Methodology article

**Comparative evaluation of gene delivery devices in primary cultures of rat hepatic stellate cells and rat myofibroblasts**

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**Abstract**

**Background:** The hepatic stellate cell is the primary cell type responsible for the excessive formation and deposition of connective tissue elements during the development of hepatic fibrosis in chronically injured liver. Culturing quiescent hepatic stellate cells on plastic causes spontaneous activation leading to a myofibroblastic phenotype similar to that seen in vivo. This provides a simple model system for studying activation and transdifferentiation of these cells. The introduction of exogenous DNA into these cells is discussed controversially mainly due to the lack of systematic analysis. Therefore, we examined comparatively five nonviral, lipid-mediated gene transfer methods and adenoviral based infection, as potential tools for efficient delivery of DNA to rat hepatic stellate cells and their transdifferentiated counterpart, i.e. myofibroblasts. Transfection conditions were determined using enhanced green fluorescent protein as a reporter expressed under the transcriptional control of the human cytomegalovirus immediate early gene 1 promoter/enhancer.

**Results:** With the use of chemically enhanced transfection methods, the highest relative efficiency was obtained with FuGENE™6 gene mediated DNA transfer. Quantitative evaluation of representative transfection experiments by flow cytometry revealed that approximately 6% of the rat hepatic stellate cells were transfected. None of the transfection methods tested was able to mediate gene delivery to rat myofibroblasts. To analyze if rat hepatic stellate cells and myofibroblasts are susceptible to adenoviral infection, we have inserted the transgenic expression cassette into a recombinant adenoviral type 5 genome as replacement for the E1 region. Viral particles of this replication-deficient Ad5-based reporter are able to infect 100% of rat hepatic stellate cells and myofibroblasts, respectively.

**Conclusions:** Our results indicate that FuGENE™6-based methods may be optimized sufficiently to offer a feasible approach for gene transfer into rat hepatic stellate cells. The data further demonstrate that adenoviral mediated transfer is a promising approach for gene delivery to these hepatic cells.

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**Background**

Transfection is the insertion of foreign molecules such as cDNAs or promoter constructs into eukaryotic cells. This method has become a powerful experimental tool for studying gene functions and to analyze the control of gene expression. Genes of interest can either be trans-
fected transiently or stable into cultured mammalian cells. Detailed protocols for efficient gene transfer to various primary cells and continuous cell lines, irrespective whether these cells are grown as monolayers or in suspension have been established during the last decades. Many methods have been developed to overcome the low transfection efficiency in differentiated cells if classical approaches, such as calcium-phosphate-DNA coprecipitates [1], diethylaminoethyl (DEAE)-dextran [2] or polylysine mediated gene transfer [3] are applied. Transfection by electroporation, microinjection, biolistic particle delivery, activated dendrimers or cationic liposomes are useful for established cell lines [4,5,6,7,8]. However, in cells of primary culture most of these transfection systems proved to be inefficient. Direct introduction of genes into rat hepatic stellate cells (rHSCs) and their transdifferentiated phenotype, i.e. the rat myofibroblasts (rMFBs) is difficult to achieve, in part due to the quiescent and fragile phenotype of rHSCs or the extracellular matrix in which the rMFBs are embedded. In normal liver, quiescent HSCs (also called Ito cells, lipocytes, fat-storing cells) are the precursor cells for MFBs, which are responsible for the dramatic increase in the synthesis of extracellular matrix proteins in cirrhotic livers [9,10]. Upon fibrogenic stimuli, HSCs become activated, a process in which they loose vitamin A granules, proliferate, change morphologically into MFBs, and increase their synthesis of extracellular matrix proteins [11, 12]. Culture of quiescent HSCs on a plastic surface also results in spontaneous activation of these cells similar to that seen in liver fibrosis in vivo. Efficient gene delivery to cultured rHSCs and rMFBs would therefore be of great interest for studying the processes involved in hepatic fibrogenesis and for gene-therapeutic devices. To compare various transfection mediators for their potential to increase the efficiency of gene delivery to rHSC and rMFB we used the reporter plasmid pEGFP-C1 expressing the enhanced green fluorescent protein (EGFP) from the jellyfish Aequorea victoria as a reporter expressed under transcriptional control of the ubiquitously active human cytomegalovirus (CMV) immediate early gene 1 promoter. Transfections were performed with the commercially available cationic liposome reagents Effectene, LipofectAmine Plus, Superfect, a classical calcium phosphate based method with and without glycerol shock, and the lipid-based reagent FuGENE™6 applying essentially the protocols given by the manufacturers. As control for evaluation of the suitability of each transfection protocol we transfected NIH/3T3 cells in parallel under the same conditions. Transfection efficiencies for the established cell line NIH/3T3 was in the expected range (Effectene ~9.5%, LipofectAmine Plus ~7.5%, Superfect ~2.5%, calcium-phosphate/-glycerol shock ~8%, calcium-phosphate/+glycerol shock ~12%, FuGENE™6 ~30%) there was obviously an overall low transfection rate of rHSCs and rMFBs.

