

Growth suppression by transforming growth factor β_1 of human small-cell lung cancer cell lines is associated with expression of the type II receptor

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Summary Nine human small-cell lung cancer cell lines were treated with transforming growth factor β_1 (TGF- β_1). Seven of the cell lines expressed receptors for transforming growth factor β (TGF- β -r) in different combinations between the three human subtypes I, II and III, and two were receptor negative. Growth suppression was induced by TGF- β_1 exclusively in the five cell lines expressing the type II receptor. For the first time growth suppression by TGF- β_1 of a cell line expressing the type II receptor without coexpression of the type I receptor is reported. No effect on growth was observed in two cell lines expressing only type III receptor and in TGF- β -r negative cell lines. In two cell lines expressing all three receptor types, growth suppression was accompanied by morphological changes. To evaluate the possible involvement of the retinoblastoma protein (pRb) in mediating the growth-suppressive effect of TGF- β_1 , the expression of functional pRb, as characterised by nuclear localisation, was examined by immunocytochemistry. Nuclear association of pRb was only seen in two of the five TGF- β_1 -responsive cell lines. These results indicate that in SCLC pRb is not required for mediation of TGF- β_1 -induced growth suppression.

The transforming growth factor betas (TGF- β) constitute a family of polypeptides which have been shown to be multi-functional regulators of basic cellular functions such as proliferation, differentiation, cell adhesion and interactions with extracellular matrix (Massagué, 1990; Roberts & Sporn, 1990; Moses, 1992).

Five different isoforms of TGF- β have been described, of which a wide range of human cells express TGF- β_1 , TGF- β_2 , and TGF- β_3 (Derynck *et al.*, 1985, 1988; de Martin *et al.*, 1987; ten Dijke *et al.*, 1988). TGF- β binds to cell membrane-bound receptors. Three types (TGF- β -rI, -II and -III) are expressed by a variety of human cells, both normal and malignant (for review see Massagué *et al.*, 1992). TGF- β -rI and -II are glycoproteins and their recent cloning revealed an intracellular serine/threonine kinase domain (Lin *et al.*, 1992; Ebner *et al.*, 1993), which is believed to play a role in the initial signal transduction. It has been proposed that TGF- β -rI and -II signal through formation of a heterodimeric complex (Wrana *et al.*, 1992). TGF- β -rIII, which is a proteoglycan (betaglycan), has also been cloned (López-Casillas *et al.*, 1991; Wang *et al.*, 1991) and apparently has no direct role in signal transduction, but may act as a capacitor for the signalling receptors. TGF- β_1 exerts a growth-suppressive effect on a wide range of normal (Massagué, 1990) and malignant human cells, including ovarian carcinoma (Berchuck *et al.*, 1992), mammary carcinoma (Knabbe *et al.*, 1987; Arteaga *et al.*, 1988), endometrial carcinoma (Boyd & Kaufman, 1990), colon carcinoma (Wu *et al.*, 1992), prostate carcinoma (Wilding *et al.*, 1989) and gastric carcinoma (Yanagihara & Tsumuraya, 1992). Only one small-cell lung cancer (SCLC) cell line has previously been reported to be growth inhibited by TGF- β (Lagadec *et al.*, 1991).

The intracellular signalling pathway of TGF- β is still obscure. The protein product of the retinoblastoma gene (pRb) has been proposed to be involved in the pathway mediating TGF- β_1 growth inhibition *in vitro* in mink lung cells (Laiho *et al.*, 1990a) and in human keratinocytes (Pietenpol *et al.*,

1990). In contrast, it has been shown that in a mammary carcinoma cell line pRb is not an obligatory component of this pathway (Ong *et al.*, 1991). Retinoblastoma protein has properties of a cell cycle regulatory factor in that its phosphorylation state varies through the cell cycle, with a highly phosphorylated form predominating in S and G₂/M, and an underphosphorylated form predominating in G₁. Aberrant, non-functional, protein products of mutated Rb genes have been characterised and shown to have lost the ability to become hyperphosphorylated and to associate with nuclear structures (Szekely *et al.*, 1991; Templeton *et al.*, 1991).

