A Swiss 3T3 Variant Cell Line Resistant to the Effects of Tumor Promoters Cannot Be Transformed by *src*

MUKUND NORI, LAURA K. SHAWVER,† AND MICHAEL J. WEBER*

Department of Microbiology, University of Virginia School of Medicine, Charlottesville, Virginia 22908

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To study the relationship between oncogenesis by v-src and normal cellular signalling pathways, we determined the effects of v-src on 3T3-TNR9 cells, a Swiss 3T3 variant which does not respond mitogenically to tumor promoters such as 12-O-tetradecanoyl-phorbol-13-acetate (TPA). We found that src was unable to transform these variant cells, whether the oncogene was introduced by infection with a murine retrovirus vector or by transfection with plasmid DNA. 3T3-TNR9 cells were not inherently resistant to transformation, since infection with similar recombinant retroviruses containing either v-ras or v-abl did induce transformation. Further analysis of Swiss 3T3 and 3T3-TNR9 cell populations infected with the v-src-containing retrovirus revealed that although the amount of v-src DNA in each was approximately the same, the level of the v-src message and protein and the overall level of phosphotyrosine expressed in the infected variants was much less than in infected parental cells. Cotransfection experiments using separate v-src and neo plasmids revealed a decrease in the number of G418-resistant colonies when transfections of TNR9 cells occurred in the presence of the src-containing plasmid, suggesting a growth inhibitory effect of v-src on 3T3-TNR9 cells, as has also been found for TPA itself. Since v-src cannot transform this variant cell line, which does not respond mitogenically to the protein kinase C agonist TPA, we suggest that src makes use of the protein kinase C pathway as part of its signalling activities.

The *src* oncogene encodes a tyrosine-specific protein kinase, $pp60^{src}$, whose enzymatic activity is thought to be necessary for transformation (12, 24, 38, 42). Cells in which $pp60^{src}$ is expressed exhibit a very large number of tyrosylphosphorylated proteins (13, 34, 38, 49), and this has made it extremely difficult to determine which of these phosphorylations is (are) functionally significant with respect to signal transduction and transformation (reviewed in references 38 and 49). Hence, although the *src* oncogene has been studied in detail for many years, the biochemical steps by which expression of *src* modifies cellular growth control mechanisms are still unclear.

Genetic approaches have often been useful in the analysis of complex biochemical pathways. Since oncogenes appear to be transduced elements of the normal machinery of growth control and subvert that machinery to induce abnormal growth and metabolism, it seemed to us that cellular variants which are altered in their responses to normal mitogenic stimuli might also be altered in their response to some oncogenes. Analysis of such variants could elucidate the signalling pathways used by various oncogenes.

In this study, we examined the effects of v-src on 3T3-TNR9 cells. TNR9 cells were derived from Swiss 3T3 cells on the basis of the inability to respond mitogenically to the tumor promoter 12-O-tetradecanoyl-13-acetate (TPA) (8), which is an analog of the normal signalling intermediate, diacylglycerol (4, 15, 16, 18, 21). 3T3-TNR9 cells contain normal levels of active protein kinase C (PKC) and respond to TPA stimulation like parental 3T3 cells with respect to most of the measured responses, including phosphorylation of p80 (5; but see reference 3), production of arachidonate metabolites, and elevation of glucose transport (9, 10). However, TPA does not induce activation of p42-mitogenactivated protein (MAP) kinase, S6 kinase, ornithine decarboxylase, or DNA synthesis in these cells (9, 19; G. L'Allemain, T. W. Sturgill, and M. J. Weber, submitted for publication). It has been reported recently that PKC is defectively down regulated in 3T3-TNR9 cells in response to long-term treatment with TPA (3). Thus, the defect in 3T3-TNR9 cells appears to reside in some early component of the tumor promoter-diacylglycerol signal transduction pathway.

Infection of 3T3-TNR9 cells with the v-src oncogene did not result in transformation, whereas infection with either of two other oncogenes, v-ras and v-abl, did. This refractoriness to src was not due to inability of the cells to take up src DNA, but rather appeared to be due, at least in part, to v-src-induced growth inhibition of these cells. Apparently, the lesion that renders these cells unable to respond mitogenically to TPA also makes them incapable of being transformed by v-src, implying that v-src activates the PKC signal transduction pathway.