Interestingly, only FuGENE™6 was able to mediate a significant gene transfer (~ 6%) to rHSCs (Fig. 1). None of the mediators tested was able to provide sufficient gene delivery in rMFBs. FACS analysis was used to estimate the percentage of transfected cells. EGFP has a single red-shifted excitation peak at 488 nm which is suitable for flow cytometric analysis by argon ion laser excitation and a standard FITC filter set [13]. The viability of transfected cells was proven by double staining with propidium iodide (PI), which selectively stains dead cells. Fluorescence signals were acquired using a 488 nm excitation and a 530 ± 30 nm emission fluorescence filter for EGFP and a 630 ± 11 nm emission fluorescence filter for PI, respectively. This "double staining" protocol allows to discriminate between viable but not transfected cells, non viable and not transfected cells, viable transfected cells and dead transfected cells. Representative results obtained after transfection with FuGENE™6 as vehicle are summarized in Fig. 2. In agreement with our microscopic results (see above) 6% of rHSC were found to be positive for EGFP. The transfected cells are viable as evidenced by the lack of PI staining. Furthermore, the
flow cytometric analysis revealed that the non viable rHSC were negative for EGFP. The fraction of about 11% dead rHSC may be due to destructive effects of trypsini-zation and washing procedures on these fragile cells. rMFB were transfectable neither with FuGENE™6 nor with the other non viral transfection systems tested. The fraction of 30% transfected NIH/3T3 cells documents the correct handling of the transfection protocol.

It was previously shown that the lipid formulation Fu-GENE™6 is able to promote DNA uptake by primary thy-rocytes and articular chondrocytes known to be refractory to most transfection systems [14, 15]. In our hands none of the five transfection systems tested was able to (>0.5%) provide significant gene delivery to rMF-Bs. Human MFBs isolated from outgrowths of cirrhotic

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Figure 1
Representative analysis of transfection efficiencies of rat hepatic stellate cells and myofibroblasts using FuGENE™6 transfection procedure. Rat hepatic stellate cells (rHSCs) were isolated from adult male Sprague-Dawley rats by the pronase-collagenase method and a single-step density gradient centrifugation. Rat myofibroblasts (rMFBs) were prepared by subculturing primary rHSCs by trypsinization at day 7 after seeding following spontaneous activation on a plastic surface. The established murine cell line NIH/3T3 served as a reporter transfection cell line. Cells were transfected with 2 µg of reporter construct pEGFP-C1 and FuGENE™6 according to the suppliers instructions. After incubation for proposed time media were changed and transfection efficiencies were monitored 48 hours after transfection. Representative phase contrast microscopy (A, C, E) and fluorescence microscopy (B, D, F) of rHSCs (A, B), rMFBs (C, D) and NIH/3T3 cells (E, F) are shown. The intensity of fluorescence varies among transfected cells indicating various levels of reporter expression.

