Effects of the combination of camptothecin and doxorubicin or etoposide on rat glioma cells and camptothecin-resistant variants

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Summary From the rat C6 glioma cell line in culture, we selected camptothecin-resistant variants by growth in the presence of increasing amounts of this drug (C6CPT10, C6CPT50 and C6CPT100, growing respectively with 10, 50 and 100 ng ml\(^{-1}\) camptothecin). The degree of resistance to camptothecin ranged between 15-fold (C6CPT10) and 30-fold (C6CPT50 and C6CPT100). The C6CPT10 cell line presented a collateral sensitivity to etoposide (3.6-fold), while the C6CPT50 and C6CPT100 cell lines were cross-resistant to etoposide (1.8-fold) The resistant lines were characterised by a two-fold reduced content and catalytic activity of topoisomerase I, and C6CPT50 and C6CPT100 presented a significant increase in topoisomerase II content and catalytic activity and a marked overexpression of P-glycoprotein. We explored the cytotoxicity of combinations of camptothecin and doxorubicin or etoposide at several molar ratios, allowing the evaluation of their synergistic or antagonistic effects on cell survival using the median effect principle. The simultaneous combination of camptothecin and doxorubicin or etoposide was additive or antagonistic in C6 cells, slightly synergistic in the C6CPT10 line and never more than additive in the C6CPT50 and C6CPT100 cell lines. The sequential combination of doxorubicin and camptothecin gave additivity in the order camptothecin → doxorubicin and antagonism in the order doxorubicin → camptothecin. Clinical protocols combining a topoisomerase I inhibitor (camptothecin) and a topoisomerase II inhibitor (doxorubicin or etoposide) could be compensated for by the topoisomerase of the other class, or to a qualitative alteration of the topoisomerase amino acid sequence leading to reduced cleavable complex formation in the presence of drugs.

Keywords: topoisomerase inhibitors; drug combinations; camptothecin resistance

The nuclear enzymes DNA topoisomerase I (top 1) and DNA topoisomerase II (top 2) represent important primary targets for the action of anticancer drugs (Liu, 1989). These enzymes are able to induce and relegate DNA strand breaks in order to allow multiple topological modifications of DNA such as relaxation of the supercoiled molecules and decatenation and unknotting of intermingled fragments (Andersen et al, 1994). Relaxation is required when the RNA and DNA polymerases operate, in order to relieve the torsional constraints of intertwined DNA strands during the operations of transcription and replication. Decatenation is required during mitosis, for the segregation of sister chromatides in the daughter cells. Top 1 induces DNA single-strand breaks and is mainly involved in DNA relaxation, whereas top 2 induces DNA double-strand breaks, creating a gap through which another DNA double strand can cross the first one (`strand passing'): therefore, the topological activities of top 2 cannot be ensured by top 1, whereas top 2 is able to provide DNA relaxation (Nittiss, 1994).

Many drugs classes have been shown to interfere with DNA topoisomerases: aminocaridine (amascarine), epipodophyllotoxins (etoposide) and anthracyclines (doxorubicin) interfere with the relaxation reaction of top 2, whereas camptothecin and its water-soluble analogues (irinotecan, topotecan) interfere with the same step of top 1 action. In both cases, the drugs stabilise the topoisomerase–DNA complex, which is often called the ‘cleavable complex’, transforming the transient breaks into permanent breaks, which is considered by the cell as a lethal lesion leading to cell death (Osheroff et al, 1994; Pommier et al, 1994). Other drugs have been shown to interfere with other catalytic steps of the DNA-topoisomerase action and present some antiproliferative properties, but none of them has yet been approved as an anticancer drug.

One of the mechanisms of resistance to topoisomerase-interfering drugs consists in the alteration of the target itself (Andoh and Okada, 1994; Robert and Larsen, 1998). A frequent feature characterising cell lines which have been rendered resistant to such drugs is the strong decrease in drug-induced cleavable complex formation, as compared to the corresponding sensitive cell lines. This may be due either to a quantitative defect in topoisomerase expression, which has been shown to occur at the mRNA and at the protein levels, or to a qualitative alteration of the topoisomerase amino acid sequence leading to reduced cleavable complex formation in the presence of drugs.