Recently, the expressions of TGF- β -r types I, II and III were examined by chemical cross-linking to TGF- β_1 in a panel of 21 human small-cell lung cancer cell lines (SCLCs) in our laboratory (Damstrup *et al.*, 1993). Different combinations of the three receptor types were expressed in seven cell lines. Using the Northern blotting technique the expression of TGF- β_1 , TGF- β_2 and TGF- β_3 mRNAs was examined and coexpression of TGF- β -r and TGF- β mRNA was found in six cell lines. Another six cell lines were found only to express TGF- β mRNA.

The expression of Rb mRNA and pRb in the panel of SCLC cell lines has been reported previously (Rygaard *et al.*, 1990). Using Western blotting, it was shown that in five of the 17 cell lines phosphorylated pRb was expressed, indicating expression of a functional retinoblastoma protein. In this study the functionality of the TGF β receptors in SCLC and the possible involvement of pRb in mediating the effect of TGF- β_1 were investigated. We characterised the effect on the growth of the seven TGF- β -r-positive and of two TGF- β -r negative cell lines of continuous treatment with exogenous TGF- β_1 , and evaluated the subcellular localisation of pRb by immunocytochemistry. The results showed that TGF- β_1 induced growth suppression in five SCLC cell lines, and that the growth suppression was found exclusively in the cell lines expressing TGF- β -r type II. We report for the first time TGF- β_1 -induced growth suppression of a cell line expressing the type II receptor without coexpression of the type I receptor. Retinoblastoma protein was not an obligatory component of the pathway mediating the effect of TGF- β_1 , since only two of the growth-suppressed cell lines also expressed functional pRb, as characterised by nuclear localisation and phosphorylation.

In the two cell lines expressing all three types of TGF- β -r, growth suppression by TGF- β_1 was accompanied by morphological changes at the light microscopical level.

Materials and methods

Cell lines

SCLC cell lines were cultured in 150 cm² flasks at 37°C, in an atmosphere of 5% carbon dioxide and 80% humidity in medium containing 10% inactivated (56°C, 30 min) fetal calf serum (FCS) (Flow Laboratories, Irvine, UK) without antibiotics. A total of nine SCLC cell lines established from six patients and characterised as SCLC cell lines in three different laboratories were examined. Three cell lines established at Dartmouth Medical School (Hanover, NH, USA) (DMS 53, DMS 114, DMS 273) (Pettengill *et al.*, 1980; Sorenson *et al.*, 1984) were cultured in Waymouth medium (Gibco, Paisley, UK). Four cell lines established at Groningen Lung Cancer Center (Groningen, The Netherlands) (GLC 3, GLC 14, GLC 16, GLC 19) (De Leij *et al.*, 1985; Berendsen *et al.*, 1988) were cultured in RPMI-1640 (Gibco), and two cell lines established in our laboratory (CPH 54A, CPH 54B) (Engelholm *et al.*, 1986) were grown in EMEM (Eagle's minimum essential medium) (Gibco). AKR-2B, a mouse fibroblast cell line that has previously been reported to be TGF- β -r positive, (Tucker *et al.*, 1984), was cultured in EMEM supplemented with 10% FCS, and used in the displacement assay described below. AKR-2B was kindly provided by Professor H.L. Moses (Vanderbilt University, TN, USA). The cells were passaged twice a week. Cells growing as monolayer cultures (CPH 54A, CPH 54B, DMS 53, DMS 114, DMS 273) were passaged with trypsin. Cells growing as floating aggregates (GLC 3, GLC 14, GLC 16, GLC 19) were allowed to sediment before replacing the medium. All cell lines were routinely checked for, and found to be free of, mycoplasma infection.

Growth factor

Porcine TGF- β_1 was purchased from British Biotechnology (Oxford, UK) and/or was a gift from Bristol-Myers Squibb (Pharmaceutical Research Institute, Seattle, WA, USA). One microgram of dried TGF- β_1 was reconstituted in 0.5 ml of 4 mM hydrochloric acid containing 2 mg ml⁻¹ BSA, and stored at +4°C. Fresh TGF- β_1 stock solution was tested for the ability to displace ¹²⁵I-labelled TGF- β_1 in AKR-2B cells using a radioreceptor assay as described elsewhere (Damstrup *et al.*, 1993). For growth assays, solutions of TGF- β_1 in culture medium containing 10% inactivated (56°C, 30 min) FCS without antibiotics were made immediately before each experiment.

