forms (LC3-II) of LC3 in PC-3 cells. The upper band (LC3-I) is the autophagy-inactive form of the protein and the lower one (LC3-II) is the active-membrane bound form. The analysis of the intensity of LC3-I and LC3-II by western blot was expressed as a ratio (LC3-II/LC3-I) of the positive control and was normalized against the respective GAPDH signal. DXR (50nM), OLEU (200 µg/mL), DXR (50nM) + OLEU (200 µg/mL)

3.5. Metabolic Profiling

Differences in the metabolic profile between different groups were revealed using NMR based metabolic profiling. A typical spectrum of cell extracts is illustrated in Figure 5.

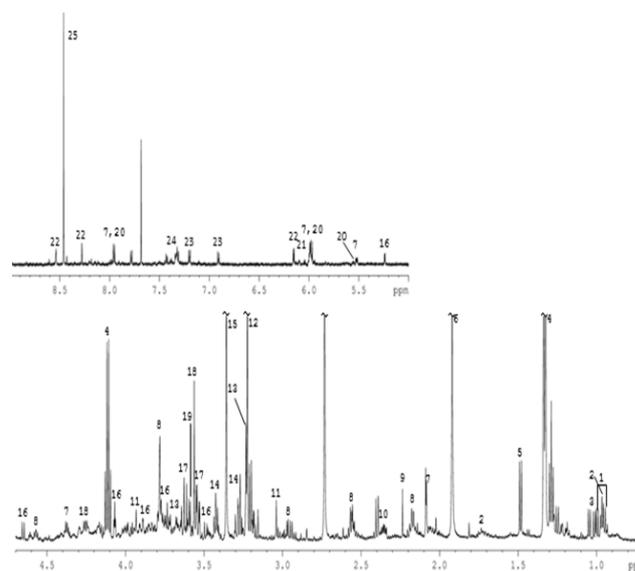


Figure 5. ^1H NMR spectrum of a representative control sample along with the assignment of metabolite signals.

[1: Isoleucine; 2: Leucine; 3: Valine; 4: Lactate; 5: Alanine; 6: Acetate; 7: UDP-N-Acetylglucosamine; 8: Glutathione; 9: Acetone; 10: Glutamate; 11: creatine or creatine phosphate; 12: O-Phosphocholine; 13: sn-Glycero-3-Phosphocholine; 14: Taurine; 15: Methanol; 16: Glucose; 17: Myo-Inositol; 18: Glycine; 19: Threonine; 20: UDP-N-Acetylgalactosamine; 21: NAD⁺; 22: AXP; 23: Tyrosine; 24: Phenylalanine; 25: Formate]

Twenty-eight metabolites were assigned deviation from controls (according to the normalized data) after different treatments is depicted by z-scores in Table 1.

Table 1. ^1H NMR chemical shifts and assignments for cell metabolites and the deviation of each treatment compared to control as depicted by z-scores (z-scores are colored as a heat map for easier interpretation). * $p < 0.05$ compared to control samples. DXR (50nM), OLEU (200 µg/mL), DXR (50nM) + OLEU (200 µg/mL)

Metabolites	δ , ^1H (ppm)	z-scores vs control		
		DXR	DXR+OLEU	OLEU
Formate	8.47-8.45	-0.01	0.71	0.09
Phenylalanine	7.45-7.43	-2.87	0.60	0.05
Tyrosine	7.21-7.19	-1.49	1.18	-4.69*
ATP	6.15-6.13	-0.86	15.15*	-5.27
NAD	6.05	-0.06	-0.06	-1.18
UDP-N-Acetylglucosamine	5.53-5.51	1.40	14.50*	-0.76
Glucose	5.23	1.12	5.83*	0.91
Myo-Inositol	4.07	-0.11	0.02	0.35
Glycine	3.57	-0.55	1.13	1.11
Taurine	3.43	-2.71*	-1.01	-0.79
sn-Glycero-Phosphocholine	3.23	-1.56	-0.14	2.03
Creatine	3.05-3.03	-1.45	0.05	1.56
Glutathione	2.55	-2.50	-4.59*	-4.10*
Succinate	2.41	1.04	0.03	0.00
Glutamate	2.35	-0.58	-0.90	-0.95
Acetate	1.93-1.91	-2.95	-4.03	-4.61
Alanine	1.49	-0.27	-1.71	-1.71
Lactate	1.33	0.61*	-0.30	-1.11
3-Hydroxy-isovalerate	1.25	5.32	0.43	2.64
Valine	1.05	0.13	0.41	-0.65
Leucine	0.97	-0.64	-0.24	-0.76
Isoleucine	0.95	0.02	0.55	-0.21



