

Rho Family GTPases Regulate Mammary Epithelium Cell Growth and Metastasis Through Distinguishable Pathways

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Abstract

Background: Relatively few genes have been shown to directly affect the metastatic phenotype of breast cancer epithelial cells *in vivo*. The Rho family of proteins, including the Rho, Rac and Cdc42 subfamilies, are related to the small GTP binding protein Ras and regulate diverse biological processes including gene transcription, cytoskeletal organization, cell proliferation and transformation. The effects of Cdc42, Rac and Rho on the actin cytoskeleton suggested a possible role for Rho proteins in cellular motility and metastasis, however a formal analysis of the role of Rho proteins in breast cancer cellular growth and metastasis *in vivo* had not previously been performed.

Materials and Methods: We generated a panel of MTLn3 rat mammary adenocarcinoma cells that expressed similar levels of dominant inhibitory mutants of Cdc42-, Rac- and Rho-dependent signaling, to examine the contribution of these GTPases to cell spreading, guided chemotaxis, and metastasis *in vivo*. The ability of Rho proteins to regulate intravasation into the peripheral blood was determined by implanting MTLn3 cell stable dominant negative lines in nude mice and measuring the formation of breast cancer

cell colonies grown from the peripheral blood. Serial sectioning of the lungs was performed to determine the presence of metastasis in mice in which mammary tumors expressing the dominant negative Rho family proteins had grown to a similar size.

Results: Cell spreading of MTLn3 cells was selectively abrogated by N17Rac1. N19RhoA and N17Cdc42 reduced the number of focal contacts (FCs) and disrupted the colocalization of vinculin with phosphotyrosine at FCs. While N17Rac1 and N17Cdc42 preferentially inhibited colony formation in soft agar, all three GTPases affected cell growth *in vivo*. To distinguish effects on tumorigenicity from intravasation into the bloodstream, implanted tumors were grown to the same size in nude mice. Each dominant inhibitory Rho protein reduced intravasation into the peripheral blood. Lung metastasis of MTLn3 cells was also abrogated by the dominant inhibitory Rho proteins, despite the presence of residual CFU.

Conclusions: These studies demonstrate for the first time a critical role for the Rho GTPases involving independent signaling pathways to limit mammary tumor cellular growth and metastasis *in vivo*.

Introduction

Rho family proteins are closely related to the small GTP binding protein Ras (1). Rho GTPases cycle between an active GTP-bound state and an inactive GDP-bound state to transduce diverse signals from cell surface receptors to intracellular targets (1). Rho proteins regulate a diverse spectrum of biological

processes including regulation of gene transcription, superoxide production, changes in cytoskeletal organization, cell proliferation and transformation (1–4). Analyses performed predominantly in fibroblasts, employing microinjection of dominant-active or dominant-negative mutants, demonstrated that specific Rho proteins regulate distinct functions (1). The functional specificity of Rho GTPases is transduced, at least in part, by an interaction with specific effector proteins that coordinate the activation of multiple signaling cascades (1). The regulation of the actin cytoskeleton and specialised cellular adhesion structures is also quite selective for distinct members of the Rho

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family. Thus, activated RhoA stimulates stress fiber formation (5), Rac1 controls growth factor-induced lamellipod formation in Swiss 3T3 fibroblasts (2) and Cdc42 regulates filopodial protrusion (6,7)

The effects of Cdc42, Rac and Rho on the actin cytoskeleton and cellular adhesion suggested a possible role for Rho proteins in cellular motility. A role for Rho proteins in migration of fibroblasts and neutrophils was demonstrated using the C3 exoenzyme and Rho-GDI (GDP dissociation inhibitor) to block Rho function (8,9). In Bac1.2F5 macrophages, inhibition of Rac and Rho function blocked cell migration in response to CSF-1, while a dominant inhibitory Cdc42 enhanced cell migration (10). Rac activity is also required for PDGF-BB-induced migration of Rat1 cells across a porous membrane (11). In T47D mammary epithelial cells, activated Rac1 and Cdc42, but not RhoA, enhanced cell migration across filters coated with collagen (12). Cellular migration through increasingly complex surfaces such as a three dimensional (3D) collagen gel requires distinct properties of Rho proteins (13). Invasion of 3D collagen matrices by Rat1 cells was inhibited by dominant negative mutants of Cdc42 and Rac1 indicating fibroblast invasion requires optimal level of activity of multiple Rho family members (13). The process of cellular metastasis *in vivo* is a still more complex process involving several distinct phases including invasion through tissue structures, intravasation and survival in the peripheral blood and the ability to adhere at a distant site. It is predicted that Rho GTPases might regulate mammary epithelial cell invasion and metastasis *in vivo*. The contribution of the Rho proteins to *in vivo* mammary tumor bioassays such as tumor blood burden and lung metastasis is therefore of fundamental importance (14,15). As the role of Rac, Rho and Cdc42 to the components governing metastasis *in vivo* had not previously been formally assessed, we employed a syngeneic model to address this question.

