Growth inhibition by transforming growth factor β (TGF- β) type I is restored in $TGF- β -resistant hepatoma cells after expression of$ TGF- β receptor type II cDNA

(human hepatoma/transfection/polyclonal antisera)

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ABSTRACT The growth of human hepatoma Hep 3B cells is potently inhibited by TGF- β_1 (ID₅₀ = 0.2 ng/ml, 8 pM). A mutant cell line was derived that was not inhibited in growth by TGF- β_1 at 5 ng/ml (200 pM) and that lacked TGF- β receptor type II $(TGF-BRII)$ gene. Transfection of the cloned $cDNA$ for human TGF- β RII to this mutant cell line restored receptor expression as well as the inhibition in growth by TGF- β_1 . In both wild-type and mutant cells stably transfected with $TGF- β RII cDNA, TGF- β RII coimmunoprecipitated with$ $TGF- β receptor type I in the presence of ligand. These$ experiments provide direct evidence for the role of $TGF- β RII$ in the inhibitory effect of TGF- β on growth and suggest that $TGF- β RII acts by means of a heterometric surface complex with$ TGF- β receptor type I.

TGF- β is a multifunctional protein that mediates cell proliferation, growth inhibition, differentiation, and other functions (1-5). Several transformed cell lines have lost their sensitivity to the inhibition by TGF- β on growth (6, 7), which is a possible mechanism for escape from normal regulation of growth by TGF- β and carcinogenesis. Loss of TGF- β mediated growth regulation could result either from loss of specific receptors for TGF- β or from alterations in the postreceptor signal-transduction pathway (6, 7).

Three cell-surface proteins have been identified through their ability to bind with high affinity and be chemically crosslinked to ¹²⁵I-labeled TGF- β_1 : types I (55 kDa), II (80 kDa), and III (280 kDa) receptors (8-19). These receptors are considered candidates for mediating signal transduction by TGF- β . Experiments using chemically mutated mink lung epithelial cells (20-22) or tumor cell lines (23) that are not inhibited in growth by $TGF-B$ indicated that $TGF-B$ receptors type I (TGF- β RI) and type II (TGF- β RII) might mediate the multiple effects of TGF- β . Such cell lines lack TGF- β RI and TGF- β RII but always retain the TGF- β receptor type III (TGF- β RIII). Several other naturally occurring TGF- β resistant tumor cell lines that lack TGF- β RII also lack TGF- β RI (2, 6, 15), and in other cases they also lack TGF- β RIII (24), but absence of the TGF- β inhibitory effect on growth most consistently correlates with loss of $TGF- β RI$ and/or TGF- β RII. These observations led to the hypothesis that TGF- β -mediated signaling involves both TGF- β RI and TGF- β RII (19, 22, 23, 25). Recently, a cDNA encoding the $TGF- β RII protein has been isolated by using an expression$ cloning strategy (26); its cytoplasmic region, when expressed in bacteria, demonstrated serine/threonine autophosphorylation activity (26). Furthermore, another, broadly expressed TGF- β receptor (type V) has also been reported to have

kinase activity (27), but its involvement in signaling by $TGF- β has not been established.$

To determine whether the TGF- β RII-encoding gene is important in modulating the inhibitory effects of TGF- β_1 on growth, an expression plasmid containing the human TGF- β RII cDNA was transfected into Hep 3B-TR cells (a cell line resistant to TGF- β_1 in that the inhibitory effect on growth is absent and one that lacks the TGF- β RII gene). We show that expression of recombinant TGF- β RII restores the inhibitory action of TGF- β_1 on growth and, in the presence of ligand, forms a noncovalent complex with TGF- β RI. These experiments provide direct evidence that responsiveness to $TGF- $\beta$$ in the transfected cells correlates with ligand binding to a heteromeric receptor complex.

