



SHORT COMMUNICATION

Differential Sensitivities of Recombinant Human Topoisomerase II α and β to Various Classes of Topoisomerase II-interacting Agents

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ABSTRACT. A series of topoisomerase-interacting antitumour agents were tested for their ability to differentially inhibit the catalytic activity of either topoisomerase (TOPO) II α or β , as judged by a DNA decatenation assay. The α form, relative to the β isoform, proved 1 to 3 times more sensitive to nonintercalating complex-stabilizing TOPO II-interacting agents (etoposide and derivatives) and up to 18 times more sensitive to non-complex-stabilizing inhibitors of TOPO II ((\pm)-1,2-bis(3,5-dioxopiperazinyl-1-yl)propane [ICRF 159] and *meso*-2,3-bis(3,5-dioxopiperazine-1-yl)butane [ICRF 193]). However, the β form of the enzyme appeared 1 to 3 times more sensitive to intercalating TOPO II-interacting agents (daunorubicin, aclarubicin and mitoxantrone). A possible implication of these data are that tumours preferentially expressing either the α or the β isoform may be differentially responsive to various classes of TOPO II-interacting agents. *BIOCHEM PHARMACOL* 56;4:503–507, 1998. © 1998 Elsevier Science Inc.

KEY WORDS. topoisomerase II α ; topoisomerase II β ; catalytic activity; antitumour agents; DNA-decatenation; topoisomerase II-interacting agents

TOPO II \dagger is an important nuclear enzyme controlling DNA topology through catalysis of a transient breakage of double-stranded DNA in an ATP-dependent fashion, allowing for the passage of double-stranded DNA followed by resealing of the DNA [1]. Relaxation of DNA supercoils by TOPO II plays a major role in DNA replication and transcription, and TOPO II plays a critical role in chromosome condensation and separation during mitosis and in the attachment of DNA loops to the nuclear matrix and chromosomal scaffold [1]. Two isoforms of TOPO II, namely α and β , have been characterised [2], and the corresponding human genes cloned [3–6] and expressed [7, 8] in yeast. Both isoforms seem to be functionally equivalent and both complement defective TOPO II mutants in yeast [9, 10]. However, whilst the β isoform is expressed throughout the cell cycle, expression of the α isoform is strictly cellular proliferation-dependent [11]. During mitosis, TOPO II α appears completely bound to the mitotic chromatin, while isoform β diffuses into the cytosol [12], suggesting a role for TOPO II α in chromosome disentanglement. However, the specific localisation of the isoforms during interphase seems controversial [12, 13]. Nevertheless, both isoforms can be

overexpressed in human tumours [13], with a higher proportion of cells in individual tumours showing an elevation in TOPO II β than in α . These data, combined with the fact that exposure of cells to etoposide [12, 14] or to ICRF 187 [14] resulted in a comparable diminution of salt-extractable TOPO II α and β , suggest that either isoform could be targeted by TOPO-interacting drugs.

TOPO II is the target of clinically useful anticancer drugs which either stabilize the cleavable complex, such as the nonintercalating agent etoposide and its derivatives and the DNA intercalating anthracyclines, or those which do not stabilize this complex, such as the catalytic inhibitor ICRF 159 and derivatives [15]. In spite of preliminary reports [2, 8, 10, 16], the question of the relative sensitivity of each isoform of the human enzyme toward TOPO-interacting agents remains open. If, indeed, there is any preferential inhibition of either isoform by specific antitumour agents, this information could be of value in designing optimal drug combinations for different tumours. Therefore, we systematically compared the effects of a series of TOPO-interacting agents on the DNA decatenation activity of the α and β forms of recombinant human TOPO II.

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\dagger Abbreviations: ICRF 159, (\pm)-1,2-bis(3,5-dioxopiperazinyl-1-yl)propane (razoxane); ICRF 187, (+)-1,2-bis(3,5-dioxopiperazinyl-1-yl)propane (dextrazoxane); ICRF 193, *meso*-2,3-bis(3,5-dioxopiperazine-1-yl)butane; and TOPO II, topoisomerase II.