In this study we employed MTLn3 cells that were originally derived from a lung metastasis of the 13762 rat mammary adenocarcinoma (16,17). The effects of Epithelial Growth Factor (EGF) on guided chemotaxis have been extensively studied in the MTLn3 cells. EGF, through binding to its receptor, functions as a chemoattractant for a number of different cell types and is thought to enhance the migration and invasion of tumor cells (18–21). EGF-mediated chemotaxis of MTLn3 cells was shown to include actin polymerization at the leading edge of the lamellipod to enable cellular extension and motility (22–24). In this study, we used MTLn3 cells stably expressing Rho inhibitory proteins to examine the role of the Rho family GTPases in coordinating cytoskeletal reorganization and in the regulation of metastasis. We demonstrate for the first time separable signaling pathways regulated by Rho GTPases that together contribute to *in vivo* mammary epithelial tumor metastasis.

Materials and Methods

Cell Culture, Proliferation and Transformation Assays

Cell culture was performed as previously described (24). Cells were grown in α -MEM (Gibco), supplemented with 5% fetal calf serum and antibiotics. For all experiments, unless otherwise mentioned, MTLn3 cells were plated in regular medium for 24 h at low density on tissue culture dishes (Falcon), or MaTek dishes (MaTek Corporation, Ashland, MA), which had been previously coated for 2 h at room temperature with rat tail collagen type 1 (Collaborative Biomedical, Bedford, MA) at 30 μ g/ml in DPBS (Gibco). MTLn3 cells were starved for 3 h prior to the experiment in α -MEM medium supplemented with 0.35% bovine serum albumin (BSA) and 12 mM HEPES (starvation medium). Stimulation was done with a final concentration of 5 nM murine EGF (Life Technologies) in starvation medium.

The plasmids pEXV, Myc-N17Rac1, Myc-N19RhoA (25), pCMV5, and Flag-N17Cdc42 (26) have been previously described. MTLn3 cells were stably transfected with the epitope-tagged Rho family GTPase dominant negative mutants and pCNeo, selected using G418 (800 ng/ml) and maintained in G418 (400 ng/ml). Expression of the mutant Rho proteins was confirmed by western blot of the epitope tag. For cellular proliferation assays, the MTLn3 cells encoding Rho family inhibitory proteins, or the parental line containing the empty vector control, (pEXV, pCMV), were plated in 6-well tissue culture plates. After six days growth in regular culture medium (α -MEM, 5% FBS) cells were trypsinized, harvested and counted using a hemocytometer. The soft agar growth assays was performed as previously described (27–29). MTLn3 stable cells (3×10^4 cells) were suspended in 3 ml of α -MEM containing 5% FBS and 0.33% SeaPlaque low-melting-temperature agarose (FMC Bioproducts). The suspension was plated on 60 mm dishes containing a 2 ml layer of solidified α -MEM, 5% FBS and 0.5% SeaPlaque agarose, in quadruplicate. The cells were allowed to settle at the interface between these layers for 30 min at 37°C. Cells were fed every 3 days by overlaying with 2 ml of complete medium containing 0.33% SeaPlaque agarose. After 15 days, the plates were examined and the colonies were counted under a Nikon Phase contrast microscope at 4 \times or 6 \times magnification. Experimental values represent the average number of foci in the 60 mm plates for each experimental condition; error bars represent the observed SEM between the 4 plates.

Immunofluorescence and Rhodamine-Phalloidin Staining of Filamentous Actin (F-actin)