MATERIALS AND METHODS

Cell Culture and Growth-Inhibition Assays. Hep 3B-TR cells were established from a human hepatoma line sensitive to TGF- β (Hep 3B-TS) cells (ATCC no. HB8064) by exposure to low TGF- β_1 concentrations (0.1-1 ng/ml) (R & D Systems, Minneapolis) without any chemical mutagens (32). Cells were routinely passaged in minimal essential medium (MEM)/ penicillin/streptomycin/10% fetal bovine serum (GIBCO/ BRL). Growth was assayed after cells were plated at 5×10^4 cells per 35-mm plastic tissue culture dish in MEM/10% fetal bovine serum. After 24 hr, medium was replaced with the same medium but containing $TGF- β_1 at various concentra$ tions; the medium was then changed every 3 days. At various intervals, cultured cells were treated with 0.02% trypsin and suspended in 1 ml of phosphate-buffered saline $(PBS)/10\%$ calf serum. Absorbance at 660 nm was measured spectrophotometrically. Control experiments demonstrated a linear correlation between Hep 3B-TS cell density and absorbance at 660 nm measured in single-cell suspensions by hemocytometer or by spectrophotometer.

Expression Plasmids and Transfection. A recombinant plasmid was constructed (26) by inserting the \approx 4.5-kbp fulllength cDNA of TGF- β RII into the EcoRI site of the mammalian expression vector pcDNAI/neo (Invitrogen, San Diego). Recombinant plasmids, which expressed both sense and antisense $TGF- β RII genes, were purified by Qiagen-pack$ 500 (Qiagen, Chatsworth, CA). Plasmid DNAs (20 μ g), rendered linear by the Sfi ^I restriction enzyme, were transfected by electroporation at 250 V and 960 μ F capacitance,

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Abbreviations: TGF- β , transforming growth factor β ; TGF- β RI, TGF- β RII, and TGF- β RIII, TGF- β receptor types I, II, and III, respectively; Hep 3B-TS, human hepatoma cell line sensitive to TGF- β ; Hep 3B-TR, Hep 3B hepatoma cell line resistant to TGF- β . §To whom reprint requests should be addressed at: E1552 BST, Liver Transplantation, Department of Surgery, University of Pittsburgh, Terrace and Lothrop Streets, Pittsburgh, PA 15213.

using the Gene Pulser (Bio-Rad). Forty-eight hours after transfection, cells were treated with Geneticin (G418) (400 μ g/ml) (GIBCO/BRL). G418-resistant clones were isolated by limiting dilution and were individually expanded. These clones were screened for expression of $TGF- β RII$ and for reappearance of TGF- β sensitivity by culturing with or without TGF- β_1 , at 0.5 ng/ml.

RNA and DNA Analysis. Total RNA $(30 \ \mu g)$ was prepared from Hep 3B-TS, Hep 3B-TR, and clone 2 (Hep 3B-TR cells stably transfected with TGF- β RII cDNA) by using RNAzol (Biotecx, Houston), and RNA samples were resolved by denaturing glyoxal-agarose gel electrophoresis (28). The separated RNAs were transferred to a Zetabind nylon membrane (Whatman) and were crosslinked with UV light (254 nm). The membrane was hybridized to a ³²P-labeled TGF-BRII cDNA probe according to standard protocols (28) before autoradiography at -70° C. A ³²P-labeled oligonucleotide hybridization probe for human 28S ribosomal RNA (Oncogene Science, Manhasset, NY) was used to compare the amounts of RNA transferred onto the membrane.

For DNA analysis, 10 - μ g samples of DNA prepared from each cell line were digested for 3 hr with either HindIII or EcoRI, electrophoresed in a 1% agarose gel, and subjected to Southern transfer and hybridization with the TGF- β RII cDNA probe, according to standard protocols (28). A human β -actin probe (29) was used to verify the digestion and transfer efficiency of DNAs.