In both cases, it has been postulated that the reduced catalytic activity following the quantitative or qualitative enzyme alteration could be compensated for by the topoisomerase of the other class, since the actions of both enzymes are partially overlapping (Tan et al, 1989). Indeed, an increase in top 2 activity has been shown to occur in cell lines resistant to camptothecin, a top 1 inhibitor (Sugimoto et al, 1990), and a collateral sensitivity of camptothecin-resistant cells to doxorubicin has been observed (Oguro et al, 1990).
MATERIALS AND METHODS

Cell culture
The cell lines used in this study originate from a chemically induced rat glioblastoma cell line, C6 (Benda et al, 1968), selected because of its properties to generate orthotopic and heterotopic tumours in Wistar rats. We have selected two variants of this line by continuous in vitro growth in the presence of stepwise increasing concentrations of camptothecin: these variants grow in the continuous presence of 10 (C6\textsubscript{CRT10}), 50 (C6\textsubscript{CRT50}) and 100 (C6\textsubscript{CRT100}) ng of camptothecin per ml of medium, respectively.

Drugs and chemicals
Doxorubicin and etoposide were obtained as chemical formulations from Pharmacia & Upjohn and Novartis, respectively. Camptothecin was purchased from Sigma Chimie (Saint-Quentin-Fallavier, France). Stock solutions were prepared periodically in distilled water or dimethylsulfoxide (for camptothecin) and stored at \(-20^\circ\text{C}\). Chemicals were of the highest commercially available quality. Top 1 and top 2 inhibitors. We wanted to explore this possibility in a cellular model developed and studied in our laboratory and its variants resistant to camptothecin, in order to determine whether clinical applications could be implied from these in vitro studies.

Cytotoxicity
Adequate numbers of cells were seeded in 10 cm\(^2\) Petri dishes and grown for at least one doubling time. Drugs were then added to the culture medium and left for the duration of one doubling time in contact with the cell monolayer. After removal of the drug-containing medium, the cell monolayers were rinsed twice with saline solution and cells were allowed to further grow with fresh medium for the duration of two more doubling times, such that the cells remained in exponential phase of growth during the entire process. Cells were then recovered after trypsinisation and counted in a Coulter Counter ZM (Coultronics, Margency, France). Each cell line was characterised by its IC\(_50\) value to a given drug, i.e. the drug concentration reducing cell number by 50\% as compared to drug-free controls. IC\(_{50}\)\(_s\) were calculated at one dose ratio for sequential combinations. The dose ratios selected corresponded to the ratios of the IC\(_{50}\) of individual drugs.

The data were analysed according to the median effect principle, as described by Chou and Talalay (1984). Combination indices were calculated on a microcomputer using the software of Chou and Chou (1987). Combination index values either lower or higher than 1 indicate synergistic or antagonistic effect, respectively.

Immunoblot analysis of topoisomerase II
Nuclear extracts were prepared from about 1 \times 10\(^7\) exponentially growing cells as already described (Montaudon et al, 1997).

Evaluation of the catalytic activities of topoisomerase II and topoisomerase I
Top 2 activity was evaluated by decatenation of a catenated DNA substrate originating from \(Trypanosoma\) kinetoplasts (kDNA, Topogen) into various relaxed DNA forms as already described (Montaudon et al, 1997). The reaction was initiated by the addition of 0.35 M NaCl nuclear extracts and stopped after a 20-min incubation at 37\(^\circ\text{C}\) by adding a denaturing solution. The samples were then electrophoresed on a 1\% agarose gel for 45 min at 80 V. A positive control for the reaction was a decatenated kDNA marker (Topogen). DNA was visualised under ultraviolet light and the various DNA forms were quantified by densitometric scanning. The catalytic activity of topoisomerase II was evaluated as the amount of ng kDNA decatenated per mg protein in the nuclear extract.
Top 1 activity was determined using the DNA relaxation assay previously described by Liu and Miller (1981) with some modifications. Different concentrations of 0.35 M NaCl nuclear extracts and 1 μg of pBKS IKKS supercoiled DNA were mixed on ice in a final volume of 20 μl. The reaction mixtures were incubated at 37°C for 30 min and the reaction stopped on ice. The samples were then submitted to electrophoresis in 1% agarose gel at 80 V for 1 hour. DNA was visualised under UV light and quantified by densitometric scanning. The catalytic activity of topoisomerase I was evaluated as the amount of ng DNA relaxed per ng protein in the nuclear extract.

**Immunoblot blot analysis of P-glycoprotein**

Electrophoresis of whole cell lysates (400 μg proteins per lane) and protein transfer on Immobilon-P membranes were carried out as already described (Huet et al, 1992). The blots were incubated with C219 monoclonal antibody at 4°C overnight. The membranes were then incubated with alkaline phosphatase-conjugated rabbit anti-mouse immunoglobulins and developed using BCIP substrate. Spots were quantified as already mentioned.