Cells were seeded onto 22 mm square glass cover slips (Becton Dickinson, Bedford, MA) coated with rat tail collagen type 1 (27 μ g/ml) in 6-well tissue culture plates. When 70–80% confluent, cells were

rinsed with $1 \times$ PBS at 37°C , fixed in 3.7% formaldehyde in buffer F (5 mM KCl, 137 mM NaCl, 4 mM NaHCO_3 , 0.4 mM KH_2PO_4 , 1.1 mM Na_2HPO_4 , 2 mM MgCl_2 , 5 mM PIPES, pH 7.2, 2 mM EGTA, 5.5 mM glucose) for 5 min at 37°C , extracted in 0.5% Triton X-100 in buffer F for 20 min at room temperature and washed in 0.1 M glycine in buffer F for 10 min at room temperature. The cover slips were washed 5 times for 5 min in $1 \times$ TBS, wicked dry and placed on Parafilm in a humidified chamber and 150 ml of 1% BSA, 10% goat serum in $1 \times$ TBS with rhodamine-phalloidin (0.5 mM) (Molecular Probes, Eugene, OR) added for 20–30 min. Excess rhodamine-phalloidin was washed from the cover slips, 5 times for 5 min with 1% BSA in $1 \times$ TBS and, if no immunofluorescent staining was carried out, the cover slips were mounted in Prolong reagent (Molecular Probes, Eugene, OR). For immunofluorescent staining, the cells were incubated at room temperature for 60 min with primary antibody after blocking. The cover slips were then washed 3 times for 10 min in 1% BSA in $1 \times$ TBS and incubated with secondary antibody for 45–60 min at room temperature before a final series of three 10 min washes in 1% BSA in $1 \times$ TBS. If two primary antibodies were used, the antibodies were added sequentially and each directly followed by incubation with their respective secondary antibodies. The cover slips were mounted as before and examined under an Olympus 1×70 inverted microscope with images recorded using a Photometrics CH1 cooled CCD camera.

Antibodies used included an anti-phosphotyrosine (P112300 polyclonal antibody, Transduction Labs), anti-vinculin (V4505, Sigma), anti-Flag antibody (Sigma, (M2)), and anti-Myc (9E10, Santa Cruz, mouse monoclonal).

Microchemotaxis Assay

Chemotactic responses were assessed using the 48-well chemotaxis chamber (Neuroprobe, Cabin, MD, USA) following the manufacturers instructions. A Nucleopore filter with $8 \mu\text{m}$ pores (Osmonics/Poretics Products, Livermore, CA, USA) was coated with rat tail collagen type 1 at a final concentration of $35.6 \mu\text{g/ml}$ in DPBS without calcium or magnesium (JRH Biosciences) for 2 hours. MTLn3 cells were cultured in MEM medium with 5% FBS and 0.5% Penicillin/Streptomycin on 10 cm plates. Medium was aspirated and replaced with growth medium containing 12 mM HEPES pH 7.4 and BSA 0.35% in MEM (termed MEMH) without serum or antibiotics for 2 hours. Cells were then harvested with 26.6 mM EDTA in DPBS, resuspended in MEMH and counted. The lower wells of the chamber were filled with $30 \mu\text{l}$ MEMH containing appropriate concentration of EGF (Life Technologies) or buffer, then the chamber was assembled incorporating the collagen-coated filter. The stock of EGF was prepared in filtered DPBS. The upper wells were then filled with MEMH containing 20,000 cells/well in a total

volume of $50 \mu\text{l}$. The chamber was incubated at 37°C for 3 h, disassembled and the upper side of the filter scraped to remove cells that had not traveled through the filter. The filters were fixed in 3.7% formaldehyde in DPBS for 30 min, washed twice in water and stained for 12–18 h in hematoxylin. The filters were then rinsed in water and mounted for viewing. Results are means \pm SEM from 5- to 8 separate experiments.

Lamellipod Extension and Spreading Assays

Cells were seeded onto 22 mm square collagen-coated glass cover slips in 6-well tissue culture plates and treated or not with 5 nM EGF for 3 min. Cells were fixed in 3.7% formaldehyde, extracted in 0.5% Triton X-100 and mounted. The extension of lamellipodia was assessed as previously described by us (22,28). All samples were examined under an Olympus 1×70 inverted microscope with images recorded using a Photometrics CH1 cooled CCD camera. A total of 300 cells were examined.

For the spreading assays, MTLn3 cells were plated (10^4 cells/well) on a collagen coated ($27 \mu\text{g/ml}$) 6-well plate. At several time points after plating, the cells were viewed using phase contrast and a $10\times$ objective. The number of spread cells was counted by visual inspection using a 5×5 grid. Cells that turned phase-dense during the spreading process and displayed at least one protrusion were counted as spread. A total of 300 cells were counted in five different fields that were selected at random. To observe the F-actin in MTLn3 cells, the cells were fixed and stained with FITC-phalloidin after plating for various times. The data were expressed as mean \pm SEM for percentages of positive cells in the five fields.