MATERIALS AND METHODS

Cell culture. Swiss 3T3 and 3T3-TNR9 cells were obtained from H. Herschman (8) and grown in Dulbecco modified Eagle medium supplemented with 10% fetal bovine serum (GIBCO Laboratories). Virus-producing cell lines were routinely maintained in the same medium with addition of 750 μ g of G418 per ml. Cell lines that produce viruses containing no oncogenes or v-src were made in our laboratory. Cell lines producing viruses containing v-myc or v-ras were obtained from K. B. Marcu (46; personal communication).

Immunological reagents. EC10, a monoclonal antibody that has a high affinity for v-src and a low affinity for murine c-src, was obtained from S. J. Parsons (39). Antiphosphotyrosine antibodies affinity purified from immune rabbit serum by using Affigel-phosphotyrosine columns were provided by A. Rossomando in our laboratory. Rabbit anti-mouse immunoglobulin was obtained from Jackson Laboratory. Radio-

^{*} Corresponding author.

[†] Present address: Triton Biosciences, Alameda, CA 94501.

active iodinated ¹²⁵I-labeled protein A was purchased from Amersham Corp.

Viruses and plasmids. src or other oncogenes were introduced into cells by either infection or transfection. Infections were performed with murine retrovirus vectors expressing both the oncogene and neo, which served as a selectable marker. The ras- and myc-expressing viruses were harvested from cells obtained from K. B. Marcu (46; personal communication). Abelson murine leukemia virus was obtained from B. Mathey-Prevot and did not carry a neo marker. The src-expressing retrovirus vector was constructed by us as follows. The src-containing plasmid pJD100 (52) was digested with restriction endonucleases *XhoI* and *EcoRI*. The 2.3-kilobase fragment generated by this digest, containing the whole v-src gene, including the splice acceptor site, was isolated. This fragment was inserted into fpGV1 (41) at the SacI-EcoRI sites of the multiple cloning region downstream of the neo gene, thus producing fpGV100, which expresses both src and neo.

Transfections of 3T3 and 3T3-TNR9 cells were performed by cotransfecting a plasmid expressing the oncogene along with plasmid pSV2neo, which allowed selection of transfectants in G418. The plasmids used were pJD100 (*src*) (52), pMC29 (MC29 *gag-myc*) (23), the expression vector fpGV1 (41), and a plasmid containing an ecotropic helper virus, pMOV3, which were all obtained from J. T. Parsons.

Transfection and infection. Transfections of cells were carried out by using the calcium phosphate precipitation method with glycerol shock (51). In cotransfection experiments, the precipitate used contained 1 μ g of test DNA mixed with 100 ng of pSV2neo, the selective marker used in these experiments. Control transfections contained either no DNA or only pSV2neo DNA. After about 3 to 4 weeks of selection with 200 μ g of G418 (Geneticin; GIBCO) per ml, colonies were picked at random for expansion into cell lines and then the dishes were stained for counting.

Virus-producing cell lines were constructed in essentially the same manner, with the following changes. The target cells used were NIH 3T3 cells. In each case, 1.5 μ g of the virus-containing plasmid and 300 ng of pMOV3, the helper virus plasmid, were used to form the precipitates. Cells were selected in medium containing 750 μ g of G418 per ml. Culture supernatants were tested for the presence of appropriate viruses by infection into either Swiss or NIH 3T3 cells as described below.

For infection, culture supernatant from the virus-producing cells was harvested and filtered through Millex-GV filters (Millipore Corp.) and 10 ml was added to the target cell culture with Polybrene at a final concentration of 2 μ g/ml. Cells were cultured until confluent and then split 1:5 in growth medium supplemented with 200 μ g of G418 per ml. Clones were obtained by the limiting dilution method (30).