Growth assay

Exponentially growing cells were harvested as described above, resuspended in PBS and centrifuged at 275 g for 5 min. A single-cell suspension was obtained by mechanical disaggregation. Cells were counted in a haemocytometer and viability was evaluated by trypan blue exclusion. Viable cells were seeded in 35 mm six-well tissue dishes (Costar, Cambridge, USA), 5–20 × 10⁴ in 3 ml of culture medium per well. Cells growing as monolayer cultures were allowed to attach for 24 h. Culture medium was removed and medium containing TGF- β_1 was added (designated day 0). Loosely adhering monolayer cultures were supplemented with TGF- β_1 -containing medium (day 0). Cells growing as floating aggregates were seeded directly in TGF- β_1 -containing medium (day 0). The cells were treated with TGF- β_1 in concentrations corresponding to absence of ligand (0 pM), the average K_D for TGF- β_1 binding to the TGF- β receptors (20 pM) (Damstrup *et al.*, 1993), saturation of the receptors (100 pM) and excess of ligand (250 pM). Fresh medium containing TGF- β_1 was added every 48 h. Growth data were based on repeated (approximately 48 h) harvest and assaying of triplicate wells of each of the four TGF- β_1 concentrations. Samples of cells growing strictly as monolayer cultures (CPH 54A and CPH 54B) were assayed for total protein (see below). All other cells were assayed for DNA content (see below). Each experiment was extended until the plateau phase of the growth

curve for the control (TGF- β_1 0 pM) was reached. All experiments were reproduced at least twice.

Protein determination

Culture medium was removed from the six-well dishes and the cells were washed twice with cold phosphate-buffered saline (PBS). The cells were solubilised in solubilisation buffer [128 mM sodium chloride, 0.25 mM EDTA, 0.5 mM Tris pH 7.5 and 1% (v/v) Triton X-100]. Protein concentration was determined using the BCA protein kit (Pierce Europe, Oud Beijerland, The Netherlands) (Smith *et al.*, 1985).

DNA determination

Medium containing cells was sampled and the plates were further harvested by trypsinisation. The cells from each well were pelleted by centrifugation (1,200 g, 10 min) and homogenised by ultrasonication in 1.0–2.0 ml of fluorimetry buffer (2.0 M sodium chloride, 10 mM Tris, 5 mM EDTA, pH 7.4). The DNA content of 20 μ l of homogenate plus 2.0 ml of fluorimetry buffer containing 0.1 μ g ml⁻¹ bisbenzimidazole (Hoechst dye no. 33258) was determined by fluorimetry (Labarca & Paigen, 1980) with an excitation wavelength of 365 nm and an emission wavelength of 460 nm (Hofer fluorimeter TKO 100).

Immunocytochemistry

Exponentially growing cells were harvested by trypsinisation and seeded on eight-well slide glasses (Flow). After 2 days the cells were fixed with 4% paraformaldehyde–PBS (10 min), washed with PBS (5 min), permeabilised with methanol (1 min) and rewashed with PBS (15 min). The cells were blocked with 0.1% BSA–PBS, and incubated with monoclonal anti-pRb antibody, PMG3-245 (PharMingen, La Jolla, CA, USA) (40 min, 37°C) in concentrations of 0, 5 and 10 μ g ml⁻¹. The cells were washed with 0.1% BSA–PBS (5 min × 3) and incubated with FITC-coupled rabbit anti-mouse immunoglobulin antibody (Dakopatts, Glostrup, Denmark), 1:30 in blocking buffer [25 mM Tris, 125 mM sodium chloride 0.1% (v/v) Tween-20, 4% BSA, 10 mM sodium azide], with 0.01 mg ml⁻¹ Evans blue (30 min, 37°C). Finally the cells were washed with PBS (5 min × 3). Examination and photography were performed with a Leitz Aristoplan microscope equipped with appropriate filters.

Statistics

The total content of protein or DNA was expressed relative to the content at day 0 to obtain a relative increment in cell number. Growth curves were constructed by plotting the relative increment as a function of time. Each growth curve was parameterised using quadratic regression on the log-transformed increments. The estimated coefficients of treated and control cells were compared using multivariate statistics and the significance level expressed as a *P*-value. Values below 0.01 were regarded as significant.