Figure 2
Representative results of flow cytometric analysis of transfected rat hepatic stellate cells and rat myofibroblasts in comparison to NIH/3T3 cells using FuGENE™6 vehicle. For this experiment rHSCs, rMFBs and NIH/3T3 cells were transfected 2 days after seeding with 2 µg reporter plasmid complexed with 5 µl FuGENE™6. FACS analysis was performed 48 hours after transfection. Cultured cells were trypsinized under standard conditions and flow cytometric measurements were performed immediately after collection of cells. Fluorescence signals were recorded with a flow cytometer FACS-Calibur (Becton Dickinson, Sparks, MD) using a 488 nm excitation and a 530 ± 30 nm emission fluorescence filter for enhanced green fluorescence protein (EGFP) and a 630 ± 11 nm emission fluorescence filter for propidium iodide (PI), respectively. Data were acquired and analyzed with the CellQuest™ software version 3.1 (Becton Dickinson). Histograms of fluorescence intensities in EGFP (A, C, E) and PI (B, D, F) channels are shown for rHSC (A, B), rMFB (C, D) and reporter cell line NIH/3T3 (E, F), respectively. To establish background for fluorescence and to set gates for data acquisition, mock-transfected cells (not shown) were used. Mean fluorescence intensity was used to calculate levels of EGFP expression. Cells that took up PI were deemed nonviable. Nontransfected cells did not show fluorescence in EGFP channel.
liver pieces and cells from the established line CFSC [16] derived from cirrhotic rHSCs were highly transfectable with all mediators (not shown). A potential reason for the observed higher transfection susceptibility of human MFBs and immortalized rMFBs might be the difference in the rate of cell proliferation. A connection between effective gene transfer and mitotic activity was recently demonstrated in various cells and is also well documented in smooth muscle cells [17, 18].

To test if adenoviral gene delivery is a suitable alternative to overcome the observed insufficient gene delivery rates we analyzed the susceptibility of rHSCs and rMFBs to adenoviruses. To facilitate the insertion of the CMV/EGFP reporter cassette within an appropriate adenoviral shuttle vector we cut pEGFP-C1 with AsnI and SspI, filled in the resulting 1.66-kbp fragment with Klenow DNA polymerase and inserted this fragment into the filled in EcoRI site of vector pΔE1sp1A (Fig. 3A) [19]. The orientation of the reporter transgene and the cloning boundaries in the resulting vector pΔE1sp1A-CMV-EGFP were confirmed by digestion with multiple restriction endonucleases and sequencing. We then cotransfected vector pΔE1sp1A-CMV-EGFP and backbone vector pJM17 [20] into the packaging cell line 293 [21], which supplies the missing adenoviral E1 gene products necessary to generate recombinant viral particles (Fig. 3B and 3C). Based on reports describing that the integrated colinear Ad5-segment spanning bp 1 to bp 4344 of Ad5 genome [22] is transcriptionally silenced or lost frequently in 293 cultures at high passage numbers we first tested whether corresponding Ad5-genes (E1A and E1B) were actively transcribed in the 293 cells used in this study. Therefore, we performed a Northern blot analysis (Fig. 4) with total RNA of 293 cells and the 2.8-kbp EcoRI/HindIII fragment of vector pAd5Sal/I containing bp 1 to bp 2804 of Ad5 genome (cf. GenBank Accession number X02996.1) [23]. The sizes of observed hybridization signals (1.3 kb and 2.6 kb) expressed at high levels are in agreement with the complexity of the primary transcripts for E1A (1134 nt) and E1B (2369 nt) genes if in addition 200 nt for poly A+ tail are estimated. In the murine control cell line NIH/3T3 transcripts of E1A or E1B are virtually absent due to the absence of Ad5 sequences in these cells. The Northern analysis therefore revealed that the 293 cells used in this study are able to produce relevant mRNAs for both trans-activator genes. Twenty four hours after cotransfection of pΔE1sp1A-CMV-EGFP and pJM17 approximately 40% of 293 cells were positive for EGFP (not shown). Six days later the typical viral foci which are able to produce recombinant viral particles were observed and 7-12 days after transfection all cells were infected by developing recombinant Ad5-CMV-EGFP particles. Viral particles were isolated, amplified once on 293 cells and purified by standard procedures.