Genomic DNA analysis. DNAs from various cell lines were extracted from confluent cultures and digested to completion by using at least 2 U of restriction endonuclease SacI per μ g of DNA. Ten micrograms of the digested DNA was loaded in each lane and separated by electrophoresis on a 0.8% agarose gel; 0.1 ng of similarly digested plasmid DNA was electrophoresed on the same gel to serve as a positive control. The DNA was transferred to a nylon membrane (Nytran; Schleicher & Schuell, Inc.) and immobilized, and the introduced v-src was detected by using a nick-translated probe. Hybridization was performed at 42°C in buffer containing 50% formamide. The filter was rinsed free of excess probe, dabbed dry, and sealed in Saran Wrap, and the results were visualized by autoradiography. **Cytoplasmic RNA analysis.** Cells from several confluent 100-mm² tissue culture dishes were harvested, and the cytoplasmic RNAs were extracted by the guanidinium iso-thiocyanate-cesium chloride method (32). Ten micrograms of RNA was loaded into each well and separated by electrophoresis through an agarose-formaldehyde gel (32). The RNA was transferred to a Nytran nylon membrane, immobilized, and hybridized with a DNA probe in buffer containing 50% formamide. The filter was rinsed free of excess probe, dabbed dry, sealed in Saran Wrap, and visualized by autoradiography. The 28S and 18S rRNAs, which were clearly visible under shortwave UV light, were used as internal size markers.

Preparation of DNA probes. The same probe was used to detect the v-*src* DNA that was introduced into the cells by infection or transfection and to detect the RNA expressed from these vectors. Plasmid pJD100, which codes for the complete Rous sarcoma virus genome, was cut with *Eco*RI, and the 3.09-kilobase fragment containing the entire *src* gene was isolated and purified. The fragment was nick translated by using [³²P]dCTP at a specific activity of at least 2×10^8 cpm/µg of DNA for use in hybridizations (32).

Protein analysis. For protein analysis, cell extracts were made in boiling electrophoresis sample buffer (29). Extracts of sister cultures were also made in cold RIPA (50 mM Tris hydrochloride [pH 7.4], 0.15 M NaCl, 1% sodium deoxycholate, 1% Triton X-100, 0.1% sodium dodecyl sulfate) and used to determine the protein concentration by the method of Markwell et al. (33). The extracts were fractionated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and electroblotted onto nitrocellulose (7). Western immunoblots to detect tyrosine-phosphorylated proteins were done as described by Kamps and Sefton (27) by using antiphosphotyrosine antibodies made in our laboratory. Western blots to detect v-src were done in the same way, except that the blocking buffer contained 3% milk and the primary antibody was EC10 (39).

Other methods. Infected and uninfected cells were cultured in soft agar to test their abilities to grow in an anchorage-independent manner (2). The mitogenic responses of various cell populations to different stimuli were analyzed by using a [³H]thymidine incorporation assay (8). The increase in 2-deoxyglucose uptake was measured (48) as a biochemical parameter of transformation.

RESULTS

Infection of Swiss 3T3 and 3T3-TNR9 cells with retroviruses carrying src or other oncogenes. Swiss 3T3 and 3T3-TNR9 cells were infected with recombinant retroviruses carrying the *neo* gene and v-src, v-myc, or v-ras. The infected cell cultures were then transferred into medium containing G418 to select for infected cells expressing neo. In all cases, except for v-src-infected 3T3-TNR9 cells, more than 50% of the cells survived the initial selection process, as determined by counting a sister culture (data not shown), indicating a high efficiency of infection. However, only 30% of the v-src-infected 3T3-TNR9 cells survived, i.e., about half of what was obtained with the other infections. As described below, we suspect that the decreased cell survival in the src-infected cultures were due to src-induced growth inhibition. In all cases, including v-src-infected 3T3-TNR9 cells, there was no evidence of further cell death after two passages in selective medium.

After six passages in selective medium, the Swiss 3T3 cell population infected with the v-src-carrying virus was mor-

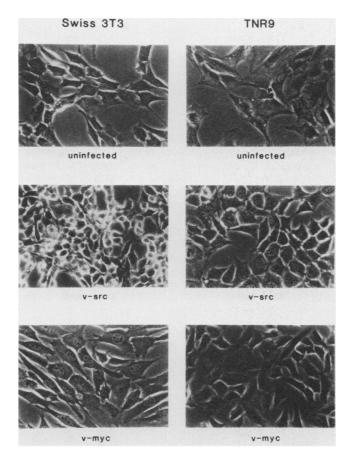


FIG. 1. Morphology of uninfected and recombinant retrovirusinfected Swiss 3T3 and 3T3-TNR9 cells. The recombinant retroviruses used carried either the *neo* gene alone under control of the simian virus 40 promoter or the v-*src* or v-*myc* oncogene under control of the Moloney murine leukemia virus long terminal repeat in a colinear manner.

phologically transformed (Fig. 1) compared with uninfected cells. At the same passage number, the 3T3-TNR9 cell population was indistinguishable from its uninfected control. There was no change in the morphology of these cells, even after more than 11 passages (data not shown). Infection with a retrovirus carrying the v-myc gene did not transform either cell line (Fig. 1).