Results

Growth suppression

The results of the TGF- β_1 treatment and the immunocytochemical investigation of pRb localisation are summarised in Table I together with previous characteristics of the examined cell lines. TGF- β_1 treatment resulted in growth suppression exclusively in the cell lines expressing TGF- β -r type II: CPH 54A and CPH 54B expressing all three types of receptors, GLC 16 and GLC 19 expressing receptor types I and II and DMS 273 expressing receptor types II and III. The growth of the two cell lines expressing only type III receptor, DMS 114 and GLC 3, and the two TGF- β -r-

Table I TGF- β receptor expression, pRb expression and phosphorylation level, nuclear localization of pRb and *in vitro* growth suppression by TGF- β_1 in nine SCLC cells

Cell line	TGF- β receptor ^a			pRb ^b	Phosphorylation level ^b	Nuclear pRb	TGF- β_1 growth suppression
	I	II	III				
CPH 54A	+	+	+	+	P	+	+
CPH 54B	+	+	+	+	P	+	+
GLC 16	+	+	-	-	c	c	+
GLC 19	+	+	-	-	c	c	+
DMS 273	-	+	+	+	NP	-	+
DMS 114	-	-	+	+	P	+	-
GLC 3	-	-	+	NT	c	c	-
DMS 53	-	-	-	+	P	+	-
GLC 14	-	-	-	-	c	c	-

^aData on TGF- β receptor expression from Damstrup *et al.* (1993). ^bData on pRb expression and phosphorylation level from Rygaard *et al.* (1990). ^cNot tested because of lack of pRb expression. -, negative; +, TGF- β receptor, expression of pRb, nuclear localisation of pRb detected, or growth suppressed by TGF- β_1 ; P, phosphorylated pRb; NP, underphosphorylated pRb; NT, not tested.

negative cell lines DMS 53 and GLC 14, was unaffected by TGF- β_1 treatment. The *P*-values for comparisons between the individual growth curves obtained after treatment with the four different concentrations of TGF- β_1 are given in Table II. These results demonstrate the significance (*P*-value below 0.01) of the above-mentioned TGF- β_1 growth suppression. The results also show that growth suppression was dose dependent within the applied range of TGF- β_1 concentrations in the cell lines GLC 16, GLC 19 and DMS 273. The growth curves of DMS 273 are shown in Figure 1. It appears that half-maximal growth suppression was obtained with 20 pM TGF- β_1 , and maximal growth suppression with 100 pM, whereas excess of ligand (250 pM) did not result in further growth suppression. The significant difference between the control and 250 pM growth curves for the receptor-negative cell line DMS 53 (Table II) reflects an apparent growth inhibition by TGF- β_1 , occurring very late in the experiment, i.e. after 200 h.

Morphological changes

The two SCLC cell lines CPH 54A and CPH 54B expressed all three receptor types (Table I), and normally have a monolayer growth morphology. In these cell lines, the TGF- β_1 -induced growth suppression was accompanied by changes in morphology, assessed by light microscopy. The cells began to aggregate, pile up and detach. The changes were not dose dependent within the applied range of TGF- β_1 concentrations. Figure 2 illustrates the changes in CPH 54A. These changes are representative of what was seen in both CPH 54A and CPH 54B. In none of the other cell lines examined did TGF- β_1 induce apparent morphological changes.

Nuclear localisation of pRb

The five cell lines previously reported to express pRb (Table I) were examined by immunocytochemistry. In four of the cell lines, CPH 54A, CPH 54B, DMS 114 and DMS 53, intense nuclear staining indicating nuclear localisation of pRb was seen in the majority of cells and a weak nuclear staining was seen in a small number of cells. In contrast, DMS 273 lacked nuclear staining, and preparations of this cell line without Evans blue cytoplasmic counterstaining showed a weak nuclear staining (data not shown). This indicated expression of a non-functional pRb. The pRb in DMS 273 also lacked the ability to become phosphorylated (Table I). Figure 3 shows representative fields of the slides of CPH 54A, DMS 114 and DMS 273.