Univariate ANOVA statistical analysis results are incorporated also in the table as indicated with the asterisk when statistical significant difference is observed. Significant alterations in DXR treatment are phenylalanine, taurine and acetate exhibiting a downregulation, while 3-Hydroxy-isovalerate exhibits an interesting upregulation primarily in DXR and secondary in OLEU treatment. Tyrosine, ATP and acetate exhibit a strong reduction for the OLEU compared to control group. More importantly substantial increase of ATP, UDP-N-Acetylglucosamine and glucose levels is observed in DXR + OLEU co-treatment as compared to control. Furthermore a concomitant reduction for glutathione and acetate is shown in all treated cell samples.

4. Discussion

To overcome advanced prostate cancer, the commonly used treatment to date is the inhibition of androgen production and/or androgen function. In the past the development of anticancer drugs has traditionally relied on *in vitro* tests aimed almost exclusively at assessing the potential of direct killing or growth inhibiting of cancer cell lines. However, accumulating evidence indicates that the *in vivo* response to anticancer therapies is mostly influenced by the toxicity and side-effects that the chemotherapeutic schemes may confer. Doxorubicin, a known topoisomerase II inhibitor, producing G2/M arrest and apoptosis in a cellular level, affects significantly the growth of prostate cancer cells. However, the severe cardiotoxicity that gradually develops, erases many questions about the "pour ou contre" of DXR applications. Therefore, many researchers have launched studies aiming at reducing the DXR-induced toxicity [19,20,21]. These studies have until now, focused on the induction of apoptosis, while the present study addresses the combination of DXR with oleuropein, a natural product, in the field of prostate cancer, with regard to the more novel mechanism of autophagy.

Nowadays, there is a strong evidence for the protective role of oleuropein against many types of cancer [22,23,24]. More specifically, Acquaviva *et al.*, have recently demonstrated the antiproliferative effect of oleuropein in the androgen sensitive LNCaP and in the androgen insensitive DU145 prostate cancer cell line which are characterized by low and moderate tumorigenicity, respectively [25]. Herein, it is studied the antiproliferative effect of oleuropein and its potent synergistic action with DXR, in the androgen insensitive p53-negative human PC-3 prostate cancer cell line, originally derived from the bone metastatic lesion of a prostate cancer patient. Because the characteristics of PC-3 cells consent with many of the clinical aspects of the hormone refractory prostatic cancer, this cell line has been widely used as a model of prostate cancer progression [15].

This study demonstrates that both DXR and oleuropein inhibit PC-3 cells proliferation in a dose-dependent manner, while the co-treatment with DXR and oleuropein results in an additive inhibition of cell proliferation even at very low doses of DXR. This finding is considered of great importance, as oleuropein is capable of lowering significantly the cytotoxic dose of DXR, while obtaining an outstanding antiproliferative effect in prostate cancer

cells. Additionally, the fact that the effectiveness of DXR in inhibiting cell proliferation is not compromised when oleuropein is present, which may not always be taken for granted, demonstrates that the anticancer activity of DXR is not alleviated by oleuropein.

Furthermore, the fact that there was no significant alteration of the cell cycle distribution and that there was a marginal increase on the rate of late apoptosis, between the cell samples treated with the combination scheme and the treatment solely with DXR, led the authors to investigate the implication of another intracellular molecular mechanism, autophagy. Indeed, the results show that DXR and OLEU produce a remarkable induction of autophagy, which is more pronounced following the co-treatment scheme. This finding was confirmed by both immunofluorescence and immunoblotting. Although autophagy has been proposed as a cell death process, the role of autophagy in cancer cell death is still in dispute. Previous studies have shown that when tumor cells are deprived of growth/survival factors, autophagy is increased to prevent the cells from dying [26]. In addition, when autophagy is prevented under these conditions, the cells undergo apoptosis. Therefore, autophagy seems to play a role in preventing cellular apoptosis from nutritional stress in cancer cells. In contrast, several other studies have demonstrated that autophagy has an anti-cancer effect. However, it is unclear whether autophagy is the mechanism for cell death or a reactive mechanism by which the cell is trying to survive the chemotherapy [27].