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MATERIALS AND METHODS

Materials

Genistein, actinomycin D and distamycin A, amsacrine, doxorubicin, aclarubicin and daunorubicin as hydrochlorides were purchased from Sigma, netropsin from Boehr-

inger Mannheim, mitoxantrone and bisantrene from Lederle, suramin from RBI, ICRF 187 hydrochloride (dexrazoxane) from Chiron and NPF (4'-demethyl-4p.fluoro-aniline-4-desoxypodophyllotoxin) from Duphar. Teniposide, Top-53 (4'-demethyl-4 β -[2-[N-(2-(N,N'-dimethylamino)ethyl)-N-methylamino]ethyl]-4-desoxypodophyllotoxin) dihydrochloride, GL-331 (4'-demethyl-4-p.nitro-aniline-4-desoxypodophyllotoxin), etopofos, etoposide and ICRF 159 (razoxane) were provided by Pierre Fabre Medicament. Intoplicine mesylate was obtained from Rhone-Poulenc Rorer and ICRF 193 was a gift from Dr. A. Creighton, St. Bartholomew's Hospital Medical College (London).

The plasmid YEpWob6 encoding human TOPO II α under the control of the GAL1 promoter was purchased from Prof. J. C. Wang (Harvard University) and YEp-Top2 β expressed in the yeast JEL1 encoding human topoisomerase II β under the control of the GAL1 promoter was purchased from Dr. C. Austin (University of Newcastle).

Production and Purification of Human Recombinant TOPO II α and β in Yeast

Production was according to published methods [8, 17]. In brief, TOPO expression was induced by switching growth from 2% glucose to 2% galactose for 24 hr. The yeast pellet was weighed, resuspended in 1 to 2 volumes of buffer B₀ and used for purification or stored at -70° . Buffer B: 50 mM Tris pH 7.7, 1 mM EDTA, 1 mM EGTA, 10% glycerol, 20 mM KF, 1 mM dithiothreitol, 0.1 mM paratitrophenylphosphate, 0.1 mM β -glycerophosphate, 1 mM benzamidine, 200 μ g/mL of aminoethylbenzenesulfonyl fluoride, 20 μ g/mL of aprotinin, 5 μ g/mL of leupeptin and pepstatin. The suffix to B indicates the KCl concentration added; e.g. B₂₀₀ signifies B plus 200 mM KCl.

Published methods were used in purification [7, 8, 17], although an additional step was introduced for TOPO II α . Briefly, the phosphocellulose eluate containing TOPO II α was adjusted to 200 mM KCl with B₀ and applied to a 1-mL heparin column (HiTrap, Pharmacia) preequilibrated with buffer B₂₀₀. Following elution with increasing concentrations of KCl, TOPO II α eluted with 500 mM KCl. Fractions of interest were stored in 50% glycerol at -70° . SDS PAGE was performed according to Laemmli [18], and gels were stained with a Silver Stain Plus kit purchased from Bio-Rad.

Assays for kDNA Decatenation Activity of TOPO II

Eighteen microliters of buffer A (50 mM Tris pH 8.0, 120 mM KCl, 0.5 mM DTT, 0.5 mM ATP, 10 mM MgCl₂) containing 200 ng of kDNA (TopoGen) and 1 U of TOPO II (the amount of enzyme resulting in complete decatenation of 200 ng of kDNA) were added to an Eppendorf tube containing 2 μ L of either vehicle (DMSO) alone or the test drug in vehicle [19]. After a 30-min incubation at 37 $^{\circ}$, the reaction mixture was analysed on a 1% agarose gel, run at 35 mA for 2 hr in Tris-borate-EDTA buffer (89 mM Tris,

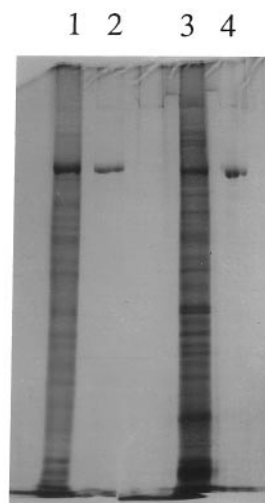


FIG. 1. Assessment of the purity of isolated TOPO II α and β by separation of the proteins on SDS PAGE followed by silver staining. Lane 1: fraction containing TOPO II β before application to the phosphocellulose column; lane 2: TOPO II β eluted from the phosphocellulose column; lane 3: fraction containing TOPO II α before application to the phosphocellulose column; and lane 4: TOPO II α eluted from the heparin column.