Receptor Affinty-Labeling Assays. Binding and crosslinking of $125I$ -labeled TGF- β_1 to cells grown on six-well trays or 100-mm dishes (Fisher) was as described (26). Crosslinked proteins were resolved by 5-10% linear-gradient SDS/PAGE under reducing conditions and exposed to XAR film (Kodak) at -70° C. To test the sensitivity of TGF- β RI to the reductant dithiothreitol, cells were incubated for 5 min at 37°C in the presence of ¹ mM dithiothreitol before adding 125I-labeled $TGF- β_1 .$

Immunoprecipitation of Receptors with Anti-TGF-8RII Polyclonal Antiserum. Cell-surface $TGF-\beta$ receptors were labeled with ¹²⁵I-labeled TGF- β_1 as described above, except that saturating concentrations of radioligand (0.3 nM) were used to amplify the receptor signal. A polyclonal rabbit antiserum specific for the C-terminal 16-amino acid epitope $(H2-D)$ of the human TGF- β RII $(H.Y.L.,$ unpublished results) was raised. Immunoglobulins (IgGs) were prepared from the crude rabbit serum by ammonium sulfate precipitation followed by DEAE chromatography (30) before use for immunoprecipitation (IP) reactions. Affinity-labeled extracts were diluted to 1-ml final volume of ice-cold IP buffer (1% Triton X-100/0.5% sodium deoxycholate/0.25% SDS/2 mM phenylmethylsulfonyl fluoride). Extracts were precleared by incubation with preimmune serum. Then, the IgG fraction was added at 30 μ g/mI alone or with an equimolar amount of peptide H2-D to inhibit antibody binding to the receptor. Protein A-Sepharose (Sigma) was used to precipitate the immunocomplexes, and the pellets were thoroughly washed in the above IP buffer, dissolved in Laemmli loading buffer, and subjected to 5-10% linear gradient SDS/PAGE before autoradiography.

RESULTS

Growth Response to TGF- β_1 . Growth of Hep 3B-TS cells was effectively inhibited by $TGF-\beta_1$; maximal inhibition of growth was achieved at 0.4 ng/ml (Figs. ¹ A and 2). The Hep 3B-TR cells were not inhibited by TGF- β_1 at concentrations of up to 5 ng/ml (Fig. $1B$).

After transfection with TGF-BRII cDNA, G418-resistant clones were expanded and examined for growth inhibition. Populations of G418-resistant cells that were transfected either with $pcDNAI/neo/TGF- β RII (sense or antisense)$

FIG. 1. Effect of TGF- β_1 on growth of Hep 3B-TS (A), Hep $3B-TR(B)$, and clone $2(C)$ cells. Values were the means of duplicate dishes, and the experiments were repeated at least three times. Concentrations of TGF- β_1 are 0 ng/ml (\bullet), 0.05 ng/ml (\diamond), 0.1 ng/ml (\Box), 0.2 ng/ml (Δ), 0.6 ng/ml (\odot), 1 ng/ml (\triangle), 2 ng/ml (\blacklozenge), and 5 ng/ml (m) (effects of concentrations as high as 5 ng/ml are shown in B).

plasmid showed little, if any, inhibition by TGF- β_1 , even at 2 ng/ml (data not shown). Consequently, individual G418 resistant clones were isolated by limiting dilution after transfection and expanded separately with or without TGF- β_1 at ¹ ng/ml. Two selected clones were inhibited in growth by TGF- β_1 (clone 2, Fig. 1C and clone 24, Fig. 2), although with decreased sensitivity compared with Hep 3B-TS cells. Individual G418-resistant clones expressing the antisense TGF- β RII cDNA were not inhibited by TGF- β_1 (data not shown).

RNA and DNA Analysis of the TGF-BRII-Encoding Gene. The Hep 3B-TS cell line expressed $TGF-BRII$ mRNA (Fig. 3, lane 1), whereas the Hep 3B-TR cells expressed none (Fig. 3, lane 2). Clone 2 (Fig. 3, lane 3) and clone 24 (data not shown)

FIG. 2. Effect of TGF- β_1 on growth of cell clones during a 6-day culture. \circ , Hep 3B-TS cells; \Box , Hep 3B-TR; \blacktriangle , clone 2; and \blacklozenge , clone 24.

did express TGF- β RII message. We did not detect TGF- β RII mRNA expression in any of the four G418 and TGF- β resistant clones tested that were derived after transfection with the sense TGF- β RII vector (data not shown).