Western Blots

Immunoprecipitation western blotting was performed as previously described using either the Flag or Myc antibodies (30,31). The abundance of Fak, Talin, and Vinculin proteins was determined by western analysis as previously described (32,33), using anti-Vinculin antibody (Sigma), anti-focal adhesion kinase (FAK) antibody (F15020 polyclonal antibody, Transduction Labs), guanine nucleotide dissociation inhibitor (GDI) antibody (Acknowledgments, 25), and anti-Talin monoclonal antibody (8d4, Sigma). Cell homogenates ($50 \mu\text{g}$) were separated in an SDS-12% polyacrylamide gel and transferred electrophoretically to a nitrocellulose membrane (Micron Separations Inc., Westborough, MA). After transfer, the gel was stained with Coomassie blue as a control for blotting efficiency. The blotting membrane was incubated for 2 h at 25°C in T-PBS (PBS + 0.05% Tween 20) buffer supplemented with 5% (wt/vol) dry milk to block non-specific binding sites. Following 6 h incubation with primary antibody at a 1:5000 dilution (GDI) in T-PBS buffer containing 0.05% (vol/vol) Tween 20, the membrane was washed with the same buffer. The membrane

was incubated with goat anti-mouse horseradish peroxidase second antibody (Santa Cruz Biotechnology, Santa Cruz, CA) and washed again. The protein was visualized by the enhanced chemiluminescence system (Kirkegaard and Perry Laboratories, Gaithersburg, MD).

In Vivo Tumor Growth, Blood Burden, and Lung Metastasis Assay

To determine the effect of dominant negative N17Rac, N19Rho and N17Cdc42 on MTLn3 cell metastasis, parental and stably transfected MTLn3 cells were injected into the subcutaneous tissue of 6- to 8-week old nude mice (strain Balb/cAnNCr-nuBR, National Cancer Institute) as previously described (25,34). Experiments were conducted under an approved protocol of the AECOM animal ethics committee. Cells were trypsinized, counted, washed twice and resuspended in ice-cold sterile PBS. Nude mice were anesthetized lightly with Methofane (Schering-Plough Animal Health) and injected with 10^6 viable cells in 0.1 ml of PBS. Tumors were palpated every 3- to 4 days after the first week and measured with calipers (mean tumor diameter) to quantitate tumor size. Animals were euthanized 5- to 8.5 weeks after cells were injected. Tumors growing at the site of injection were measured and then removed and rinsed in ice-cold PBS. Samples of tumor tissue were rapidly frozen in liquid nitrogen for further analysis.

Tumor cell blood burden was determined as previously described (35) by placing nude mice with a 5 week old tumor under methofane anesthesia and removing 1 ml of blood via heart puncture. The blood was then spun at 5,000 rpm and the serum layer and buffy coat region were plated into α -MEM growth medium. The following day, plates were rinsed $2\times$ with Dulbecco's PBS (Gibco) to remove red blood cells and regular growth medium (as above) was added. After six days, all clones in the dish were counted. For experiments in which the Rho family dominant inhibitory MTLn3 cells were grown to the same size as vector controls, the subcutaneous tumors were grown for 8.5 weeks.

For measurement of metastasis, excised lungs were placed in 3.7% formaldehyde, mounted in paraffin, sectioned and stained with H&E. Serial slices of 5 μ M thickness were methodically viewed throughout the entire lung with sections viewed using a $20\times$ objective and all visible metastases in a section containing more than 5 cells were counted in each section.

Results

Effects of Rho Proteins on EGF-induced Stress Fiber Formation

To investigate the role of Rho family GTPases in MTLn3 metastatic behavior *in vivo*, we established stably transfected lines. The expression of the various Rho GTPases was confirmed by western blotting of the epitope tagged construct (Fig. 1). Cells extracts