To test the possibility that 3T3-TNR9 cells were intrinsically resistant to transformation, they were infected with a recombinant retrovirus carrying the v-ras oncogene or with Abelson murine leukemia virus, which expresses the v-abl oncogene. Both of these oncogenes were able to transform the 3T3-TNR9 cells (Fig. 2), indicating that these cells are not completely refractory to transformation.

The populations of v-src-infected Swiss 3T3 and 3T3-TNR9 cells which had been selected in G418 were cloned by limiting dilution. Of the 24 clones isolated from the 3T3v-src population, 23 showed morphological transformation, whereas of the 48 clones isolated from the 3T3-TNR9-v-src population, none were morphologically transformed.

The infected populations and clones were tested for two other parameters of transformation, (i) ability to grow in an anchorage-independent manner in soft agar and (ii) increased 2-deoxyglucose uptake. In all cases, morphological transfor-

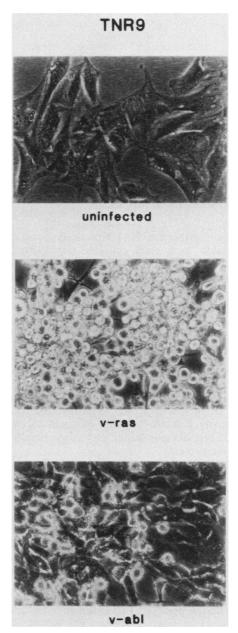


FIG. 2. Morphology of uninfected and v-ras- or v-abl-infected 3T3-TNR9 cells.

mation correlated extremely well with growth in soft agar and 2-deoxyglucose uptake (Table 1).

Molecular characteristics of Swiss 3T3 and 3T3-TNR9 cells infected with v-src. The presence or proviral DNA in each of the infected cell populations was determined by Southern blot analysis of the respective genomic DNAs (Fig. 3A). Relative proviral copy numbers were determined from the intensities of the proviral bands on the autoradiogram. The data show that the intensities of the proviral bands in the Swiss 3T3-v-src (lane 2) and the 3T3-TNR9-v-src (lane 4) populations were comparable, indicating that both cell lines were infected to roughly the same extent and that the failure of TNR9 cells to become transformed was not due to failure to take up virus. DNAs from uninfected Swiss 3T3 cells (lane 1) and 3T3-TNR9 cells (lane 3) exhibited no v-src DNA.

Expression of the v-src gene from these proviruses was

TABLE 1. Correlation of morphological transformation with
ability to grow in an anchorage-independent manner
and increased 2-deoxyglucose uptake

Oncogene	Morphology ^a		Colony formation in soft agar ^b		2-Deoxyglucose uptake ^c	
	3T3	TNR9	3T3	TNR9	3T3	TNR9
v-src	Т	NT	+	_	3.9	1.2
v-ras	Т	Т	+	+	3.5	3.1
v-myc	NT	NT	-	-	1.0	0.8

^a T, Transformed; NT, not transformed.

b +, Colonies formed in soft agar; -, colonies not formed in soft agar.

^c Fold increase in [³H]2-deoxyglucose uptake compared with that of corresponding uninfected cells.

tested in a Northern (RNA) blot analysis of cytoplasmic RNA isolated from each cell line (Fig. 3B). There was considerably less v-src mRNA in the 3T3-TNR9-derived population (lane 4) than in the corresponding one from Swiss 3T3 cells (lane 2), although both cell populations were apparently infected to the same extent (Fig. 3A, lanes 2 and 4).