Southern analysis showed that the Hep 3B-TS cells contained $TGF-BRII$ genomic sequences, whereas the Hep 3B-TR cells did not (Fig. 4). Transfection of $TGF- β RII cDNA$ into Hep 3B-TR cells resulted in appearance of the recombinant TGF- β RII-specific DNA sequences. With either Hind-III or EcoRI restriction enzymes, Hep 3B-TS cells and clones 2 and 24 showed different patterns on Southern blots. Thus, the TGF- β -sensitive phenotype of clones 2 and 24 was not due to cell contamination by Hep 3B-TS cells. The different Southern patterns in clones 2 and 24 resulted from independent integration events of the TGF- β RII cDNA in the host cell genome, thus proving the clonal origin of these two cell

Chemical Crosslinking of 125 I-Labeled TGF- β_1 to Cell- \mathcal{C}_{new} Crossinial of 12I-1Labeled TGF- ρ_1 to Cell**urface Receptors.** Hep 3B-TS cells express three detectable
CE a kinding experience TCE and $(55, 1.$ D-1. TCE and $(90, 1.00)$ TGF- β -binding proteins, TGF- β RI (55 kDa), TGF- β RII (80 kDa), and TGF- β RIII (280 kDa) (Fig. 5A, lanes 1 and 2). $TGF- β RI and TGF- β RII were more abundantly expressed.$ TGF-,3RI and TGF-j3RII were more abundantly expressed than TGF-pixili. Binding to each of these proteins was saturable, as indicated by their disappearance in the presence
of 1 nM (20-fold excess) unlabeled TGF- β_1 (data not shown). Hep 3B-TR cells expressed no detectable TGF- β RII or TGF- β RII were found, as well as several additional binding proteins of found, as well as several additional binding proteins of μ undetermined origin, two of which (named DDT-R, 50 and 42 μ

FIG. 3. Expression of TGF- β RII mRNA, determined by RNA hybridization. Each lane contained 30 μ g of total RNA prepared from Hep 3B-TS cells (lane 1), Hep 3B-TR cells (lane 2), or clone 2 (lane 3). Equivalent RNA loading and transfer were confirmed by subsequent probing with ^a human 28S ribosomal RNA oligonucleotide. Positions of TGF- β RII (labeled T β R-II) and 28S RNAs are indicated by arrows.

FIG. 4. DNA analysis by Southern hybridization. Ten micrograms of DNA prepared from Hep 3B-TS cells, Hep 3B-TR cells, clone 2, or clone 24 were digested by either HindIII (H) or EcoRl (E) for 3 hr at 37°C before electrophoresis, and the filter was hybridized with the full-length human TGF- β RII cDNA. Control hybridization of the filter with an actin DNA probe (data not shown) revealed equal intensities of bands in all four DNAs, indicating equal loading and transfer of DNA in all lanes.

kDa) were the most abundant species (Fig. 5A, lane 3). These small binding proteins are not TGF- β RI because TGF- β binding to TGF- β RI is abrogated by treatment with dithiothreitol (25) (Fig. 5A, lane 2), whereas dithiothreitol treatment had no effect on the DTT-R proteins (Fig. 5A, lane 4).
Furthermore, binding of ¹²⁵I-labeled TGF- β_1 to cell-surface determore, binding of $\frac{1}{2}$ 125 $\frac{1}{2}$ to cell-surface T_{G} -pRI and T_{G} -pRII in Hep 3B-TS cells was not competed against by 100-fold excess TGF-P2; yet binding to the DTT-R species in Hep 3B-TR was readily competitive and to the same extent as was binding of ^{125}I -labeled TGF- β_1 to TGF- β RIII (data not shown). After transfection of TGF- β RII DNA into Hep 3B-TR cells, expression of TGF-*p*RII was barely detectable at the cell surface by chemical crosslinking of 125 I-labeled TGF- β_1 in either clone 2 (Fig. 5A, lanes 5 and 6) or clone 24 (data not shown), even though the growth of these two transfected clones was inhibited by TGF- β_1 , although less than that of the parental Hep 3B-TS cells.

