DNA fragmentation induced by the antimitotic drug estramustine in malignant rat glioma but not in normal brain – suggesting an apoptotic cell death

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Summary Estramustine, a combination of 17β -oestradiol and nor-nitrogen mustard, has been shown to be metabolised and to induce specific antiproliferative effects in malignant glioma, including arrest of glioma cells in the G_2/M phase of the cell cycle, damage to cell membranes and DNA and induction of free oxygen radicals. To evaluate further the effects of estramustine, an *in vivo* rat glioma model using inbred BD-IX rats and the BT4C cell line was set up. In order to detect cells with fragmented DNA, tumour and brain specimens were, following fixation for histological examination, processed for *in situ* end labelling (ISEL) with biotin-labelled nucleotides. Fresh tissue fragments were also used for DNA integrity analysis on agarose gels. It was demonstrated that estramustine induced clusters of ISEL-positive cells and a pronounced typical fragmentation of DNA 0.5-8 h after treatment. In tumours examined 24 or 94 h after estramustine treatment, and in untreated tumours, only occasional single ISEL-positive cells were scattered in the tumour. DNA from normal brain tissue did not display any visible sign of fragmentation. These changes are indicative of programmed cell death induced by estramustine in glioma cells but not in normal brain tissue. Further studies are, however, needed to establish in detail the mechanisms of cell death following treatment with the antimitotic drug estramustine.

Keywords: glioma; estramustine; chemotherapy; apoptosis; DNA

Estramustine (EaM), a fusion molecule between 17β-oestradiol and nor-nitrogen mustard used in the treatment of prostatic carcinoma, has been proposed to be of interest in the management of malignant glioma. EaM has been shown to exert dose-dependent antiproliferative effects in glioma cells in vitro (von Schoultz et al., 1988; Piepmeier et al., 1993) and in vivo (Bergenheim et al., 1993a, 1994). The mechanism of action involves the microtubule system (Hartley-Asp, 1984; Stearns, 1988; Bjermer et al., 1988; Dahllöf et al., 1993) with arrest of glioma cells in the G_2/M phase of the cell cycle (von Schoultz et al., 1988). The activity seems to be independent of the hormone and alkylating moiety of EaM (Murphy et al., 1984; Tew and Sterns, 1987). However, other targets have also been suggested to be of importance in explaining the cytotoxicity, such as effects on cell membrane-coupled events (Henriksson et al., 1990; Engström et al., 1991; Sandström et al., 1994) and DNA (von Schoultz, 1991; Piepmeier et al., 1993), while others emphasise that non-DNA targets such as nuclear protein matrix are of importance (Tew et al., 1983; Hartley-Asp and Kruse, 1986; Pienta and Lehr, 1993). Therefore, we now find it of interest to report that EaM induces signs of a rapid and transient apoptotic cell death in malignant rat glioma cells but not in the normal brain tissue of rats.

Materials and methods

Animals

A rat glioma model using inbred BD- IX rats and the BT4C glioma cell line was set up (Bergenheim *et al.*, 1994). The malignant glioma cells were grown as a monolayer for 1 week before implantation. Cells growing in log phase were harvested and trypsinised before being spun down and diluted in MEM supplemented with 5% BD IX rat serum to give 20 000 cells $5 \mu l^{-1}$. BD IX rats, 8-14 weeks old, were anaesthetised by i.p. administration of 1.8 ml kg^{-1} of a 1:1 mixture of Hypnorm (fluanisonum 10 mg ml⁻¹ and fentany-

lum 0.2 mg ml^{-1}) and Dormicum (midazolam 5 mg ml⁻¹). Twenty-thousand viable cells in 5 µl were transplanted under stereotactic conditions, 3.5 mm to the right of bregma at 4.5 mm depth, in the caudate nucleus. A microsyringe was used (22S Ga needle; Unimetrics, Shorewood, IL, USA), allowing at least 5 min for injection and withdrawal of the needle to prevent cellular reflux and extracerebral spread of tumour cells. The drill hole was closed with bone wax. To ensure cell viability the cell suspension was kept on ice during the implantation procedure and the viability of the cell suspension was continuously controlled by staining with trypan blue. After implantation the rats were fed *ad libitum*.