Discussion

In a previous study we reported that different combinations of the three TGF- β -r types were expressed by a panel of 21 SCLC cell lines (Damstrup *et al.*, 1993). These findings enabled us to investigate a possible functional diversity of the receptor types, and in the present study we examined the effect of TGF- β_1 on the growth of seven TGF- β -r-expressing and two TGF- β -r-negative SCLC cell lines. TGF- β_1 induced growth suppression in two cell lines expressing all three receptor types, two cell lines expressing types I and II receptors and one cell line expressing receptor types II and III. The growth of two cell lines expressing only the type III receptor and the two receptor-negative cell lines were not suppressed by TGF- β_1 (Table I). These results showed that

Table II Estimated *P*-values for comparisons of growth curves obtained after *in vitro* treatment of nine SCLC cell lines with different concentrations of TGF- β_1 (0, 20, 100 and 250 pM). Values below 0.01 are considered significant

Cell line	0/20 pM	0/100 pM	0/250 pM	20/100 pM	20/250 pM	100/250 pM
CPH 54A	0.00	0.00	0.00	0.04	0.13	0.71
CPH 54B	0.00	0.00	0.00	0.02	0.00	0.03
GLC 16	0.00	0.00	0.00	0.79	0.00	0.00
GLC 19	0.55	0.00	0.00	0.00	0.01	0.13
DMS 273	0.00	0.00	0.00	0.00	0.00	0.58
DMS 114	0.91	0.56	0.15	0.44	0.12	0.44
GLC 3	0.93	0.83	1.00	0.98	0.95	0.84
DMS 53	0.09	0.14	0.00	0.98	0.03	0.01
GLC 14	0.62	0.70	0.45	0.92	0.89	0.90

0/20 pM, 0/100 pM and 0/250 pM, comparison of growth curves for control cells (0 pM) and treated cells (20, 100 and 250 pM respectively). 20/100 pM, 20/250 pM and 100/250 pM, comparison growth curves of treated cells.

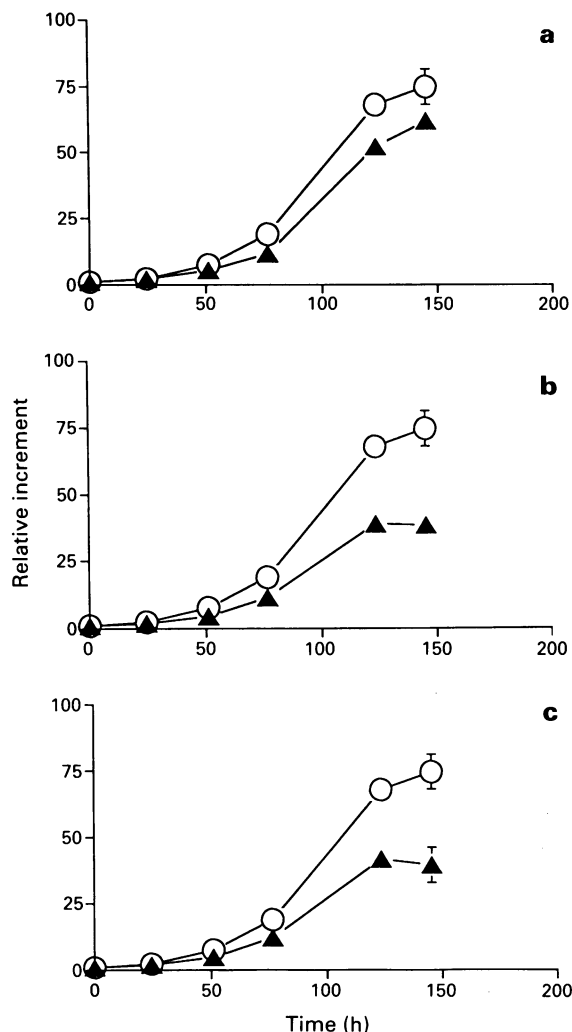


Figure 1 Growth curves for the SCLC cell line DMS 273, obtained after *in vitro* treatment with the following TGF- β_1 concentrations: **a**, 20 pM; **b**, 100 pM; **c**, 250 pM. The relative increment in cell number was plotted as a function of time. Values are mean of triplets \pm s.d. \circ , untreated cells; \blacktriangle , treated with TGF- β_1 .