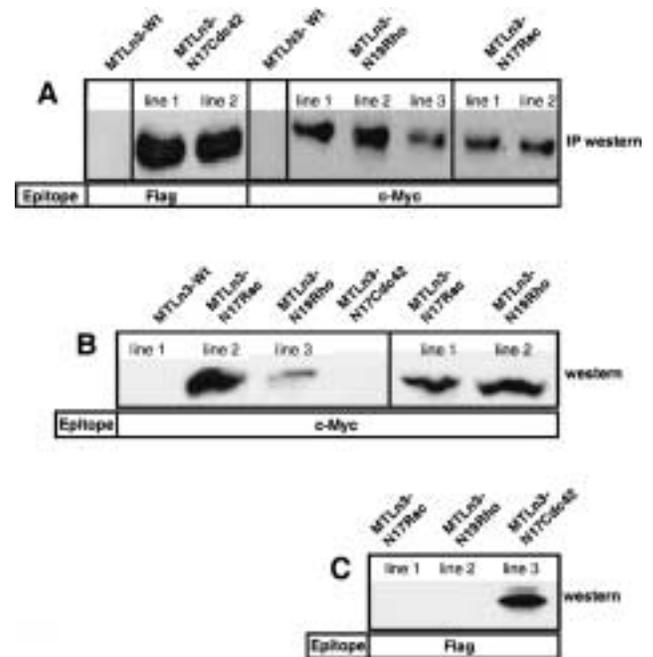


Fig. 1. Expression of dominant inhibitory Rho-family mutants. (A) MTLn3 cells were stably transfected with N17Rac1, N19RhoA, N17Cdc42, pEXV (control vector for Rac and Rho) and pCMV5 (control vector for Cdc42). Cell extracts from each cell line were immunoprecipitated with Flag (pCMV5 and N17Cdc42) or Myc (pEXV, N17Rac1 and N19RhoA) antibodies, separated by SDS-PAGE and identified by western analysis with anti-Flag antibody or anti-Myc antibody. (B,C) MTLn3 cell extracts were separated by SDS-PAGE and western blotting was performed with an antibody to the Flag or Myc epitope.

were either subjected to immunoprecipitation with an antibody to the epitope (Myc or Flag) with subsequent western blotting (Fig. 1A), or subjected to direct western blotting (Fig. 1B,C).

Tyrosine Phosphorylation of Focal Contacts Involves Rho and Cdc42

Many cultured cells form specialized structures known as focal adhesions to link them to the underlying extracellular matrix (ECM). To assess the focal adhesions, MTLn3 cells were plated on the ECM protein, collagen. Focal contacts (FCs) contain a number of cytoskeletal and submembranal anchor proteins including vinculin, paxillin and talin (36–38), which link the transmembrane integrin receptors to the actin cytoskeleton. FCs are also enriched in signal transduction molecules including focal adhesion kinase (FAK) (39). A characteristic of FCs in fibroblasts is their high level of tyrosine phosphorylation (36). The assembly and tyrosine phosphorylation (pTyr) of proteins in FCs is regulated by cytoplasmic factors including Rho, caldesmon as well as by microtubular integrity (37,40–45). We assessed the distribution of pTyr in the FCs of MTLn3 cells (Fig. 2). FCs were identified by their high abundance of vinculin by double immunofluorescence staining

with antibodies to pTyr and to vinculin. Cells grown to 70–80% confluence on collagen-coated glass cover slips were treated with 5 nM EGF or vehicle for either 2 or 4 h with similar results. In the parental vector control MTLn3 cells, pTyr co-localized with vinculin in the basal state with a modest reduction upon EGF treatment (Fig. 2A), consistent with findings in NR6 fibroblasts (46). A modest reduction in vinculin

staining FCs was observed upon EGF treatment at 2 hours. The N17Rac1 lines displayed abundant pTyr and vinculin, which co-localized upon image merging (Fig. 2C). In contrast, vinculin staining FCs were reduced and co-localization with pTyr disrupted in the N19RhoA (Fig. 2D) and N17Cdc42 lines (Fig. 2B). Western analysis showed comparable total cellular vinculin abundance in the parental and Rho-protein