Estramustine treatment

Estramustine EaM (Kabi Pharmacia, Helsingborg, Sweden) was dissolved in a vehicle of 20 g% ethanol and then mixed with 78 g% castor oil with 2 g of Tween 80 to give the concentration 10 mg ml⁻¹ (Eklöv *et al.*, 1992). EaM was given intraperitoneally, 20 mg kg⁻¹ as a single injection at day 24 after implantation. At that time the tumours have an estimated volume of 160 mm³ and do not contain necrotic areas (Bergenheim et al., 1994). The rats were divided into nine subgroups containing 3-6 animals in each group. At 0.5, 1, 2, 4, 8, 24 and 96 h after treatment with EaM the animals were killed and the tissues were handled as described below. Two control groups were used: one received no treatment and one group was treated with the vehicle alone. The brains were immediately, after decapitation, taken out and the tumours were carefully dissected out under the microscope. Normal and tumour tissues were immediately treated as described below.

In situ end labelling (ISEL)

Several pieces of tumour tissue and non-tumour-containing brain tissue from each animal were fixed and embedded in paraffin, and sections $3-4 \,\mu m$ thick were made according to routine histology procedures. In situ detection of apoptotic cells was visualised as follows according to the protocol by Wijsman *et al.* (1993). After deparaffinisation and rehydra-

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tion, the sections were heated twice in SSC (sodium chloride 17.5 g, sodium citrate 8.8 g l⁻¹ water, pH 7.0; Merck Darmstadt, Germany) at 80°C for 20 min and subsequently washed thoroughly in distilled water. To enable enzymatic incorporation of nucleotides, the sections were digested in 0.5% pepsin in hydrochloric acid (pH 2) for 15 min with gentle shaking in a 37°C water bath. The digestion was stopped by washing several times in tap water and then washed in buffer for 5 min. After drying, the sections were incubated for 1 h at 15°C with buffer containing 0.01 mM dATP, dGTP, dCTP and 0.01 mM biotin dUTP (Boehringer Mannheim, Mannheim, Germany) along with 4 U ml⁻¹ DNA polymerase 1 (Sigma, USA). Endogenous peroxidase was blocked for 5 min in PBS with 0.1% hydrogen peroxide, and the sections were then washed twice for 5 min in PBS (0.1% hydrogen peroxide). The sections were then incubated with avidin dissolved in PBS with 1% BSA (bovine serum albumin) and 0.5% Tween 20 (Boehringer Mannheim, Mannheim, Germany) for 30 min at room temperature before developing with diaminobenzidine. In negative controls, DNA polymerase was excluded from the nucleotide polymerase mix. Normal rat prostate, 3 days after castration, was used as a positive control. At that time a significant number of the epithelial cells display characteristic features of apoptotic cells (Dive and Wyllie, 1993). The number of ISEL-positive cells was quantified in the light microscope expressed as cells per surface area (Table I). Five randomly chosen areas $(5 \times 10^{-3} \text{ cm}^2)$ in each section were counted. Sections were also processed for routine staining with haematoxylin and eosin.

Analysis of DNA integrity

The analysis of DNA integrity has recently been described in detail (Tounekti et al., 1993). Briefly, whole tissue was frozen in liquid nitrogen and ground with a homogeniser. Tissue fragments were suspended in digestion buffer (100 mM sodium chloride, 10 mM Tris-HCl pH 8, 25 mM EDTA pH 8, 0.5% sodium dodecyl sulphate and 0.1 mg ml⁻¹ proteinase K) and incubated at 50°C for 12 h and each sample was treated with DNAse free RNAse (0.5 mg ml^{-1}) for 1 h. The samples were heated to 70°C. Loading buffer (0.25% Orange G, 30 mM EDTA, 15% Ficoll) was added to each sample at a 1:2 (v/v) ratio before loading into the 1.5% agarose gel containing $0.1 \,\mu g \, m l^{-1}$ ethidium bromide. Electrophoresis was carried out in 0.1 M TBE (0.045 M Trisborate, 0.001 M EDTA) at 21 V for 14-16 h on 1.5% agarose gels and viewed by transillumination with UV light and photographed. λ HindIII standard was used as molecular size standard. All chemicals were from Sigma.

Results

Histological examination of the untreated BT4C tumour displayed a polymorphic and dense cellular appearance with features of a gliosarcoma. Mitoses were frequent, and abnormal mitotic figures were also found. The growth behaviour was invasive with a clear tendency for perivascular growth with nests of tumour cells developing in the normal brain.