only in the cell lines expressing TGF- β -r type II did treatment with TGF- β_1 result in growth suppression, and this suggested that in SCLC TGF- β -r type II mediated the growth-suppressive effect of TGF- β_1 . The maximum growth inhibition was approximately the same in all the responding cell lines, and as expected the inhibitory effect emerged later in the slower growing cell lines. This indicated that in SCLC no obligatory coexpression between TGF- β -rII and type I or III was required for mediation of TGF- β_1 growth suppression. However, this panel of cell lines did not include cells expressing TGF- β -rII or TGF- β -rI alone, or TGF- β -rI coexpressed with TGF- β -rIII. Therefore, it could not be established whether these receptor profiles were capable of mediating a response to TGF- β_1 . An apparent growth inhibition of the TGF- β -r-negative cell line DMS 53 did occur very late in the experiment with high TGF- β_1 concentrations. A possible explanation could be that this cell line expressed TGF- β -r in a concentration below the detection limit with chemical cross-linking and radioreceptor assays.

A few reports have elucidated the issue of functional diversity and/or correlation among the TGF- β receptors, and it has become apparent that different mechanisms operate in different model systems. In a recent study, Geiser *et al.* (1992) used a human bladder carcinoma and a human colon adenocarcinoma, which were both resistant to the growth-inhibitory action of TGF- β but responded to TGF- β by producing increased levels of mRNA for extracellular matrix proteins. When fused, a non-tumorigenic hybrid was produced, in which the effect of TGF- β on growth was restored,

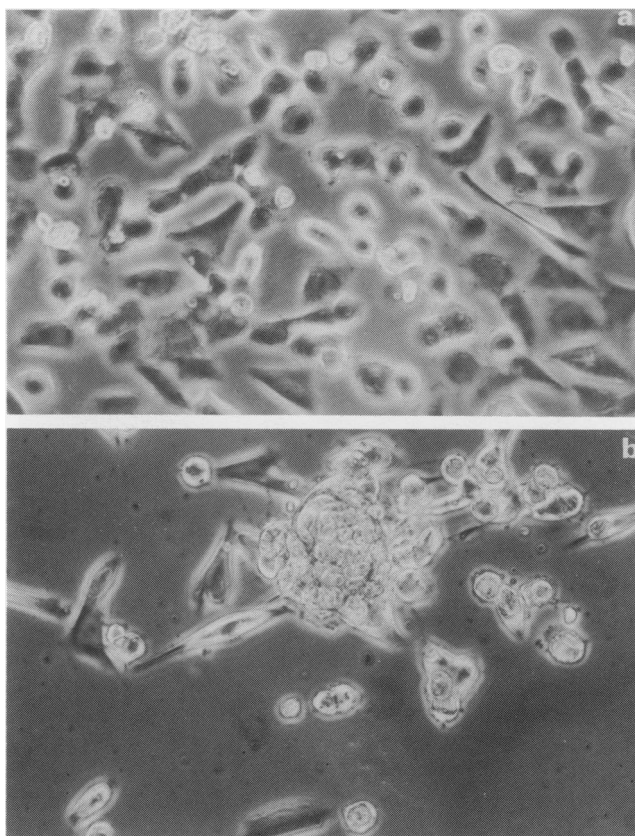


Figure 2 TGF- β_1 -induced morphological changes in the SCLC cell line CPH 54A. Photomicrograph after 140 h of growth. **a**, Without TGF- β_1 . **b**, In the presence of TGF- β_1 , 100 pM. These changes are representative of what was seen in both CPH 54A and CPH 54B, after treatment with either 20 pM, 100 pM or 250 pM TGF- β_1 . Magnification \times 60.

accompanied by a marked increase in TGF- β -rII expression. In another report, transfection of hepatoma cells with the type II receptor cDNA resulted in restoration of growth suppression by TGF- β_1 (Inagaki *et al.*, 1993). Our present results support these findings, and emphasise the role of TGF- β -r type II in mediating growth inhibition, because the SCLC cell line DMS 273 represents the first reported case of growth suppression by TGF- β_1 mediated by the type II receptor without coexpression of the type I receptor (Table I and Figure 1).