Immunoprecipitation with Anti-TGF-BRII Polyclonal Se $rum.$ Anti-TGF- β RII polyclonal serum immunoprecipitated labeled TGF- β RII from Hep 3B-TS and clone 2 cells incubated with ¹²⁵I-labeled TGF- β_1 and then subjected to crosslinking (Fig. $5B$, lanes 1 and 5). Immunoprecipitable receptors could not be detected in similarly labeled Hep 3B-TR cells, even after prolonged autoradiography (Fig. $5B$, lane 3). The levels of immunoprecipitated TGF- β RII from clone 2 were at least 10-fold lower than those in Hep 3B-TS cells. Both precipitates contained a labeled species with identical electrophoretic mobility as $TGF- β RI (Fig. 5B, lanes)$ 1 and 5) and characteristic sensitivity to dithiothreitol (data not shown). As a control, peptide H2-D, against which the antiserum was raised, suppressed the immunoprecipitation of both TGF- β RII and TGF- β RI (Fig. 5B, lanes 2 and 6). The efficient coimmunoprecipitation of TGF- β RI with TGF- $\frac{1}{10}$ community community contract $\frac{1}{10}$ and $\frac{1}{10}$ $\sum_{n=1}^{\infty}$ (SCR) with 0.25% SDS during immunoprecipitation, suggests a strong interaction of the two receptors on the cell surface when they bind ligand. Under these conditions very small amounts of labeled TGF-BRIII coimmunoprecipitated, indicating a much weaker interaction between ligand-bound cell-surface TGF- β RII and TGF- β RIII, as we have consistently seen in a variety of other cell lines $(A.M.,$ unpublished work). Because the coimmunoprecipitated receptors resolve on the reducing SDS/PAGE, the receptor complexes are not on the reducing $SDS/PAGE$, the receptor complexes are not interesting and C/SDS before addition linked covalently. Boiling samples in 1% SDS before addition of the anti-TGF- β RII serum abolishes immunoprecipitation

FIG. 5. (A) Chemical crosslinking of ¹²⁵I-labeled TGF- β_1 to Hep 3B-TS, Hep 3B-TR, and transfected clone 2. Hep 3B-TS (lanes ¹ and rep 3B-TR (lanes 3 and 4), clone 2 (lanes 3 and 6) cells were
cubated with 50 pM of ^{125}I -labeled TGF- β_1 and crosslinked with disuccinimidyl suberate before analysis on SDS/PAGE. In lanes 2, 4, and ⁶ cells were treated with ¹ mM dithiothreitol for ⁵ min at 37°C before binding of ¹²⁵I-labeled TGF- β_1 . Autoradiograms were ex-
posed for 12 days except for lanes 3 and 4, which were exposed for p_{out} for p_{out} is except for lanes 3 and 4, which were exposed for days. The different TGF- β receptor subtypes and the additional different β . dithiothreitol (DTT)-resistant (DTT-R) binding proteins in Hep 3B-TR are indicated, as are molecular mass markers (lane 7). (B) Immunoprecipitation with anti-TGF- β RII serum. Hep 3B-TS (lanes 1 and 2), Hep 3B-TR (lanes 3 and 4), and clone 2 (lanes 5 and 6) were \lim_{unit} 2), Hep 3B-TR (lanes 3 and 4), and clone 2 (lanes 3 and 6) were
finity-labeled with 0.3 nM ¹²⁵I-labeled TGF- β_1 . Detergent extracts from these cells were immunoprecipitated with anti-TGF- β RII serum in the presence (lanes 2, 4, and 6) or absence (lanes 1, 3, and 5) of equimolar amounts of immunogenic peptide H2-D. Autoradiography was done for 12 days. The two TGF- β receptor subtypes and molecular mass markers (lane 7) are also indicated.