89 mM borate, 2 mM EDTA, pH 8.3). Gels stained with ethidium bromide were photographed under UV. Assays to determine inhibition of decatenation were carried out on at least three separate occasions when definite activity was detected, using a range of concentrations lower than the highest tested of 100 μ M.

RESULTS AND DISCUSSION

TOPO II α and II β were purified to homogeneity (Fig. 1). Whilst the purity of the β isoform seemed satisfactory after chromatography on a phosphocellulose column (Fig. 1, lane 2), the α form, though apparently pure on a Coomassie-stained gel, was contaminated by proteins of lower molecular mass detected by silver staining (data not shown). Therefore, a second chromatographic step on a heparin column was added, leading to purification to apparent homogeneity (Fig. 1, lane 4).

To compare the pharmacological profiles of the two enzymes, the minimal amount of each necessary to achieve full decatenation of 200 ng of kDNA was determined, from three independent experiments, as 16 ng (0.1 pmol) and 8 ng (0.05 pmol) per assay for the α and β isoforms respectively.

The first experiments were performed with both the minimal defined enzyme concentration and the slightly lower one, so as to rule out any possible excess of enzyme which might affect the results. As shown in Fig. 2, lowering the minimal amount of enzyme needed for full decatenation from 16 to 14 ng per assay led to incomplete decatenation of the control (lane T3) and to an increased sensitivity to the inhibitor (lanes d and e, shown here for aclarubicin). The consistency of the results obtained led to our using

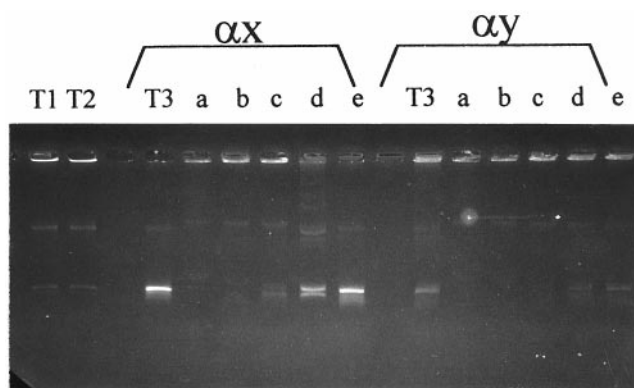


FIG. 2. Effects of aclarubicin on the catalytic activity of TOPO II α . T1: no enzyme; T2: no enzyme + 10^{-4} M aclarubicin; T3: enzyme without drug; a–e: enzyme plus aclarubicin (a: 1.0×10^{-4} M; b: 3.2×10^{-4} M; c: 1.0×10^{-5} M; d: 3.2×10^{-6} M; e: 1.0×10^{-6} M); αx : 16 ng of enzyme (minimal amount required to obtain the complete decatenation of 200 ng of kDNA/assay); αy : 14 ng of enzyme/assay. Note in each of the T3 lanes that while decatenation is complete with the concentration \times (16 ng) of enzyme, this is not the case with the lower concentration y (14 ng) of enzyme. Moreover, this slight change in amount of enzyme led to a dramatic shift in drug sensitivity (compare lanes d and e in gels αx and αy , respectively). In gel αx , the EC_{50} value derived from this experiment is 5.6×10^{-6} M, which is the log mean between the last active (1.0×10^{-5} M) and the first inactive (3.2×10^{-6} M) concentration.

only the minimal enzyme concentration in the assays detailed below.

The ratio of the EC_{50} values obtained against TOPO II β as opposed to α , abbreviated as the β/α ratio, was used as an indicator of the “preference” of a test product for either the TOPO II α or β isozyme. Therefore, values greater than 1 reflect a preference for the α isozyme.

Epipodophyllotoxin derivatives inhibited DNA decatenation catalysed by TOPO II and, as a class, showed an apparent preference for isoform α versus β with values for the β/α ratio ranging from 1 to 3 (Table 1). Both GL-331, the most potent inhibitor of both TOPO II α and β , with EC_{50} values of 5.6 and 18 μ M respectively, and etoposide showed a stronger inhibition of α over β , with a β/α ratio of 3. Teniposide and Top-53 also showed a preference for TOPO II α with a β/α ratio >1 and >2 , respectively. On the other hand, NPF appeared equipotent against both isozymes. In our hands, the prodrug etopofos was inactive against both isoforms at concentrations up to 100 μ M.