The abilities of the various infected cell populations to stably express the $pp60^{v-src}$ protein were determined by Western blot analysis. Whole-cell extracts were separated by polyacrylamide gel electrophoresis and electroblotted onto nitrocellulose, and the relative amount of $pp60^{v-src}$ expressed was detected by using $p60^{src}$ -specific monoclonal antibody EC10 (39; Fig. 3C). The infected 3T3 cells expressed easily detectable levels of $pp60^{v-src}$, whereas the infected 3T3-TNR9 population expressed barely detectable levels of the protein.

To determine the in vivo tyrosyl protein kinase activity of the pp 60^{v-src} expressed in these cells, total cell lysates were electrophoresed and analyzed by Western blots using affinity-purified antiphosphotyrosine antibodies (Fig. 3D). Both v-src-infected cell populations exhibited increased levels of tyrosyl-phosphorylated proteins compared with uninfected cells. The extent of this increase correlated with the level of pp 60^{v-src} expressed by the individual cell line (Fig. 3C). Hence, the pp 60^{v-src} expressed in all of these infected cell lines was enzymatically active.

It seemed possible that the extremely low level of enzymatically active pp60^{v-src} expressed in v-src-infected 3T3-TNR9 cells might be a consequence of a genetic defect in the proviruses in these cells, affecting src transcription and/or activity. If, as we suspect (see below), src expression is growth inhibitory in 3T3-TNR9 cells, then the only cells which would grow in a TNR9 population infected with src would be those infected with defective viruses. To test whether the infected but untransformed TNR9-src cells contained functional proviruses, genomic DNA from vsrc-infected cells was digested with restriction endonuclease SacI, which excises the complete provirus. The digested DNA was then transfected into fresh Swiss 3T3 cells, and transfectants were selected by using G418. Clones were picked at random and observed for morphological transformation. The number of G418-resistant transfectants per plate was approximately the same whether DNA was obtained from cells infected with viruses carrying either the neo gene alone or colinear v-src and neo (Table 2). No G418-resistant transfectants were obtained when DNA from uninfected Swiss 3T3 or 3T3-TNR9 cells was used. The percentage of morphologically transformed clones observed in the population transfected with DNA from 3T3-TNR9-v-src cells was

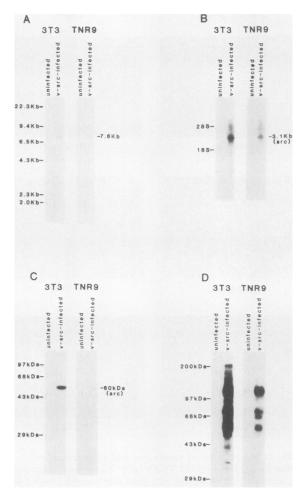


FIG. 3. Analyses of uninfected and v-src-infected Swiss 3T3 and 3T3-TNR9 populations. DNA (10 μ g) from each cell population was digested with SacI and subjected to Southern blot analysis (A); RNA (10 μ g) was subjected to Northern blot analysis (B). The membranes were probed with a piece of ³²P-labeled v-src DNA isolated from plasmid pJD100. Whole-cell extracts from uninfected and v-src-infected Swiss 3T3 and 3T3-TNR9 populations were subjected to Western immunoblot analyses with either an anti-src monoclonal antibody (EC10) to detect the level of p60^{src} expression (C) or affinity-purified polyclonal antiphosphotyrosine antibodies to detect tyrosyl-phosphorylated proteins as a measure of src activity (D). Kb, Kilobases; kDa, kilodaltons.

the same as that seen in the population transfected with DNA from Swiss 3T3-v-src cells. These results indicate that infected 3T3-TNR9 cells, even though they were untransformed, carried proviruses capable of transforming 3T3 cells.

Inhibition of 3T3-TNR9 cell colony formation by src expression. To determine whether src expression was growth inhibitory in 3T3-TNR9 cells, we ascertained whether cotransfection with src would reduce the ability of neo to generate G418-resistant colonies in 3T3 and 3T3-TNR9 cells. We reasoned that if src was indeed growth inhibitory in 3T3-TNR9 cells, its presence should reduce the ability of neo DNA to give rise to G418-resistant colonies in TNR9 cells but not in 3T3 cells. The results of this study are shown in Table 3. Swiss 3T3 and 3T3-TNR9 cells were either transfected with the neo gene alone or cotransfected with neo and v-src. Cotransfection with v-myc served as a control for

DNA source	No. of Neo ^r colonies/plate ^a	No. of transformants/ total (%)	
Swiss 3T3	0	NA ^b	
Swiss 3T3-neo	217	0/22 (0.0)	
Swiss 3T3-src	239	24/35 (68.6)	
3T3-TNR9	0	NA	
3T3-TNR9-neo	184	0/17 (0.0)	
3T3-TNR9-src	206	17/25 (68.0)	

^a Mean of two experiments.