of TGF- β RI but does not abolish that of TGF- β RII (data not shown). These results indicate that the coimmunoprecipitation of TGF- β RI with TGF- β RII is from interactions between the two receptor types in the presence of $TGF-\beta$, although the possibility that the antiserum recognized native but not denatured TGF- β RI cannot be fully excluded.

DISCUSSION

Characterization of a TGF- β -Resistant Hepatoma Cell Line. Hep 3B-TR cells were established by exposing Hep 3B-TS cells to stepwise increases in TGF- β_1 concentration (0.1-1 ng/ml) without the use of any chemical mutagen. These cells show resistance to the inhibitory action of TGF- β_1 on growth at concentrations as high as 5 ng/ml. The resistant cell line expressed a distinctive pattern of cell-surface $TGF- β -binding$ proteins, as determined by chemical crosslinking of 1251 labeled TGF- β_1 . No apparent band(s) comigrating with TGF- β RII were detectable despite the presence of a faint smear in that region of the gel that resulted from high TGF- β RIII levels. Hep 3B-TR also had decreased levels of cell-surface binding by TGF- β RI and increased TGF- β RIII binding. An additional set of TGF- β -binding proteins was detected in this cell, among which two low-molecular-mass proteins (DTT-R, ⁵⁰ and ⁴² kDa) were the most abundant. We confirmed that these were not identical to TGF- β RI, based on their different electrophoretic mobilities, their different sensitivities to dithiothreitol, and their different binding affinities for 1251 labeled TGF- β_1 in the presence of excess amounts of competitor TGF- β_2 . Specifically, a brief preincubation of cells with 1 mM dithiothreitol abolished the ability of TGF- β RI to bind ¹²⁵I-labeled TGF- β_1 without perturbing binding to TGF- β RII or TGF- β RIII. TGF- β_2 effectively abolished ¹²⁵I-TGF- β_1 binding to these two proteins, similar to its action on TGF- β RIII and characteristically different from that on TGF- β RI and TGF- β RII.

 DNA analysis showed a loss of the TGF- β RII gene in Hep 3B-TR cells and consequent loss of TGF- β RII mRNA. Our data suggested that the loss of the TGF- β RII gene is the mechanism by which this hepatoma cell line lost its sensitivity to the inhibitory actions of $TGF- β_1 on growth. This is$ a different situation, compared with other $TGF- β -resistant$ cells, such as chemically induced lung epithelial cell mutants (22) or naturally occurring solid tumor cell lines (23), which lose responsiveness to TGF- β by acquiring point mutations in TGF- β RII protein (25). G_F -prii protein (25).

inibitory Effect of T GF- β_1 on Growth is Restored After Expression of TGF- β RII in Hep 3B-TR Cells. Here we demonstrate the reestablishment of growth inhibition by TGF- β_1 in Hep 3B-TR cells after transfection with an expression vector for TGF- β RII cDNA. In different G418-resistant clones the expression of TGF- β RII mRNA and the TGF- β RII cell-surface protein correlated perfectly with the response to TGF- β_1 of inhibiting growth. Specifically, after transfection of TGF- β RII cDNA into Hep 3B-TR cells, we established two clones (2 and 24) the growth of which was potently inhibited by TGF- β_1 . RNA analysis showed that clones 2 and 24 expressed TGF- β RII mRNA. Chemical crosslinking of ¹²⁵I-labeled TGF- β_1 followed by immunoprecipitation with antiserum against TGF- β RII showed that the TGF- β RII protein had also reappeared. Most G418-resistant clones derived from transfection with the sense TGF- β RII vector tested negative for both TGF- β RII mRNA expression and inhibition of growth by TGF- β_1 . Similarly, none of the G418-resistant clones isolated after transfection with the antisense TGF- β RII vector responded to TGF- β_1 . We conclude that expression of the $TGF- β RII cDNA in Hep 3B-TR$ cells restored cell-surface receptor levels adequate for the modulation of growth by TGF- β_1 .