Genistein, an inhibitor of TOPO II which stabilizes the cleavable complex [15], showed equivalent effects on both enzymes. Within the class of DNA-intercalating TOPO II inhibitors [15], notably daunorubicin, aclarubicin and mitoxantrone (Table 1), a preference for β over α is apparent, with the β/α ratio ranging from 0.3 to 1. Amongst those inhibitors of TOPO II which do not stabilize the cleavable complex [15], suramin showed equivalent effects on both enzymes, whereas ICRF 159 and ICRF 193 displayed preferential inhibition of TOPO II α over β , with values of >18 for the β/α ratio. ICRF 187, however, exhibited a more modest value for this ratio (Table 1).

Amongst the dual TOPO I/II inhibitors, the DNA-intercalator intoplicine [15] (like the TOPO II specific intercalators) favored the β form, with a β/α ratio of 0.3, while actinomycin D, another intercalator, and distamycin A, a minor groove-binder, appeared neutral with ratios of 1 (Table 1).

In summary, TOPO II α appears 1 to 3 times more sensitive to the nonintercalating TOPO II-interacting agents tested (epipodophyllotoxin derivatives) and up to 18 times more sensitive to the specific catalytic inhibitors of TOPO II (ICRF 159 and derivatives), as compared to the β form. However, the β form of the enzyme appears 1 to 3 times more sensitive to intercalating agents than the α isoform. This latter fact seems to rule out any potentially unrecognized bias in our assays which might favor activity against the α as compared to the β forms of the enzyme. Moreover, data presented here are consistent with those reported both for the mouse enzyme (pointing to a greater effectiveness of teniposide toward the TOPO II α form [2]) and for the human enzyme using a DNA-cleavage assay: the α form was described as either slightly more sensitive than β to epirubicin and etoposide [16] or as sensitive as the β form to teniposide and etoposide [8]. Amsacrine has been reported to induce more cleavage with TOPO II β [8] and mitoxantrone has displayed a fourfold preference for β [10], in line with the generally higher sensitivity of the β form to intercalating agents that we observed in this study. In a DNA relaxation assay, suramin proved equipotent against both isoforms, as described in our decatenation assays, whilst merbarone showed a 2- to 3-fold selectivity for TOPO II α [10].

These data identify an apparently clear-cut differential sensitivity between TOPO II α and β in terms of decatenation activity from a purely biochemical point of view. However, whether they can be considered in any way predictive of the *in vitro* or *in vivo* relative effectiveness of these drugs against either form of TOPO II remains to be established. Indeed, results at the cellular level appear rather contradictory. For example, Brown *et al.* [20] pointed to a negative correlation between levels of TOPO II β and resistance to etoposide in six lymphoma lines (which proved insignificant), while Houlbrook *et al.* [21] described a correlation among seven breast cancer lines between levels of TOPO II β and sensitivities to etoposide. However, Withoff *et al.* [22] showed in one tumour cell line that resistance to teniposide was associated with a decrease in TOPO II α , but not the II β protein. Interestingly, in yeast strains where either human TOPO II α or β was expressed instead of the native yeast enzyme, cell killing with etoposide (as with doxorubicin and mitoxantrone) was greater in strains expressing the α form, whereas amsacrine produced comparable levels of killing in both strains [10]. Therefore, evidence at the cellular level of any definite specific targeting of TOPO II-interacting drugs is still awaited. Such data may result from studies of “knock-out” cell lines expressing only a single enzyme isoform.

TABLE 1. Effects of a series of TOPO II-interacting agents on the catalytic activity of recombinant TOPO II α and β