**Evaluation of DNA damage in C6 cells**

Analysis of drug-induced DNA strand breaks was performed using the filter alkaline elution technique (Kohn, 1991) as already described (Montaudon et al, 1997). The cells were labelled with [methyl-3H]thymidine (0.1 mCi ml⁻¹, Amersham Pharmacia Biotech), then treated with different drug concentrations for 2 h at 37°C. Cell pellets were resuspended and layered onto polycarbonate membranes of 25 mm diameter and 2 mm pore size (Nucleapore, purchased from Schumacher-DMF, Gonesse, France). The cells were lysed in situ for 1 h in the presence or absence of 0.5 mg ml⁻¹ proteinase K (Roche, Meylan, France), and the DNA on the filter was eluted with tetrapropylammonium hydroxide (Kodak, purchased from Touzart et Quentin-en-Yvelines, France), pH 12.5. Elution was carried out at 0.044 ml min⁻¹ for a total of 15 h. The radioactivity on the filters and collected fractions was determined in a liquid scintillator. It was then possible to calculate the fraction of DNA retained on the filter versus the elution time. The slopes of each elution profile were used to determine the elution rate constant for a given drug concentration. The radiation-induced elution rate constants had previously generated standard curves (Montaudon et al, 1997) allowing the determination of the radiation equivalence of each drug concentration. Results of drug-induced DNA damage were expressed as equivalent radiation dose in Gray (Gy).

**HPLC evaluation of camptothecin cellular accumulation**

Evaluation of camptothecin concentration in cells was performed by HPLC as already described (Rivory and Robert, 1994). Cell extracts were obtained in methanol/acetonitrile (50/50, v/v) containing 1% HCl. Separation was carried out on a C-18 reversed-phase column (Nova-Pak, Radial Pak, Waters, Saint-Quentin-en-Yvelines, France) with a mobile phase consisting of a mixture of acetonitrile and 0.075 M ammonium acetate buffer, pH 6.0, containing 5 mM tetrabutylammonium phosphate (Pic-A, Waters). This mobile phase was delivered isocratically at a flow rate of 1.5 ml min⁻¹ with a Spectra Systems P4000 XR pump (Thermo Quest, Les Ulis France). Fluorometric detection was carried out with excitation and emission wavelengths set at 355 and 515 nm respectively, using the Spectra Systems FL 3000 (Thermo Quest) detector. Peaks were quantified by reference to a standard calibration curve obtained by spiking known amounts of drugs in untreated cell extracts, thanks to the PC1000 software (Thermo Quest).

**RESULTS**

**Characterisation of camptothecin-resistant cell lines**

We selected the three C6 sublines, C6_{CPT50} and C6_{CPT100} by continuous exposure to stepwise increasing concentrations of camptothecin up to 10 ng ml⁻¹ (C6_{CPT50}), 50 ng ml⁻¹ (C6_{CPT100}) and 100 ng ml⁻¹ (C6_{CPT100}), these concentrations being then maintained in the cell culture medium. Table 1 presents the IC₅₀ of several anticancer drugs in these lines. The C6_{CPT50} line is characterised by a 15-fold resistance to camptothecin and a collateral sensitivity to etoposide (3.6-fold) without any change in doxorubicin cytotoxicity and a slight resistance to vincristine. The C6_{CPT100} and C6_{CPT10} cell lines are characterised by a 30-fold resistance to camptothecin and a slight cross resistance to both top 2 inhibitors and to vincristine.

Top 1, top 2α and P-glycoprotein were quantified by immunoblotting of cellular extracts (Figure 1). There was a significant 50% decrease in top 1 amount in C6_{CPT10} and a 70% decrease in C6_{CPT50} and C6_{CPT100} cells. In contrast, there was a 2-fold increase in top 2α level in C6_{CPT50} and C6_{CPT100} cells and no significant change in its level in C6_{CPT10} cells.

Catalytic activities of top 1 and top 2 were studied in C6, C6_{CPT10} and C6_{CPT50} cells (Table 2). There was a decrease in top 1 relaxation activity in both camptothecin-resistant cell lines as compared to the C6 cell line, and a significant increase in top 2 decatenation activity occurring only in the C6_{CPT50} line.

Camptothecin accumulation was evaluated in C6, C6_{CPT10} and C6_{CPT50} cells by HPLC using fluorescence detection. After 4-hour incubations at the concentration of 50 μM, no significant difference in drug accumulation was observed between the 3 cell lines (0.242 ± 0.20 nmole per 10⁶ cells in wild-type cells versus 0.222 ± 0.22 in C6_{CPT10} cells and 0.206 ± 0.20 in C6_{CPT50} cells).