To test the susceptibility of cultured rHSC/rMFB we infected them two days after seeding using a multiplicity of infection (MOI) of 10. Analysis of the cells for forty-eight hours after infection revealed that 100% of the cells (rHSCs and rMFBs) were positive for EGFP providing evidence that adenoviral based vectors are very effective in gene delivery to these cells in culture (Fig. 5). However, it was surprising that the use of adenovirus at a MOI of 10 was sufficient for complete infection. It is well established that the internalization of adenoviral particles is promoted by integrins [24]. Also, it was previously demonstrated that HSC and MFB express high levels of various types of integrins [25, 26]. Therefore, it is most likely that the observed high adenoviral susceptibility of HSC and MFB may be caused by high-level expression of integrins. Interestingly, also the relative intensities of fluorescence signals in these cells were extremely high compared to those observed after chemically enhanced gene delivery (compare Fig. 5B and Fig. 1B). Infected cells continued to express the product of the transgene (EGFP) for at least 2 weeks. It is most likely that the termination of transgene expression after 2 weeks is related to the loss of episomal transgene, which is not incorporated into the host genome and, therefore, is lost with cell division. Partial cellular disintegration occurred (not shown) at later stages most likely due to the intrinsic cytotoxicity of high EGFP content or the overexpression of some viral proteins [27]. Independent experiments revealed further that infection of these sinusoidal hepatic cells is highly reproducible, irrespective whether infection is performed in the presence of 2%, 5% or 10% fetal calf serum (not shown).

Based on these data we conclude that a MOI of 10 is sufficient to achieve an infection of 100% in cultured rHSC and rMFB, respectively, under most experimental conditions.

**Conclusions**

In agreement with other reports on gene delivery to cultured rHSCs by adenoviral based techniques our data show that this method is straightforward particularly when high efficiency of gene transfer is required [28,29,30,31]. Furthermore, our report indicates that introduction of foreign DNA even into rMFBs is possible by use of adenoviral based vector systems. Because of the considerable interest on the rHSC/rMFB transition as a cell culture model for liver fibrogenesis the improvement of efficiency of gene delivery to these cells should facilitate applications such as reproducible reporter vector assays, or bulk expression of signalling proteins for biochemical or cell biological assays.
Figure 3
Construction of a replication-defective recombinant adenovirus expressing enhanced green fluorescent protein under transcriptional control of the human cytomegalovirus immediate-early gene 1 promoter/enhancer. (A) In the mammalian reporter vector pEGFP-C1 the enhanced green fluorescent protein (EGFP) is expressed under control of the human cytomegalovirus immediate-early gene 1 promoter (P_{CMV}). Abbreviations are: Kan/Neo, kanamycin/neomycin resistance genes; Amp, ampicillin resistance gene; pUC ori, E. coli origin of replication; SV40 ori, simian virus 40 origin of replication; SV40 polyA, simian virus 40 polyadenylation signal; HSVTK polyA, herpes simplex thymidine kinase polyadenylation signal. The adenoviral shuttle vector p∆E1sp1A contains Ad5 sequences from bp 22 (0 μm) to bp 5790 (16.1 μm) with a deletion of E1 sequences (∆E1) from bp 342 to bp 3523 (1.0 - 9.8 μm) and a selectable ampicillin resistance gene (Amp). A multiple cloning site (MCS) containing unique restriction sites for ClaI, BamHI, XhoI, XbaI, EcoRV, EcoRI, HindIII, Sall, and BglII is embedded in the Ad5-sequences. For construction shuttle vector p∆E1sp1A-CMV-EGFP the 1656-bp Asnl/SspI fragment from plasmid pEGFP-C1 was filled in by Klenow DNA polymerase and cloned into the EcoRI digested and filled in p∆E1sp1A vector. (B) For integration of the reporter cassette from pE1∆sp1A-CMV-EGFP into the Ad5 backbone plasmid vector pJM17 both plasmids were cotransfected into human embryo kidney cell line 293 leading to homologous recombination between common Ad5 regions. Generation of recombinant viral particles were visualized by an increase of EGFP-positive cells and by viral focus formation in fluorescence microscopy. (C) Released replication-defective viral particles are infectious and are capable to deliver the CMV-EGFP cassette to target cells. The nucleotide sequence of the cloned vector p∆E1sp1A-CMV-EGFP is deposited in GenBank (Accession number AF288620).
Additional studies will be required to determine the optimal in vivo conditions for adenoviral gene transfer to rHSCs/rMFBs.