Cancer cells often modify their metabolism by promoting diverse biosynthetic pathways to adapt to the environment and to ensure their rapid proliferation. Hence, a growing appreciation of the role of autophagy in controlling cellular metabolism has fuelled immense interest in elucidating how dysfunctional autophagy influences metabolic disorders and metabolic adaptation in diseases such as cancer. Furthermore, autophagy can mobilize diverse cellular energy stores to replenish metabolites during both normal and stressed conditions. Recent studies delineate a direct crosstalk between metabolic enzymes and autophagy and that, as an intracellular recycling system, autophagy is highly important for cell metabolism and nutrient allocation. Therefore, it is highly supported that there is a direct regulation of autophagy by the provided cell metabolic context [28]. In the present study, treated cell samples exhibited significant metabolite alterations, which highly correlate with the cell autophagic response. Among the identified metabolites, those involved in energy production (ATP, glucose and glutamate), amino acid metabolism (glycine, alanine, valine, leucine and tyrosine), as well as in oxidative profile (glutathione), of rapidly dividing tumor cells seemed to be starring in separating the groups DXR, OLEU, DXR+OLEU and Control. These results indicate that several specific metabolic pathways are disturbed in prostate cancer tissue. Sustained autophagy under conditions of protracted starvation has been proposed to lead to cell death; thus, the survival or death consequences of autophagy are condition-dependent.

Growth in a hostile environment, inefficient utilization of glucose and defective autophagy predict that prostate cancers may be particularly sensitive to therapies that inflict metabolic stress. It is, therefore, hypothesized that

prostate cancer is metabolically fragile because of dependence on glycolysis and impaired autophagy [10]. Specifically, among the treated cells, the downregulation of taurine and glutamate, characteristics of protein catabolism taking place in autophagy, correlate well with the proposed mechanism of action of the treatments tested. Moreover, the concomitant downregulation of reduced glutathione not only highlights the antioxidant role of DXR and the even stronger antioxidant effect of OLEU, but also confirms the high endogenous glutathione levels of cancer cells as a result of the cancer-mediated cell stress [29]. It is also noteworthy, that human prostate cancer cell lines of varying degrees of aggressive behavior have distinct redox properties and that each cell line shows distinct cytotoxic responses to low-molecular-weight redox-modulating compounds [30].

An observation of utmost interest in the metabolic profiling analysis is the outstanding increase of UDP-N-Acetylglucosamine and ATP along with glucose in DXR and OLEU co-treatment compared to controls. Although ATP is mostly used to provide energy for cellular processes, it also directly links energy metabolism to signaling. Indeed, Kawaguchi *et al.* recently demonstrated that cancer cells promote autophagy to overcome the energy shortage from glycolysis, the major energy source, by maintaining mitochondrial activity to produce ATP necessary for survival [31]. These data show that mitochondria are not simply the power plants of cells generating ATP, but are closely related to several types of cell death and autophagy. Uridine diphosphate N-acetylglucosamine (UDP-GlcNAc) on the other hand, which is substantially regulated by the combination scheme (DXR+OLEU) as well, is the high energy donor substrate for O-linked N-acetylglucosamine. UDP-GlcNAc sits at the nexus of glucose, nitrogen, fatty acid and nucleic acid metabolic pathways, all of which dynamically influence its cellular concentration. Mammals respond to nutrient excess and different other forms of stress (heat shock, oxidative, osmotic, ER, glucose) by addition of O-linked N-acetylglucosamine at serine and threonine residues of nucleocytoplasmic proteins, which is directly associated with autophagy and with regulation of endoplasmic reticulum stress and cell viability [32,33]. Glycosylation can alter protein function and has a key role in many important biological processes including cell adhesion, migration, interactions with the cell matrix, immune surveillance, cell signaling and cellular metabolism; altered glycosylation in prostate cancer in particular, might modify some or all of these processes [34,35,36]. More specifically, *Itkonen et al.*, showed that UDP-GlcNAc is activated in PC-3 prostate cancer cells and can act as a modulator of prostate cancer growth, which reinforces the notion that these often overlooked protein modifications have the potential to improve risk stratification and therapeutic strategies in patients with prostate cancer [37].

Finally, our results agree with those, recently reported, suggesting the beneficial effects of oleuropein against cancer, ageing, neurodegeneration, diabetes and diseases implying autophagy dysfunction [38,39]. Further understanding of the molecular mechanisms which are activated in response to different stimuli is therefore

important towards the pharmaceutical development of novel treatment methods for prostate cancer.

Concluding remarks: The results of the present study provide provocative evidence that oleuropein is a powerful sensitizer of doxorubicin-induced killing of prostate cancer. Because it is a bioactive natural product which is anyway obtained by food consumption, it is regarded as a very promising alternative, not only for the lack of toxicity when administered in a co-treatment scheme with doxorubicin, but also for providing alleviation of the side-effects of DXR in cancer patients.

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