**Cytotoxicity of combinations of camptothecin and top 2 inhibitors**

The simultaneous use of a top 1 and a top 2 inhibitor at a fixed molar ratio over a large range of concentrations allowed the evaluation of their possible synergistic or antagonistic effects on cell survival of the various C6 cell lines. Plotting log (S/F/SF) as a function of log (dose) gave in all cases a linear regression coefficient

| IC₅₀ of the anticancer drugs tested (nM) | Camptothecin | Doxorubicin | Etoposide | Vincristine |
|---------------------------------------|--------------|-------------|-----------|------------|
| C6                                   | 61.1 ± 11.2  | 21.2 ± 2.1  | 207 ± 21  | 3.7 ± 0.7  |
| C6_{CPT10}                           | 890 ± 31     | 20.9 ± 7.4  | 56.9 ± 3.4| 10.6 ± 1.8 |
| C6_{CPT50}                           | 1596 ± 116   | 94.5 ± 5.7  | 366 ± 26  | 31.4 ± 2.4 |
| C6_{CPT100}                          | 1866 ± 72    | 88.4 ± 16   | 334 ± 13  | 25.5 ± 1.4 |

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British Journal of Cancer (2001) 85(7), 1077–1083

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Figure 1 Western blots of the nuclear extracts (for top 1 and top 2) and of total cell lysates (for P-glycoprotein) of C6 cells and camptothecin-resistant variants. Extracts were submitted to electrophoresis and blotting as described in Material and Methods. Revelation was obtained after incubation with specific primary antibodies followed by antibody detection.

Table 2 Catalytic activities of topoisomerases I and II in C6 cells and camptothecin-resistant variants.

|          | Topoisomerase I | Topoisomerase II |
|----------|----------------|-----------------|
| C6       | 7.03 ± 2.38    | 65.1 ± 2.3      |
| C6CPT10  | 2.56 ± 1.02*   | 70.5 ± 2.3*     |
| C6CPT50  | 1.03 ± 0.24**  | 93.4 ± 1.8***   |

Results are expressed as ng DNA relaxed per ng protein (top 1) or as ng DNA decatenated per µg protein (top 2). Significance as follows: *: P < 0.05; **: P < 0.01; ***: P < 0.001.

Figure 2 Analysis of combined effects of camptothecin and doxorubicin administered simultaneously as 24-h exposures to C6 cells and camptothecin-resistant variants. The dose ratios used for cytotoxicity experiments are indicated on the graphs. Using the Chou and Talalay (1984) median-effect principle revealed, at the IC50s of the combinations, synergism (combination index < 1), additivity (combination index not significantly different from 1) or antagonism (combination index > 1) as indicated. Note that the ordinate scales are different for each graph.
0.95, which justifies the use of the median-effect principle for the evaluation of synergistic or antagonistic interactions.

Simultaneous exposures to camptothecin and doxorubicin appeared additive or antagonistic in the C6 line (Figure 2), with the antagonistic effect increasing as a function of the increase in the proportion of camptothecin in the mixture. This association was clearly synergistic in the C6CPT10 line but was never more than additive in the C6CPT50 and the C6CPT100 cell lines.

Simultaneous exposures to camptothecin and etoposide were strictly additive in the C6 cell line (Figure 3). In the C6CPT10 cell line, the combination was slightly but consistently synergistic but additivity only was obtained in the C6CPT50 and C6CPT100 cell lines. In all cell lines, sequential exposures to camptothecin and doxorubicin appeared additive when camptothecin was given first and antagonistic when doxorubicin was given first (Figure 4).

**DNA damage induced by camptothecin and etoposide in the C6 cell line**

The DNA damage induced by etoposide, alone and in combination with camptothecin, was evaluated in the C6 cell line as a complement, at the molecular level, of the cytotoxicity experiments. With etoposide at 1 and 2 µM, there was a concentration-dependent increase in DNA damage, from 0.94 ± 0.17 to 1.62 ± 0.15 Gy equivalents. The combination of etoposide and camptothecin, both used at 1 µM concentration, induced DNA damage equivalent to that obtained with etoposide at 2 µM (1.53 ± 0.06 Gy equivalents), corresponding well with the additive cytotoxicity observed in this cell line at equimolar concentrations.

**DISCUSSION**

The cell lines we selected for resistance to camptothecin display in fact several alterations contributing to a complex phenotype of cross-resistance. All of them display a reduction in top 1 content and catalytic activity, a feature relatively frequent in cell lines selected for resistance to top 1 inhibitors (Andoh and Okada, 1994). These cell lines present in addition 2 alterations which appear related to top 2 – interfering drugs: an increase in the level of top 2α and increased catalytic activity of top 2, generally considered as leading to an increased sensitivity to drugs such as etoposide and doxorubicin (Robert and Larsen, 1998), and an overexpression of P-glycoprotein, which in contrast contributes to resistance to the same drugs (Endicott and Ling, 1989).