A model for functional interrelation between TGF- β -rI and -rII was recently presented (Wrana *et al.*, 1992), based on studies of mutated mink lung epithelial cells (Boyd & Massagué, 1989; Laiho *et al.*, 1990a,b, 1991; Wrana *et al.*, 1992). According to this model TGF- β -rI and TGF- β -rII form a heterodimeric complex for signalling of the various effects of TGF- β , including suppression of growth. As described above, our present results concerning the SCLC cell line DMS 273 do not agree with this heteromeric TGF- β receptor model. However, the possibility exists that DMS 273 expressed TGF- β -rI at a concentration below the detection limit of the chemical cross-linking assay (Damstrup *et al.*, 1993).

TGF- β is supposed to be a central regulator in the normal coordination of growth and differentiation, through autocrine and paracrine mechanisms. A perturbation of the balance between negative and positive growth regulators could lead to an increased proliferative potential, and contribute to the neoplastic phenotype (Moses *et al.*, 1988). We previously reported coexpression of TGF- β and TGF- β -r in six of 21 examined SCLC cell lines (Damstrup *et al.*, 1993). One of the cell lines whose growth was in this study suppressed by TGF- β_1 to the same degree as the others (GLC 16) did not express TGF- β mRNA of any subtype. Two of the coexpressing cell lines expressed only the type III receptor (DMS 114 and GLC 3) and their growth was not suppressed

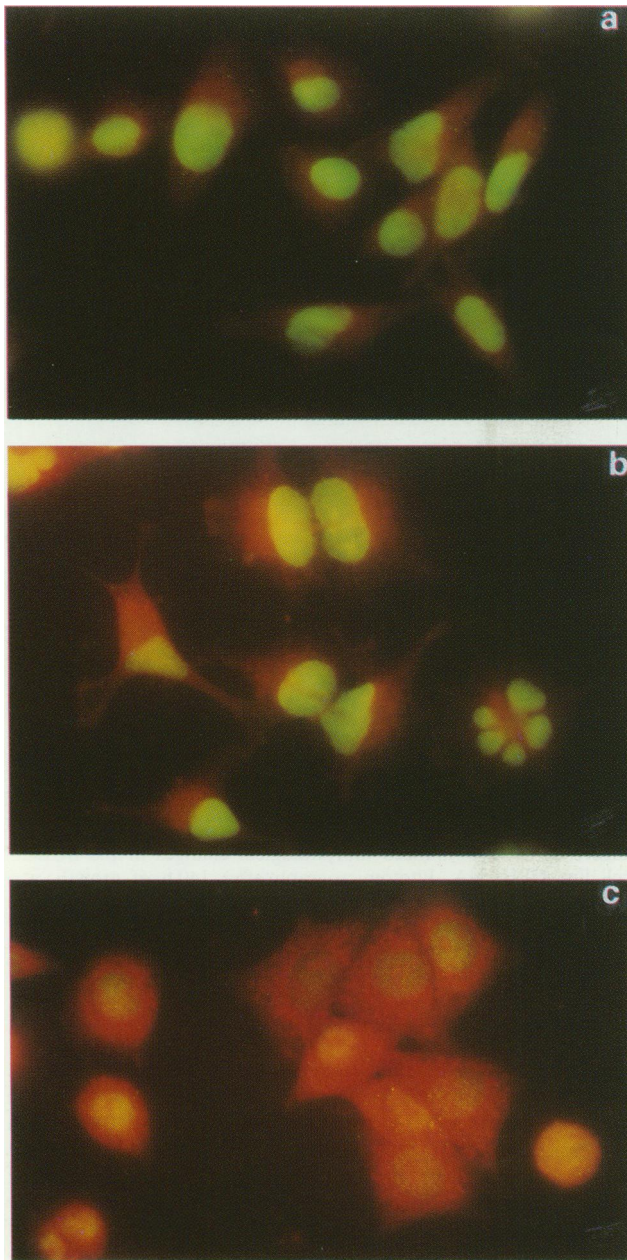


Figure 3 Immunocytochemical detection of Rb protein in SCLC cell lines, using a monoclonal anti-pRb antibody, and cytoplasmic counterstaining with Evans blue. **a**, CPH 54B; **b**, DMS 114; **c**, DMS 273. These fields of view are representative of what was seen in the whole slides. Magnification $\times 800$.

by TGF- β_1 . It is possible that in these three cell lines, together with the cell lines that expressed TGF- β mRNA but no receptors (Damstrup *et al.*, 1993), loss of autocrine growth inhibition by TGF- β contributes to the malignant phenotype. It can, however, be concluded from the present result that this putative mechanism in tumour progression is not a general phenomenon in SCLC.