The pattern of TGF- β -binding proteins in clones 2 and 24 was virtually indistinguishable from that of the parental Hep 3B-TR cell line, when assayed by chemical crosslinking of ¹²⁵I-labeled TGF- β_1 . The inability to detect TGF- β RII on the cell surface of clones 2 and 24 by this technique is from expression of a low number of receptors, as proven by immunoprecipitation with an antiserum against TGF- β RII. Clones 2 and 24, but not parental Hep 3B-TR cells, exhibited

a surface protein of mobility identical to $TGF- β RII but at$ decreased levels compared with those in Hep 3B-TS cells. The lower levels of TGF- β RII on the cell surface correlate with the decreased sensitivity of the transfected clones to TGF- β_1 compared with the parental Hep 3B-TS cell line. Use of specific antiserum to identify $TGF-\beta$ receptors, as exemplified here for TGF- β RII, greatly improves the sensitivity of detection. Even for cells in which $TGF- β receptors cannot be$ visualized by the conventional binding-crosslinking technique (Fig. 5A), immunoprecipitation allows their clear detection (Fig. SB). This technical advance may prove useful in reevaluating the expression patterns of $TGF- β receptors in$ various cells that, as assayed by binding-crosslinking, appeared to lack such surface molecules. This technique is especially important in cases of $TGF- β -response cells with$ very low cell-surface receptor numbers. Our experiments with clones 2 and 24 reinforce the hypothesis that very low levels of cell-surface receptors are adequate for $TGF- $\beta$$ mediated growth arrest, as has been seen in many other cell types (1, 18, 31).

 $TGF-{\beta}RII$ and $TGF-{\beta}RI$ Form a Cell-Surface Complex with Bound TGF- β . Coimmunoprecipitation of TGF- β RI and TGF-BRII by antiserum against TGF-BRII in Hep 3B-TS cells and clone 2 indicates the formation of a complex between TGF- β RI and TGF- β RII on the cell surface. The facts that TGF- β 1 is a disulfide-linked homodimer and that the coimmunoprecipitated receptors were revealed after chemical crosslinking to TGF- β_1 led to the model of two receptor species held in physical proximity via binding to each of the TGF- β_1 monomers in the dimer. TGF- β RI expression was not detectable by immunoprecipitation with anti-TGF- β RII serum in Hep $3B$ -TR cells that lack TGF- β RII, further supporting the idea that these two receptors are complexed together. Similar results have been reported (25) in mink lung cell mutants transfected with the same recombinant TGF- β RII cDNA. Thus, we conclude that TGF- β RI and TGF- β RII form a complex that may mediate the signal transduced by TGF- β_1 in this cell system. However, that TGF- β RII alone might mediate the signal by TGF- β is not excluded (14). Different types of signals could be generated by homodimers of TGF- β RI or TGF- β RII and heterooligomers of TGF- β RI and TGF- β RII. Hence, different cells might require different types of such intracellular signals for TGF- ,B-mediated inhibition of growth. Additionally, distinct receptor combinations may be used by a single cell type to mediate distinct growth-inhibitory and gene-regulatory effects in response to TGF- β .

The present results highlight the importance of growth regulation by TGF- β in the process of hepatic tumor development. Although Hep 3B-TS is a tumor cell line, it responds to TGF- β and expresses all three receptors for TGF- β . Our experiments clearly show that resistance to TGF- β can be achieved by in vitro selection of this cell type. Addressing the question of tumorigenicity and the differing biology of the parental, resistant, and transfected cell lines now becomes imperative.

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