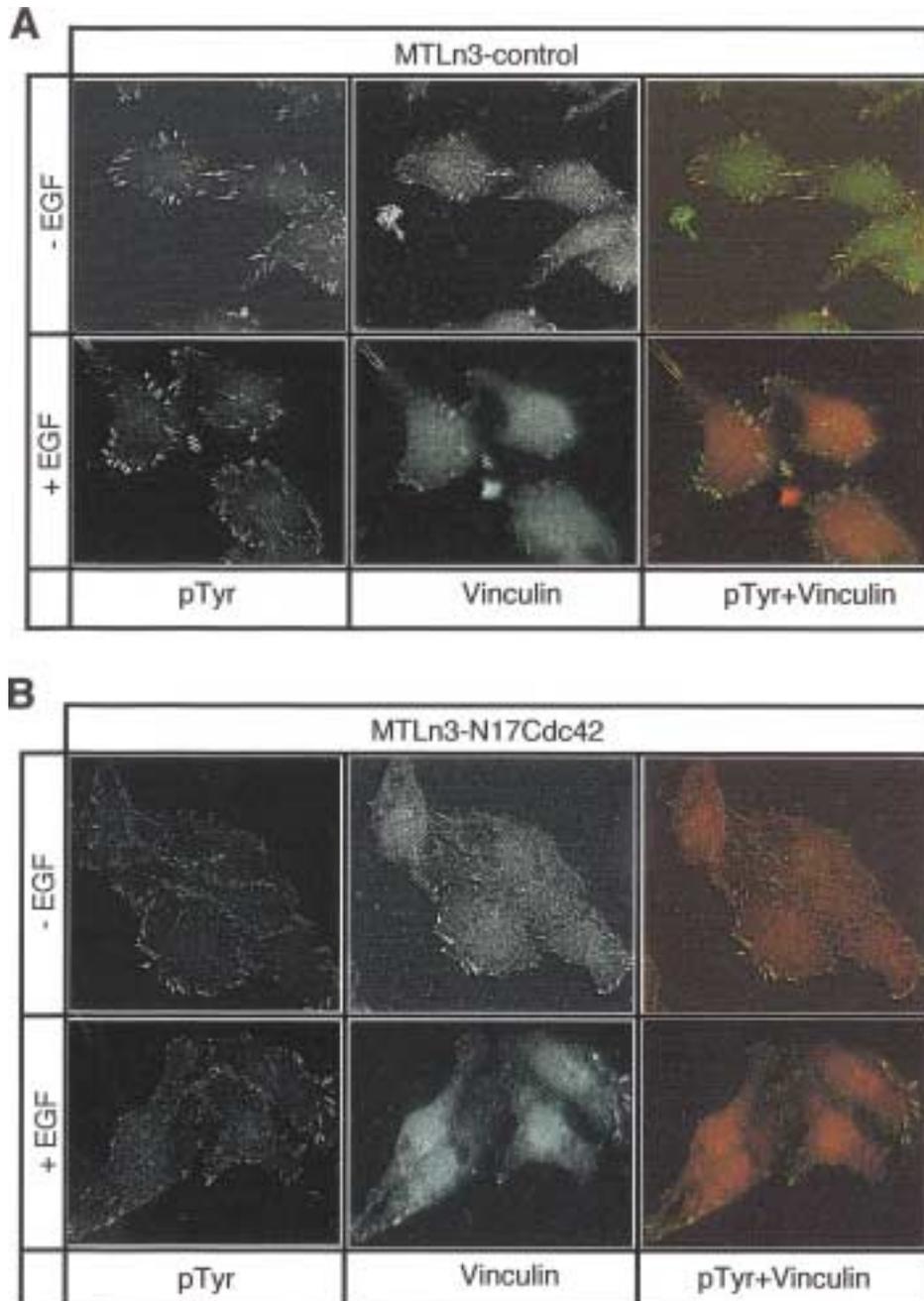


Fig. 2. Phosphotyrosine immunofluorescence at focal adhesions of MTLn3 Rho family mutant stable lines. Cells were seeded onto 22 mm square collagen-coated glass cover slips and treated with EGF (5 nM) for 4 h. Immunostaining for pTyr and Vinculin were recorded using a cooled CCD camera and the images merged (PY + Vinc). Results are shown for MTLn3-control (A), MTLn3-N17Cdc42 (B), MTLn3-N17Rac1 (C), and MTLn3-N19RhoA (D). Inhibition of pTyr IF is found in (B), and (D).