^b NA, Not applicable.

nonspecific effects of cotransfection. Transfectants were selected by using G418, and the number of G418-resistant colonies was determined. In transfections with neo alone. the number of G418-resistant colonies per plate obtained from the 3T3-TNR9 population was the same as that obtained from the Swiss 3T3 population, indicating that both the parental and the variant cells were equally capable of being transfected and expressing the neo gene. Approximately the same numbers of G418-resistant colonies were obtained when Swiss 3T3 cells were cotransfected with neo and v-src, and some of these colonies were morphologically transformed, indicating that src does not have a toxic or growth-inhibitory effect in 3T3 cells. However, there was a drastic reduction in the number of G418-resistant colonies obtained when 3T3-TNR9 cells were similarly cotransfected with neo and v-src. Furthermore, none of these colonies were morphologically transformed nor did they contain any v-src DNA (as determined by Southern blot analysis [data not shown]). This reduction in the number of G418-resistant colonies observed in 3T3-TNR9 cells was not a general phenomenon, since it was not seen when the cells were cotransfected with neo and v-myc, thus suggesting that it was specific to the effects of v-src. Moreover, a transformation-defective mutant of src (dl155 [38]) did not cause a reduction in the number of Neo^r colonies of 3T3-TNR9 cells and did result in the generation of colonies containing src DNA, indicating that these effects are dependent on functionally active src. These data provide strong evidence that v-src expression is growth inhibitory in 3T3-TNR9 cells.

Growth inhibition by TPA. If the phenotypic effects of *src* in TNR9 cells result from the same genetic change which affects their responsiveness to TPA, one would expect that TPA, like *src*, would also be growth inhibitory in these cells. Although the 3T3-TNR9 line was derived on the basis of its mitogenic nonresponsiveness to TPA (8), we found that TPA was, in fact, growth inhibitory in these cells (Table 4),

 TABLE 3. Analysis of cotransfection of neo and v-src into Swiss 3T3 and 3T3-TNR9 cells

	Mean ± SD no. of Neo ^r colonies		
Plasmid content(s)	Swiss 3T3	TNR9	
None (control [no plasmid])	0	0	
neo	120 ± 5	115 ± 20	
neo + v-src	105 ± 18	12 ± 8	
neo + v-myc	130 ± 20	120 ± 13	
neo + v-src-dl155	95 ± 10	128 ± 17	

 TABLE 4. Fold increase in [³H]thymidine incorporation in response to TPA and other mitogens

Cell line	Fold increase in [³ H]thymidine incorporation caused by:			
	ТРА	Epidermal growth factor	Serum	
Swiss 3T3	58	8–10	8-15	
3T3-TNR9	0.2-0.5	4-6	5–12	

reducing the rate of thymidine incorporation into DNA by two- to threefold in the variant while, under identical conditions, stimulating thymidine incorporation in the parent cells. In fact, the original report of Butler-Gralla and Herschman (8), which described 3T3-TNR9 cells, presented evidence that TPA inhibited DNA synthesis in those cells. Thus, these findings strengthen the suggestion that *src* expression is growth inhibitory in 3T3-TNR9 cells and that the inability of the cells to be transformed by *src* is related to their defective responsiveness to TPA.