 Table I
 Quantitative analysis of ISEL (in situ end labelling)-positive cells following estramustine treatment analysed at various time intervals

Time point (h)	Number of ISEL-positive cells $(5 \times 10^{-3} \text{ cm}^2)$ Mean \pm s.e.m. (range)
0	0 (0-0)
0.5	47 ± 7 (30–60)
1	$197 \pm 17 (160 - 230)$
2	231 ± 25 (200-264)
4	523 ± 37 (410-600)
8	453 ± 33 (400-490)
24	3 ± 1 (2-4)
96	0 (0-0)

No signs of inflammatory cells or necrosis were seen within the tumour. At the border of the tumour adjacent to normal tissue mononuclear inflammatory cells were occasionally seen. In order to detect apoptotic cells with fragmented DNA, tissue sections were processed for *in situ* end labelling (ISEL) with biotin-labelled nucleotides. In untreated control tumours occasional single ISEL-positive cells with morphologically fragmented nuclei were seen scattered in the tumour. At 0.5, 1, 2, 4, or 8 h after estramustine treatment clusters of ISEL-positive cells were observed in the tumours (Figure 1). The tumour tissue between these clusters contained only a few labelled cells. The cell clusters varied in size and some contained only a few whereas others contained up to 100 labelled cells. Several of the labelled cells had a fragmented



Figure 1 In situ end-labelled (ISEL) sections from malignant rat glioma 4 h after estramustine administration to the animals: (a) $400 \times \text{magnification}$, (b) $800 \times \text{magnification}$ and (c) non-treated tumour tissue. Note the heavily ISEL-labelled cluster of tumour cells. The normal brain from the same animal and the non-treated tumour tissue were almost without detectable ISEL cells.

nucleus suggesting that they were apoptotic. We could not detect any clear-cut signs of focal tissue necrosis. The number of ISEL-positive cells and number of clusters increased and reached peak values 4-8 h following estramustine treatment. At 24 h only a very few positive cells were seen (Table I). Ninety-four hours after estramustine treatment clusters of ISEL-positive cells were not observed and tumour morphology seemed similar to non-treated controls.

Estramustine caused a typical pronounced fragmentation of DNA in the intracranially situated glioma tumour tissue visible 30 min following the single injection as visualised in the gel analysis (Figure 2). A similar pattern was seen 1, 2, 4 or 8 h after the EaM delivery to the animals, whereas there was no sign of DNA ladder 24 or 96 h later. DNA from the normal brain tissue and untreated tumours did not display any visible sign of fragmentation at all.

Discussion

In the present study, morphological and molecular observations for the first time suggest that estramustine (EaM) induces signs of apoptotic cell death in malignant glioma cells but not in non-tumour-containing brain tissue. The degradation of genomic DNA into low molecular weight DNA (<1000 bp) is considered as a hallmark of apoptosis (see, for example, Dive and Wyllie, 1993; Isaacs, 1994). The electrophoretic DNA analysis demonstrated a marked transient DNA degradation in the glioma tissue from EaMtreated rats in contrast with that from non-treated animals and in normal brain tissue from both treated and non-treated animals. The *in situ* end labelling was used to facilitate and enhance the accuracy of detecting apoptotic cells in the tissues (Wijsman *et al.*, 1993) using the fact that the 3'-hydroxyl ends of the terminal deoxynucleotides in the double-stranded DNA fragments produced during programmed cell death can be labelled with biotinylated deoxynucleotides. Cells undergoing programmed cell death incorporate biotinylated nucleotides into their nuclear DNA and can therefore be identified (Gaurieli *et al.*, 1993; Gorcyzca *et al.*, 1993; Wijsman *et al.*, 1993), however necrotic cells are also stained.

It can thus, of course, be speculated whether the observed clusters of cells with fragmented DNA is caused by apoptotic or necrotic cell death. However, the demonstration of ISELpositive cells with fragmented nuclei suggests that at least some of the cells in the treated tumours die of apoptosis. Moreover, the pattern of DNA degradation with stereotypic nucleosomal-sized fragment is characteristic of apoptotic cell death. The response was also time dependent and disappeared without signs of inflammatory reaction or other signs of necrosis. The rapid onset, i.e. detected within 30 min, may suggest that the induction of apoptosis by EaM is a late event in the signal pathway or that the machinery of programmed death in this particular glioma is already in some way turned on and only needs to be triggered to be enhanced by EaM. However, this must be further studied before any final conclusions can be drawn.