Test compound	Inhibition of decatenation						EC ₅₀ Ratio IIβ/IIα	
	For TOPO IIα Concentrations (μM)			For TOPO IIβ Concentrations (μM)				
	Range of first active concentration	Range of last inactive concentration	EC ₅₀	Range of first active concentration	Range of last inactive concentration	EC ₅₀		
(N)								
A. TOPO II cleavable complex stabilizing agents								
Nonintercalating agents								
Etoposide	32–100	10–32	32	100–>100	32–100	100	(4)	3
Teniposide	100–>100	32–100	100	100–>100	32–100	>100	(4)	>1
Etopofos	>100	>100	>100	>100	≥100	>100	(2)	—
GL-331	3.2–10	1–3.2	5.6	32	10	18	(3)	3
Top-53	100–>100	32–100	56	>100	≥100	>100	(3)	>2
NPF	32–100	10–32	56	32–100	10–32	56	(3)	1
Genistein	100–>100	32–100	100	100–>100	32–100	100	(2)	1
Intercalating agents								
Doxorubicin	0.32–1	0.1–0.32	0.56	1.0	0.32	0.56	(4)	1
Daunorubicin	1–3.2	0.32–1	1.8	1.0	0.32	0.56	(3)	0.3
Amsacrine	100–>100	32–100	100	100–>100	32–100	100	(4)	1
Aclarubicin	3.2–10	1–3.2	5.6	3.2–10	1–3.2	1.8	(3)	0.3
Mitoxantrone	1–3.2	0.32–1	1.0	0.32–1	0.1–0.32	0.32	(4)	0.3
Bisantrene	3.2–10	1–3.2	1.8	1–3.2	0.32–1	1.0	(4)	0.6
B. Catalytic Inhibitors								
ICRF 187 (Dexrazoxane)	100	32	56	>100	≥100	>100	(3)	>2
ICRF 159 (Razoxane)	10	3.2	5.6	>100	≥100	>100	(3)	>18
ICRF 193	1–3.2	0.32–1	1.0	32–100	10–32	18	(4)	18
Suramin	10–32	3.2–10	5.6	10–32	3.2–10	5.6	(3)	1
C. Dual Inhibitors of TOPO I & II								
Intoplicine	3.2	1	1.8	1	0.32	0.56	(3)	0.3
Actinomycin	3.2–10	1–3.2	5.6	3.2–10	1–3.2	5.6	(3)	1
Distamycin A	1–3.2	0.32–1	0.56	1–3.2	0.32–1	0.56	(3)	1

EC₅₀: the effective concentration at which 50% of the assays are positive for some visible inhibitory activity. Generally, this was the intermediate concentration between the last inactive concentration and the first active one, calculated as the log mean of these two concentrations. The last inactive concentration is defined as the concentration at which 1/3 or 0/3 assays (i.e. under 50%) are positive for inhibitory activity, and the first active concentration is that concentration at which 2/3 assays (i.e. over 50%) are active in terms of inhibition. The solvent used was DMSO to a final concentration of 1%. N = number of experiments.

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References

1. Roca J, The mechanisms of DNA topoisomerases. *TIBS* **20**: 156–160, 1995.
2. Drake FH, Hoffmann GA, Bartus HF, Mattern MR, Croke ST and Mirabelli CK, Biochemical and pharmacological properties of p170 and p180 forms of topoisomerase II. *Biochemistry* **28**: 8154–8160, 1989.
3. Tsai-Pflugfelder M, Liu LF, Liu AA, Tewey KM, Whang-Peng J, Knutsen T, Huebner K, Croce CM and Wang JC, Cloning and sequencing of cDNA encoding human DNA topoisomerase II and localization of the gene to chromosome region 17q21–22. *Proc Natl Acad Sci USA* **85**: 7177–7181, 1988.
4. Chung, TDY, Drake FH, Tan KB, Per SR, Croke ST and Mirabelli CK, Characterization and immunologic detection of cDNA clones encoding two human DNA topoisomerase II isozymes. *Proc Natl Acad Sci USA* **86**: 9431–9435, 1989.
5. Jenkins JR, Ayton P, Jones T, Davies SL, Simmons DL, Harris AL, Sheer D and Hickson I, Isolation of cDNA clones encoding the β isozyme of human DNA topoisomerase II and localisation of the gene to chromosome 3P24. *Nucleic Acid Res* **20**: 5587–5592, 1992.
6. Austin CA, Sng JH, Patel S and Fischer LM, Novel HeLa topoisomerase II is the II β isoform: Complete coding sequence and homology with other type II topoisomerases. *Biochim Biophys Acta* **1172**: 283–291, 1993.
7. Wasserman RA, Austin CA, Fisher LM and Wang JC, Use of yeast in the study of anticancer drugs targeting DNA topoisomerases: Expression of a functional recombinant human DNA topoisomerase II α in yeast. *Cancer Res* **53**: 3591–3596, 1993.
8. Austin CA, Marsh KL, Wasserman RA, Willmore E, Sayer PJ, Wang JC and Fischer LM, Expression, domain structure, and enzymatic properties of an active recombinant human DNA topoisomerase II β . *J Biol Chem* **270**: 15739–15746, 1995.
9. Jensen S, Redwood CS, Jenkins JR, Andersen AH and Hickson ID, Human DNA topoisomerases α and β can functionally substitute for yeast TOP2 in chromosome segregation and recombination. *Mol Gen Genet* **252**: 79–86, 1996.