DISCUSSION

Signalling pathways used by src. To identify which of the cellular signalling systems involved in normal growth control are altered by pp60^{v-src}, we tested the effects of v-src on cellular variants defective in normal mitogenic responses. We have reported previously that src is able to transform 3T3-NR6 cells (47), which lack epidermal growth factor receptors (40), indicating that these receptors are not necessary for src-induced transformation. In this study, we examined the effects of src on 3T3-TNR9, a variant cell line which was isolated on the basis of its inability to respond mitogenically to TPA and which thus appears to be defective in some component of the PKC signalling pathway. We found that v-src was unable to transform these cells, and we interpreted these results as implying that pp60^{v-src} activates the TPA-diacylglycerol signalling pathway as part of the transformation process. This conclusion is consistent with earlier results demonstrating that src expression stimulates phosphatidylinositol turnover (17, 26, 35, 50) and that inhibition of PKC blocks induction of a transformation-related gene by v-src (44). To analyze these results further, it is necessary to review what is currently known about the 3T3-TNR9 variant.

Although the genetic defect in the 3T3-TNR9 variant has not been identified, the cells have been characterized extensively at the physiological and biochemical levels (8-10, 19, 20; L'Allemain et al., submitted). 3T3-TNR9 cells contain normal levels of functional PKC (6), and many of the early responses to TPA which occur in the parental 3T3 cells also occur in the variant, including phosphorylation of p80 and p22 (5; but see reference 3) and increased prostaglandin biosynthesis (10) and glucose transport (9). Furthermore, several genes that are specifically inducible by TPA in Swiss 3T3 cells are also expressed in the same inducible manner in 3T3-TNR9 cells (31). The earliest known TPA-induced response which appears to be defective in TNR9 cells is phosphorylation and activation of MAP kinase, a serinethreonine protein kinase which is activated by tyrosine phosphorylation within 5 to 10 min of treatment of 3T3 cells with TPA or other mitogenic agents (L'Allemain et al., submitted). MAP kinase is present in TNR9 cells and can be activated by platelet-derived growth factor, indicating that the defect in the variant cells resides specifically in the regulation of this activation by TPA. MAP kinase, when phosphorylated and activated, can, in turn, phosphorylate and activate a ribosomal S6 kinase in vitro (45). TPA is unable to induce activation of S6 kinase in 3T3-TNR9 cells (as it can in parental 3T3 cells) (19), consistent with the notion that MAP kinase participates in a kinase cascade that regulates protein synthesis.

Our suggestion that $pp60^{v-src}$ activates a TPA-diacylglycerol signalling pathway is consistent with the fact that transformation by *src* induces increased activation of MAP kinase (A. Rossomando and M. J. Weber, unpublished data) and of S6 kinase (19), as does treatment with TPA. Moreover, these same TPA-induced responses are the ones which are defective in 3T3-TNR9 cells, which cannot be transformed by *src*.

The correlation between mitogenic nonresponsiveness to TPA and resistance to transformation by *src* in 3T3-TNR9 cells is probably not adventitious: Colburn and collaborators have isolated a variant of JB6 epithelial cells which is resistant to TPA-induced transformation (11), and these cells also cannot be transformed by *src* (personal communication).

Transformation by other oncogenes. Although 3T3-TNR9 cells were not transformed by *src*, they were transformed by ras or abl, thus indicating that the specific transformation mechanisms used by src are blocked in TNR9 cells and that there is no intrinsic block to transformation in these cells. The finding that ras can transform these cells is consistent with experiments by Stacey and colleagues (43) which placed ras downstream from src in the transformation process. Presumably, the defect in 3T3-TNR9 cells lies upstream of ras and downstream of src. The ability of 3T3-TNR9 cells to be transformed by *abl* indicates that this tyrosyl protein kinase oncogene uses a signalling pathway different from that of *src*. This agrees with earlier results showing that 3T3-NR6 cells (which lack epidermal growth factor receptors) (40) could be transformed to a tumorigenic state by src but not by abl (47). Interestingly, we have obtained preliminary evidence that the fps oncogene, which is closely related to src, is also unable to transform 3T3-TNR9 cells (Rossomando and Weber, unpublished data). These findings indicate that these variant cells are specifically refractory to transformation by members of the src family of oncogenes.