The development of these 2 opposite mechanisms probably involves 2 independent phenomena. On the one hand, P-glycoprotein has been consistently found to be overexpressed in response to various kinds of stress, such as irradiation (Hill et al, 1990) or cytotoxic compounds not even expelled by this pump (Chaudhary and Roninson, 1993). In our camptothecin-resistant cell lines, there is a slight increase in P-glycoprotein expression at the first level of selection (probably responsible for the 3-fold resistance of this cell line to vincristine) and an important increase in this expression in the most resistant lines, C6CPT10 and C6CPT100. Despite this overexpression, the C6CPT10 line presents no reduced accumulation of camptothecin after 4 hours of incubation with this drug, confirming that camptothecin is not a substrate for P-glycoprotein (Chen et al, 1991).

On the other hand, the increase in top 2α level observed in the most resistant cell lines, as well as the increase in catalytic activity also detected in all resistant lines, is a feature that has been already
observed in cell lines selected with top 1 inhibitors (Sugimoto et al, 1990). It has been interpreted as a regulatory mechanism for compensating the top 1 down-regulation in such cell lines. This slight, but significant increase in top 2 activity is likely to be responsible for the significant hypersensitivity of the C6CPT100 cell line to etoposide and the unchanged sensitivity of this line to doxorubicin. In this line, the level of expression of P-glycoprotein would be too low to exert an important effect on the efflux of these drugs, in addition to the fact that etoposide is a relatively poor substrate for P-glycoprotein. In the C6CPT100 and C6CPT1000, the P-glycoprotein-mediated drug efflux would be enough to counteract the collateral intrinsic sensitivity to etoposide and doxorubicin due to top 2 overexpression.

Such complex drug-resistant phenotypes, implying concomitant alterations of several determinants of drug activity, is frequently observed during the selection process of resistant cells in vitro (‘multifactorial’ resistance, Rabier et al, 1991) and renders relatively complex the interpretation of cytotoxicity data obtained during drug combinations. In our cell models, the combination of a top 1 inhibitor (camptothecin) and a top 2 inhibitor (etoposide or doxorubicin) is not more than additive in wild-type cells. A slight, but significant synergistic effect was observed in the C6CPT10 cell line when doxorubicin or etoposide were combined with camptothecin, using several drug ratios corresponding to the ratio of the respective IC50s of the drugs in the different cell lines. This synergistic effect can be interpreted as a consequence of the increased top 2 activity in this cell line. In contrast, in the C6CPT100 and the C6CPT1000 lines, this synergy was lost, as P-glycoprotein prevents the top 2 increase to sensitize the cells to doxorubicin or etoposide.

Several studies have compared the effect of combinations of top 1 and top 2 inhibitors in different types of cell lines. Kaufmann (1991) observed an important antagonism exerted by camptothecin on the cytotoxicity of etoposide in HL60 cells. Most of the subsequent studies found at best an additive effect between camptothecin (or a camptothecin analogue such as SN-38 or topotecan) and etoposide or doxorubicin (Bertrand et al, 1992; Stahl et al, 1997), but a schedule-dependency was sometimes observed: when the top 1 inhibitor was administered before the top 2 inhibitor, a synergy could be observed (Masumoto et al, 1995; Bonner and Kozelsky, 1996). In the present study, such combinations appeared synergistic only in the C6CPT10 cell line, the only one presenting simultaneously no major change in top 2 content and activity, and a quite moderate P-glycoprotein expression. In the other cell lines, combinations of camptothecin and doxorubicin or etoposide appeared generally additive. In the C6 cell line, combinations of camptothecin and doxorubicin were even antagonistic whereas combinations of camptothecin and etoposide were additive. It must be kept in mind that doxorubicin is an intercalator acting at the level of the cleavable complex, whereas etoposide does not directly interfere with DNA. The steric hindrance resulting from the concomitant presence of doxorubicin and camptothecin on the same DNA sites might explain why their combinations are antagonistic.

This in vitro evaluation of combinations of a top 1 and a top 2 inhibitor does not allow to predict what would be the results of such combinations in vivo or in the clinical setting. However, it suggests that combining top 1 and top 2 inhibitors might be more interesting in documented situations of resistance to top 1 inhibitors not accompanied by P-glycoprotein overexpression. The narrow therapeutic window offered for such combinations may,
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ACKNOWLEDGEMENTS

This work was supported by grants from the Ligue Nationale contre le Cancer, Comités de la Charente et de la Charente-Maritime.

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