**Material and Methods**

**Isolation and culture of rat hepatic stellate cells**

All animal protocols were in full compliance with the guidelines for animal care and were approved by the Animal Care Committee from the government. Adult male Sprague-Dawley rats (Harlan Winkelmann GmbH, Borchen, Germany), which had free access to altromin chow (Altromin GmbH, Lage, Germany) and water were anaesthetized with an intramuscular injection of 100 mg/kg body weight ketaminhydrochloride (Ketavet®; SANOFI-CEVA GmbH, Düsseldorf, Germany) and 7 mg/kg body weight xylazinhydrochloride (Rompun®; Bayer-Vital, Leverkusen, Germany). rHSCs were isolated by the pronase-collagenase method, a two-step method for preparation of rat liver cells comprising a first step in which the liver is pre-perfused with a Ca\(^{2+}\)-free medium in order to remove intercellular Ca\(^{2+}\), and a second step in which the liver is perfused with a Ca\(^{2+}\) requiring collagenase mixture [32, 33, 34]. Briefly, livers were perfused at 10 ml/minute with 200 ml Hanks balanced salt solution (HBSS) without Ca\(^{2+}\) and Mg\(^{2+}\) (PAA Laboratories GmbH, Linz, Austria), 100 ml solution E [0.35% (w/v) pronase E in HBSS with Ca\(^{2+}\) and Mg\(^{2+}\)] followed by recirculating perfusion for approximately 30 minutes with 60 ml solution F [0.015% (w/v) collagenase in HBSS with Ca\(^{2+}\) and Mg\(^{2+}\)]. Thereafter, livers were dissected under permanent pH control (pH 7.3) in 100 ml HBSS with Ca\(^{2+}\) and Mg\(^{2+}\) containing 10 mg DNase, type II (Roche, Mannheim, Germany). The cell suspension was subsequently filtered through nylon mesh (250 µm and 100 µm) and centrifuged for 7 minutes at 450 × g at 4°C. The resulting pellet was washed in ice cold solution G containing 0.25% (w/v) bovine serum albumin in HBSS/+Ca\(^{2+}\)/+Mg\(^{2+}\), rHSCs were purified by single-step density gradient centrifugation with 8.25% (w/v) Nycodenz® (Nycomed Pharma AS, Oslo, Norway) as described in detail elsewhere [33]. rHSCs were carefully collected by aspiration of the white top layer of the gradient. The purity of cell preparations was assessed by light microscopic appearance, vitamin A autofluorescence of the cells and positive immunofluorescence stainings [35]. rMFBs were prepared by subculturing primary rHSCs by trypsinization at day 7 after seeding following spontaneous activation on the plastic surface. rHSCs and rMFBs were cultured in Dulbecco's modified Eagle medium (DMEM, BioWhittaker Europe, Verviers, Belgium) containing 4 mM L-glutamine and 10% fetal calf serum (Se-Romed, Biochrom KG, Berlin, Germany). Additionally,
all culture media were supplemented with penicillin (100 IU/ml) and streptomycin (100 µg/ml), respectively. All cultures were maintained at 37°C and 5% CO₂ in a humidified atmosphere.