It is of interest to recall earlier studies with EaM which have shown effects that, in the light of current knowledge, can be considered features of the process leading to apoptosis: membrane-coupled events with early influence on ion fluxes (Engström *et al.*, 1991; Sandström *et al.*, 1994) and formation of bleb-like projections observed in glioma cells (von Schoultz *et al.*, 1990; Engström *et al.*, 1991). Bleb formation has been correlated with loss of cell viability (Noseda *et al.*, 1989) and may be related to lipid peroxidation via free oxygen radicals (Noronha-Dutra *et al.*, 1988). Fragmentation of DNA and disturbed DNA synthesis have



Figure 2 DNA integrity analysis with agarose gel electrophoresis according to the description in Materials and methods. Samples of normal brain tissue and estramustine-treated and non-treated control tumours marked at various time intervals were analysed after (0.5, 1, 2, 4, 8, 24 and 96 h) EaM delivery. Note the clearly visible DNA ladder in the EaM-treated brain tumour specimens. (a) Lanes 1-6, 0.5 h treatment with EaM (odd numbers = normal tissue, even numbers = tumour tissue); lanes 7-12 l h treatment with EaM; lanes 13-18, 2 h treatment with EaM; lane 19, λ *Hind*III standard. The arrow shows 2.3 and 2.0 kb fragments; the double arrow shows the 0.5 kb fragment. (b) Lanes 1-6, 4 h treatment with EaM (odd numbers = normal tissue, even numbers = tumour tissue) lanes 7-10, 8 h treatment with EaM; lanes 11-16, 24 h treatment with EaM; lanes 17-20, control, non-treated tumour tissue; lane 21, λ *Hind*III standard. The arrow shows 12.3 and 2.0 kb fragments; the double arrow shows 2.3 and 2.0 kb fragment with EaM (odd numbers = normal tissue, even numbers = tumour tissue); lane 7, λ *Hind*III standard. The arrow shows 2.3 and 2.0 kb fragment, (c) Lanes 1-6, 9 h treatment with EaM (odd numbers = normal tissue, even numbers = tumour tissue); lane 7, λ *Hind*III standard. The arrow shows 2.3 and 2.0 kb fragments; the double arrow shows the 0.5 kb fragment. (d) Lanes 1-6, 7 h treatment with only the vehicle solution (odd numbers = normal tissue, even numbers = tumour tissue); lane 7, λ *Hind*III standard. The arrow shows 2.3 and 2.0 kb fragments; the double arrow shows the 0.5 kb fragment. (d) Lanes 1-6, λ h fragments; the double arrow shows the 0.5 kb fragment. (d) Lanes 1-6, λ h fragments; the double arrow shows the 0.5 kb fragment. (d) Lanes 1-6, λ h hindIII standard. The arrow shows 2.3 and 2.0 kb fragments; the double arrow shows the 0.5 kb fragment. (d) Lanes 1-6, λ h hindIII standard. The arrow shows 2.3 and 2.0 kb fragments; the double arrow shows the 0.5 kb fragment. (d) Lanes 1-6, λ h

also been observed in glioma cells after exposure to EaM (von Schoultz *et al.*, 1991; Piepmeier *et al.*, 1993). The observation of a rapid induction of free oxygen radicals following EaM treatment (Henriksson *et al.*, 1990) must also be emphasised since free radicals are proposed to be involved in the late steps in the pathways leading to apoptosis (Hockenbery *et al.*, 1993).

In conclusion, the antimitotic drug estramustine induced clearly visible early damage of DNA in tumour tissue but not in brain tissue from the same animals. This difference in reaction between glioma and brain tissue is of special interest and must be further analysed. EaM has previously been shown to penetrate the blood-tumour barrier and to accumulate both in human brain tumours (Bergenheim *et al.*, 1993b) and in brain tumours in rats (Bergenheim *et al.*, 1994). In this respect it is also of interest to recall the recently published results that EaM affects microtubule integrity and displays cytotoxic action only in glioma cells but not in astrocytes (Yoshida *et al.*, 1994). Although there is strong evidence that the mechanism of fragmentation of DNA is

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most likely programmed cell death, further studies are needed to evaluate the changes within the tumour cells that result in cell death. However, the results support the observations (von Schoultz *et al.*, 1991; Piepmeier, 1993) that the antimitotic drug estramustine can specifically induce alteration in DNA integrity. This is still, however, in agreement with the earlier proposed main mechanism of EaM's antimitotic action being interaction with the microtubule system and nuclear matrix (Tew, 1983; Murphy *et al.*, 1984; Tew and Stearns, 1987; Pienta and Lehr, 1993). The cell death induced by EaM may primarily be a membrane/cytoskeletal-triggered apoptotic cell death rather than a direct chemical interaction with the DNA.

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