10. Meczes EL, Marsh KL, Fisher LM, Rogers MP and Austin CA, Complementation of temperature-sensitive topoisomerase II mutations in *Saccharomyces cerevisiae* by a human Top2 β construct allows the study of topoisomerase II β inhibitors in yeast. *Cancer Chemother Pharmacol* **39**: 367–375, 1997.
11. Woessner RD, Mattern MR, Mirabelli CK, Johnson RK and Drake FH, Proliferation- and cell-dependent differences in expression of the 170 kDa and 180 kDa forms of topoisomerase II in NIH-3T3 cells. *Cell Growth Diff* **2**: 209–214, 1991.
12. Meyer KN, Kjeldsen E, Straub T, Knudsen BR, Hickson ID, Kikuchi A, Kreipe H and Boege F, Cell cycle-coupled relocation of types I and II topoisomerases and modulation of catalytic enzyme activities. *J Cell Biol* **136**: 775–788, 1997.
13. Turley H, Comley M, Houlbrook S, Nozaki A, Hickson ID, Gatter K and Harris AL, The distribution and expression of the two isoforms of DNA topoisomerase II in normal and neoplastic human tissues. *Br J Cancer* **75**: 1340–1346, 1997.
14. Sehested M and Jensen PB, Mapping of DNA topoisomerase II poisons (etoposide, clerocidin) and catalytic inhibitors (aclerubicin, ICRF-187) to four distinct steps in the topoisomerase II catalytic cycle. *Biochem Pharmacol* **51**: 879–886, 1996.
15. Capranico G and Zunino F, Antitumor inhibitors of DNA topoisomerases. *Curr Pharm Design* **1**: 1–14, 1995.
16. Cornaroti M, Tinelli S, Willmore E, Zunino F, Fischer LM, Austin CA and Capranico G, Drug sensitivity and sequence specificity of human recombinant DNA topoisomerases II α (p170) and II β (p180). *Mol Pharmacol* **50**: 1463–1471, 1996.
17. Worland ST and Wang JC, Inducible overexpression and active site mapping of DNA topoisomerase II from the yeast *Saccharomyces cerevisiae*. *J Biol Chem* **264**: 4412–4416, 1989.
18. Laemmli UK, Cleavage of structural proteins during assembly of the head of bacteriophage T₄. *Nature* **227**: 680–688, 1970.
19. Kragh Larsen A, Grondard L, Couprie J, Desoize B, Comoe L, Jardillier J-C and Riou J-F, The antileukemic alkaloid fagaronine is an inhibitor of DNA topoisomerases I and II. *Biochem Pharmacol* **46**: 1403–1412, 1993.
20. Brown GA, McPherson JP, Gu L, Hedley DW, Toso R, Deuchars KL, Freedman MH and Goldenberg GJ, Relationship of DNA topoisomerase II α and β expression to cytotoxicity of antineoplastic agents in human acute lymphoblastic leukemia cell lines. *Cancer Res* **55**: 78–82, 1995.
21. Houlbrook S, Addison CM, Davies SL, Carmichael J, Stratford IJ, Harris AL and Hickson ID, Relationship between expression of topoisomerase II isoforms and intrinsic sensitivity to topoisomerase II inhibitors in breast cell cancer cell lines. *Br J Cancer* **72**: 1454–1461, 1995.
22. Withoff S, de Vries EGE, Keith WN, Nienhuis EF, van de Graaf WTA, Uges DRA and Mulder NH, Differential expression of DNA topoisomerase II α and β in P-gp and MRP-negative VM26, mAMSA and mitoxantrone-resistant sublines of the human SCLC cell line GLC₄. *Br J Cancer* **74**: 1869–1876, 1996.