The murine fibroblast cell line NIH/3T3 [36] was purchased from the American Type Culture Collection (Rockville, MD) and maintained in culture at 37°C in DMEM supplemented with 10% fetal calf serum, 4 mM L-glutamine, penicillin (100 IU/ml) and streptomycin (100 µg/ml). All transfections of NIH/3T3 cells with various transfection agents were performed at approximately 60% confluency.

Transfection of hepatic stellate cells
The reporter construct pEGFP-C1 (GenBank Accession number U55763) was purchased from CLONTECH (Palo Alto, CA). rHSCs were transfected two days after seeding, rMFBs cells two days after subculturing (see above) with 2 µg plasmid DNA. The DNA was isolated by a standard alkaline extraction method or a cesium chloride gradient purification protocol. Transfections were performed with the commercially available cationic liposome reagents Effectene (Qiagen, Hilden, Germany), Superfect (Qiagen), the lipid-based reagent FUGENE™6 (Roche), and a classical calcium phosphate based method (Promega, Madison, WI) with and without glycerol shock, following essentially the instructions of the manufacturers. For glycerol shock the medium was removed, cells were rinsed in HBSS/-Ca²⁺/-Mg²⁺ for 2 minutes at 37°C. Thereafter, glycerol solution was removed and cells were washed three times with HBSS/-Ca²⁺/-Mg²⁺ prior addition of regular growth medium. Transfection efficiencies were monitored 48 hours later by fluorescence microscopy and flow cytometry analysis.

Microscopic data documentation
Microscopic images shown are from representative experiments. For visual observation of transfected and infected cells expressing EGFP, we typically used a 40 × objective lens and a 10 × eyepiece lens. Documentation of representative eye fields had been performed with a HV-C20 digital camera (HITACHI Denshi Ltd., Tokyo, Japan) and the DISKUS software package version 4.14 (Hilgers, Königswinter, Germany).

Flow Cytometry
Flow cytometric measurements were performed immediately after collection of cultured cells. Cells were trypsinized under standard conditions and washed once with HBSS/-Ca²⁺/-Mg²⁺. To get informations about viability, collected cells were additionally stained with propidium iodide (PI) (Sigma, Deisenhofen, Germany). Fluorescence signals were recorded with a flow cytometer FACS-Calibur (Becton Dickinson, Sparks, MD) using a 488 nm excitation and a 530 ± 30 nm emission fluorescence filter for EGFP and a 630 ± 11 nm emission fluorescence filter for PI, respectively. Data were acquired and analyzed with the CellQuest™ software version 3.1 (Becton Dickinson).