Induction of altered morphology by TGF- β was previously described (Fanger *et al.*, 1986; Koyasu *et al.*, 1988; Boyd & Kaufman, 1990), and could be expected from its effect on the expression of a wide variety of structural proteins (reviewed in Massagué, 1990). The morphological changes reported varied among different cell types. Two of the cell lines in this study (CPH 54A and CPH 54B) responded to treatment with TGF- β_1 with morphological changes, in addition to growth suppression. These cell lines, which normally grow as monolayer cultures, began to aggregate, pile up and detach in the presence of TGF- β_1 . The nature of these morphological changes is not known but will be the subject of

further investigation. In one experiment we removed TGF- β_1 -containing medium from CPH 54B and added fresh medium to evaluate if the morphological and growth-suppressive responses were reversible. The cells regained the growth rate of the control cells. An increase in number of cells with normal growth morphology was seen, whereas the number of aggregates persisted (data not shown). The morphological response to TGF- β_1 could not be correlated to any pattern of receptor expression, though none of the other cell lines responded with altered morphology as evaluated with light microscopy. CPH 54A and CPH 54B were the only cell lines in this study which expressed all three receptor types (Table I), but given the knowledge that TGF- β -rIII apparently has no direct role in signal transduction, and the finding that the growth suppression was equal in all responsive cell lines, it seemed unlikely that the coexpression of TGF- β -rIII in CPH 54A and CPH 54B, alone should determine this difference.

We also examined the possible involvement of the retinoblastoma protein in mediating the growth inhibition of TGF- β_1 . Previously the expression of Rb mRNA and pRb in our panel of SCLC cell lines was characterised (Rygaard *et al.*, 1990). Western blotting showed expression of pRb in five of the nine cell lines examined in this study (Table I), but in DMS 273 the protein was not phosphorylated. Given the fact that pRb was isolated from exponentially growing cells, and thus representative of all cell cycle phases, the dephosphorylation of the protein in DMS 273 indicated that it was non-functional (Templeton *et al.*, 1991). Using immunocytochemistry, we further investigated the functional state of pRb by evaluating the amount of nuclear localisation of the protein (Table I). Only CPH 54A, CPH 54B, DMS 114 and DMS 53 showed predominantly nuclear staining (Figure 3). Aberrant, non-functional, protein products of mutated Rb genes have been characterised and shown to have lost the ability to become hyperphosphorylated and to associate with nuclear structures (Szekely *et al.*, 1991; Templeton *et al.*, 1991). Our present results showed that an SCLC cell line expressing non-functional pRb (DMS 273) or cell lines without pRb expression (GLC 16 and GLC 19) can be growth inhibited by TGF- β_1 . These findings indicated that in these cell lines there was no correlation between responsiveness to the growth-suppressive effect of TGF- β_1 and expression of functional pRb, and strongly suggest that in SCLC pRb is not an obligatory component in the TGF- β_1 growth suppression pathway. This conclusion is in agreement with the finding that TGF- β could inhibit the growth of mammary carcinoma cell lines in the complete absence of pRb expression (Ong *et al.*, 1991). In contrast, evidence has been provided that pRb plays a part in the TGF- β signalling pathway in mink lung epithelial cells (Laiho *et al.*, 1990b) and in human skin keratinocytes (Pietenpol *et al.*, 1990). It was found that TGF- β_1 inhibition of proliferation involves suppression of *c-myc* expression and is abrogated by pRb-binding viral transforming proteins (Pietenpol *et al.*, 1990). The *c-myc* promoter region mediating this effect, called the TGF- β control element, is also required for pRb suppression of *c-myc* (Pietenpol *et al.*, 1991). The role of pRb in the TGF- β signalling pathway is apparently not definitive, and TGF- β signal transduction therefore probably functions through different mechanisms in different cell types.

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Abbreviations: SCLC, small-cell lung cancer; FCS, fetal calf serum; FITC, fluorescein isothiocyanate; PBS, phosphate-buffered saline; BSA, bovine serum albumin; Rb, retinoblastoma; pRb, retinoblastoma protein; TGF- β , transforming growth factor β ; TGF- β -r, transforming growth factor β receptor; K_D , dissociation constant.

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