Mechanism of resistance to transformation. What is the underlying mechanism which prevents the derivation of src-transformed 3T3-TNR9 cells? It is clear that the variant cells are capable of being infected with retrovirus vectors that carry the src gene, and thus the defect is not at the level of uptake and integration of oncogene DNA. However, expression of the src gene was drastically lower in infected 3T3-TNR9 cells than in infected 3T3 cells. This could occur because the variant cells have some mechanism which inhibits expression of the src gene-a result which would be surprising, since no such inhibition occurred when the ras oncogene was introduced by using a very similar vector. An alternative possibility is that src expression is growth inhibitory in 3T3-TNR9 cells. In this case, src serves as a negative selective agent, and the only cells which grow in the culture are the ones with low *src* expression. To test this hypothesis, we cotransfected *src* and *neo* on separate plasmids into 3T3-TNR9 cells and found that src reduced the number of G418-resistant colonies obtained and that the antibioticresistant colonies obtained did not have src DNA. This result can best be explained if the cells which took up src DNA were inhibited in their growth. Presumably, the reason why infection with a retrovirus vector containing src and neo did give rise to cells containing src DNA was because of the higher efficiency of DNA uptake from the retrovirus vector, combined with the fact that the oncogene and the antibiotic resistance gene were linked in the same DNA in the viral vector. Thus, infection with the retroviral vector forced all of the cells to take up viral DNA and the only survivors were ones which expressed the src oncogene poorly. This was likely the basis for the fact that fewer cells were observed in 3T3-TNR9 cultures following infection with the src-carrying virus than in 3T3 cells infected with the same virus or in either culture infected with ras- or myc-containing viruses.

The most direct way to test the growth-inhibitory effects of src in 3T3-TNR9 cells would be to generate a derivative of this variant cell containing a conditionally regulated src gene. One could then, in theory, grow such cells under restrictive conditions and shift them to permissive conditions to observe the phenotypic effects of src expression in this cell background. However, our attempts to perform such experiments by using either src under the control of the mouse mammary tumor virus promoter or by using temperature-conditional src mutants (tsLA24 and tsLA29 [53]) have been unsuccessful. The regulated src gene behaves essentially like the wild type. We suspect that even low levels of leakiness in the regulation are sufficient to cause src-induced growth inhibition. Indeed, we have found substantial tyrosine phosphorylation induced by tsLA29src, even at the restrictive temperature (28; data not shown). However, it is important to point out that the tyrosine kinase activity of p60^{v-src} may not be the sole determinant of its growth inhibitory properties.

Growth inhibition or toxicity is not unique to the src gene; other oncogenes have been shown to be inhibitory or lethal in specific cell types. For example, v-abl is lethal to some BALB/c 3T3 cell lines (54), high concentrations of mos have been shown to coincide with cell mortality in Moloney murine sarcoma virus-infected NIH 3T3 cells (37), and overexpression of Ha-ras causes growth arrest in some cases (22). Expression of src does not transform myocytes (25, 36) and induces differentiation rather than transformation in PC12 pheochromocytoma cells (1). Recently, it has been reported (14) that C127 cells could not be transformed by v-src and other tyrosine kinase oncogenes, v-fms and trk, although they could be transformed by v-ras and by serinethreonine kinase oncogenes, v-mos and v-raf. However, C127-v-src cells, unlike the 3T3-TNR9-v-src cells reported here, acquire a transformed phenotype upon continued culture. Furthermore, the response of C127 cells to phorbol ester treatment is unknown, as is the relationship, if one exists, between this response to TPA and resistance to transformation by v-src. Although growth inhibition by oncogenes has been reported before, this is the first case in which growth inhibition by an oncogene correlated with growth inhibition in response to a defined growth-regulatory agent, namely, the tumor promoter TPA. The most parsimonious explanation for these results is that src expression stimulates functionally significant activity in the PKC signalling pathway.

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ADDENDUM

While this report was under review, Biemann and Erikson (3) reported that the p80 PKC substrate was decreased in

3T3-TNR9 cells, in contrast to our earlier results (5). They also reported that 3T3-TNR9 cells, in contrast to parental 3T3 cells, were defective in the ability to down regulate PKC in response to chronic TPA treatment. This defective response was seen only at high cell densities, and the researchers reported that their 3T3-TNR9 cells grew to a very high density, forming multiple cell layers. In our experience, 3T3-TNR9 cells do not ordinarily grow to very high densities or form multiple layers unless they are beginning to become transformed—which occurs spontaneously upon extended passaging. Because p80 phosphorylation and PKC down regulation may well be affected by malignant transformation, it seems possible to us that the results of Biemann and Erikson are secondary consequences of these phenotypic changes.

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