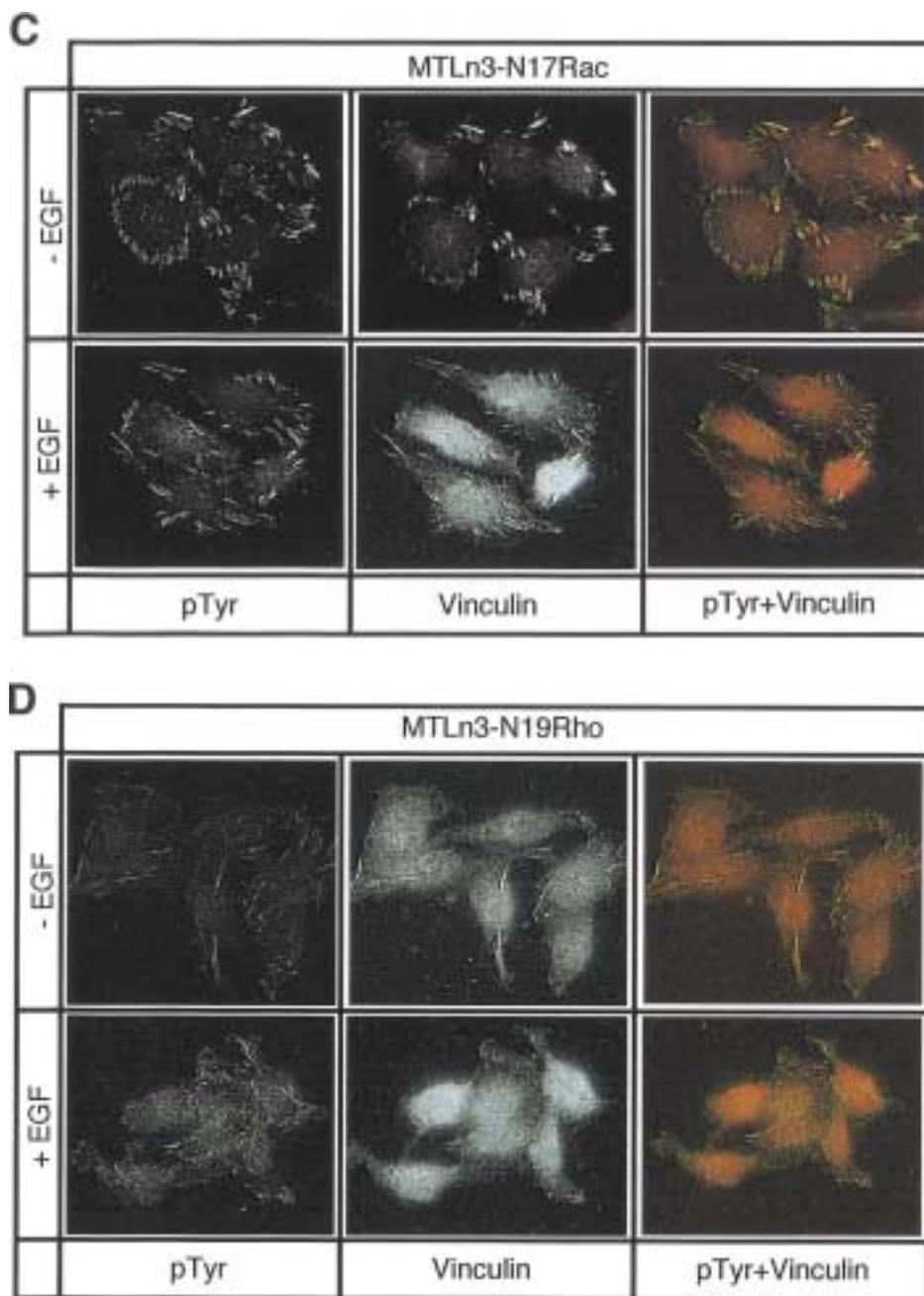


Fig. 2. (Continued)

expressing cell lines (data not shown). These studies suggest that both Rho and Cdc42 activity contribute to the assembly and formation of tyrosine phosphorylated FCs in mammary adenocarcinoma cells and are consistent with the observations that tyrosine phosphorylation and assembly of FCs involves RhoA activity (44).

EGF-induced Chemotaxis

In recent studies using T47D mammary epithelial cells, activating mutants of Cdc42 or Rac1 promoted

the motile phenotype of these cells (12). The chemotactic response of MTLn3 cells to EGF corresponds with enhanced lamellipod formation (22). We therefore examined the role of the Rho GTPases in EGF-induced guided chemotaxis of MTLn3 cells expressing Rho GTPases using microchemotaxis chambers and increasing concentrations of EGF (Fig. 3). The dominant inhibitory RhoA protein reduced EGF-mediated cell migration by 80%, both at 1 or 10 nM EGF. EGF-induced migration of MTLn3 cells was also reduced by 50% to 60% by