Construction of replication-defective recombinant adenoviruses
Plasmids pΔE1sp1A [19] and pJM17 [20] for construction of adenoviral vector were obtained from Microbix Biosystems Inc. (Toronto, Ontario, Canada). For construction the adenoviral reporter shuttle vector pΔE1sp1A-CMV-EGFP the 1656-bp AsnI/SspI fragment from plasmid pEGFP-C1 was filled in by Klenow DNA polymerase and cloned into the EcoRI digested and Klenow filled in pΔE1sp1A. The orientation of the CMV-EGFP expression cassette in pΔE1sp1A-CMV-EGFP and the integrity of cloning boundaries were verified by restriction analysis and sequencing with flanking primers 5’-d(GCGTAACCAGTAAAGATTTT)-3’ and 5’-d(GCGCACCATCAATGCTGGAG)-3’, respectively. Primers were obtained from MWG- Biotech AG (Ebersberg, Germany) and sequence reactions were done with the ABI PRISM BigDye® termination reaction kit (PE Applied Biosystems, Weiterstadt, Germany). Integration of vector sequences from pΔE1sp1A-CMV-EGFP into adenoviral backbone vector pJM17 for production of replication-deficient virus particles were performed by in vitro homologous recombination in the human embryo kidney cell line 293, which constitutively express the E1 gene products required for propagation of recombinant adenoviruses [21]. Briefly, approximately 5 × 10⁵ 293 cells were seeded in six-well dishes and cotransfected with 2 µg of plasmid pΔE1sp1A-CMV-EGFP, 2 µg pJM17 and 10 µl of the FuGENE™6 reagent. After 16 hours at 37°C, the DNA containing the transfection mix was removed, and 2 ml of growth medium was added and cells were cultured for 10-12 days with subsequent addition of fresh medium. Generation of recombinant viral particles were visualized by increase of EGFP-positive cells and by viral foci formation in fluorescence microscopy. After total infection viral particles were released from cells by three rounds of a freeze-thaw cycle. Viral particles were separated from cell debris by centrifugation at 3000 rpm for 10 minutes. To generate higher titer viral stocks, 293 cells were re-infected at a multiplicity of infection (MOI) of 1 and grown for 3-4 days, at which time viruses were harvested as described above.

Adenoviral infection
Viral stock solutions were titered on 293 cells in growth medium containing 10% fetal calf serum following stand-
ard procedures [37] and kept frozen at -20°C until use. For infection of rHSCs/rMFBs with adenoviral particles 10^5 cells were seeded in six-well dishes with 2 ml medium and cells were infected 2 days later with 500 µl viral stock containing approximately 10^6 plaque forming units of Ad5-CMV-EGFP.

RNA isolation and Northern analysis

Cells from human cell line 293 and murine cell line NIH/3T3 were rinsed with ice cold HBSS/−Ca^2+/-Mg^2+, and then lysed in 1 ml per 10^6 cells of lysis buffer containing 25 mM sodium acetate pH 6.0, 4 M guanidine thiocyanate and 0.835% (v/v) β-mercaptoethanol. The lysate was adjusted to 8 ml with lysis buffer, pressed three times through a 20-g needle, layered onto a 4-ml cushion of a solution containing 25 mM sodium acetate pH 6.0 and 5.7 M cesium chloride, and then centrifuged for approximately 24 h at 21°C and 25000 rpm in a Beckman SW41-type rotor. RNA pellets were resuspended in 300 mM sodium acetate (pH 6.0), ethanol precipitated and resuspended in water. The concentration of RNA was determined by absorbance at 260 nm.

For Northern analysis, 30 µg of total cellular RNAs were separated by electrophoresis in a 1.2% (w/v) denaturing agarose gel, transferred to a Hybond-N membrane (Amersham Pharmacia Biotech, Freiburg, Germany) and fixed by baking for 2 h at 80°C. Blots were pre-hybridized for at least 3 h, and then hybridized for 16 h at 37°C to [α-32P]dCTP-labelled probes (Multiprime DNA labelling system; Amersham) in a buffer containing 50% (v/v) formamide, 6 × SSC, 5 × Denhardt’s solution [0.1% (w/v) Ficoll 400, 0.1% (w/v) BSA, 0.1% polyvinylpyrrolidone], 5 mM EDTA, 0.5% (w/v) SDS, and 100 µg/ml sheared denatured herring sperm DNA (Roche). Filters were washed once at 55°C for 20 minutes in a solution containing 2 × SSC, 1 mM EDTA, and 0.1% (w/v) SDS, then twice at 50°C for 20 minutes in a solution containing 0.4 × SSC, 1 mM EDTA, and 0.1% (w/v) SDS. Autoradiographs were exposed for indicated times to Kodak X-OMAT AR films at -80°C using intensifying screens. As an internal standard (loading control) the blots were rehybridized with a GAPDH specific cDNA.

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