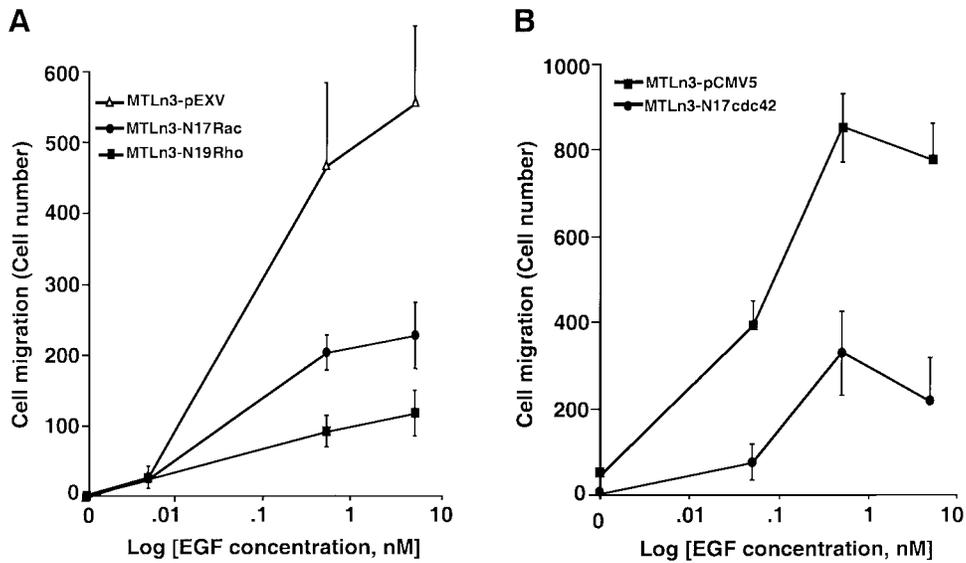


Fig. 3. EGF-induced chemotaxis. Chemotactic responses to EGF were assessed in a 48-well microchemotaxis chamber as described in Materials and Methods. Cell migration was determined for the Rho family dominant inhibitory cell lines, with comparison made to the empty vector control line. The results are mean \pm SEM from 8 and 5 separate experiments, (A and B) respectively.

the dominant inhibitory Rac1 (Fig. 3A) and Cdc42 (Fig. 3B) proteins compared with empty vector control lines.

Cell Proliferation and Anchorage-independent Growth in MTLn3 Expressing Inhibitory Rho GTPases

The proliferation of MTLn3 cell lines was measured over 6 days in culture as described in Materials and Methods (Fig. 4). Cell proliferation rates were inhibited by 50% using the dominant inhibitory N17Cdc42 and N17Rac1. In contrast, inhibition of cell proliferation using the N19RhoA mutant was less than 20%, although stress fiber formation was inhibited and tyrosine phosphorylation of focal adhesions was disrupted. These studies suggest that among the Rho family proteins, RhoA regulates the actin cytoskeleton but has modest effects only on cellular proliferation in the MTLn3 cells.

To extend these studies on cell growth, we analyzed contact-independent growth of these cells in soft agar. Previous studies suggested an important role for Rho family proteins (Rac and Rho) in Ras- and Erb-B2-induced anchorage-independent growth of Rat1 cells (25,47,48). Compared with cells stably containing the empty vector (pEXV or pCMV5) which formed large colonies in soft agar (Fig. 5B), the colonies of N17Rac1-MTLn3 and N17Cdc42-MTLn3 were significantly reduced in both size and number (Fig. 5A,B). In contrast, there was no significant reduction in the number or size of N19RhoA foci in soft agar (Fig. 5A,B). Western blotting confirmed the sustained expression of the Myc epitope-tagged RhoA in N19RhoA-MTLn3 cells.

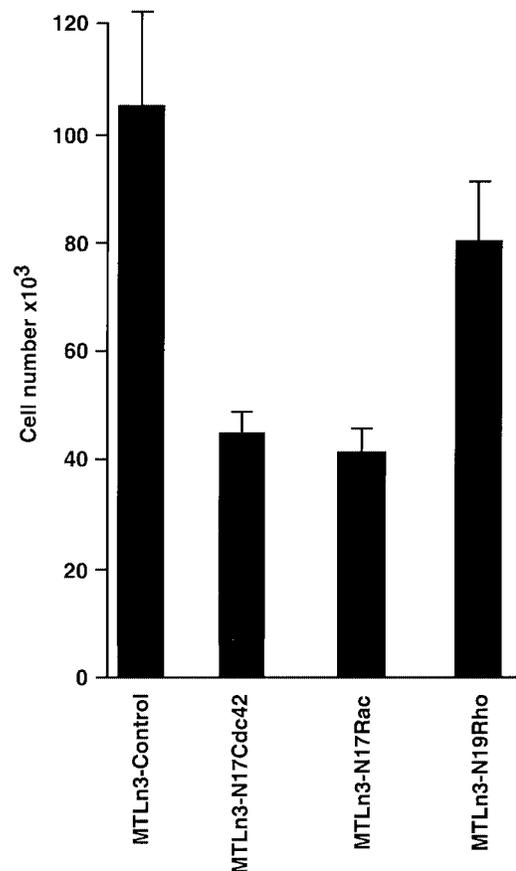


Fig. 4. Cell proliferation rates in MTLn3 inhibitory Rho mutant lines. The MTLn3 cell lines or the lines containing the empty vector controls (pEXV, pCMV5) were plated in 6-well tissue culture plates. After 6 days growth in regular medium (α -MEM, 5% FBS) cells were trypsinized, harvested and counted using a hemocytometer. The data are shown as mean \pm SEM.

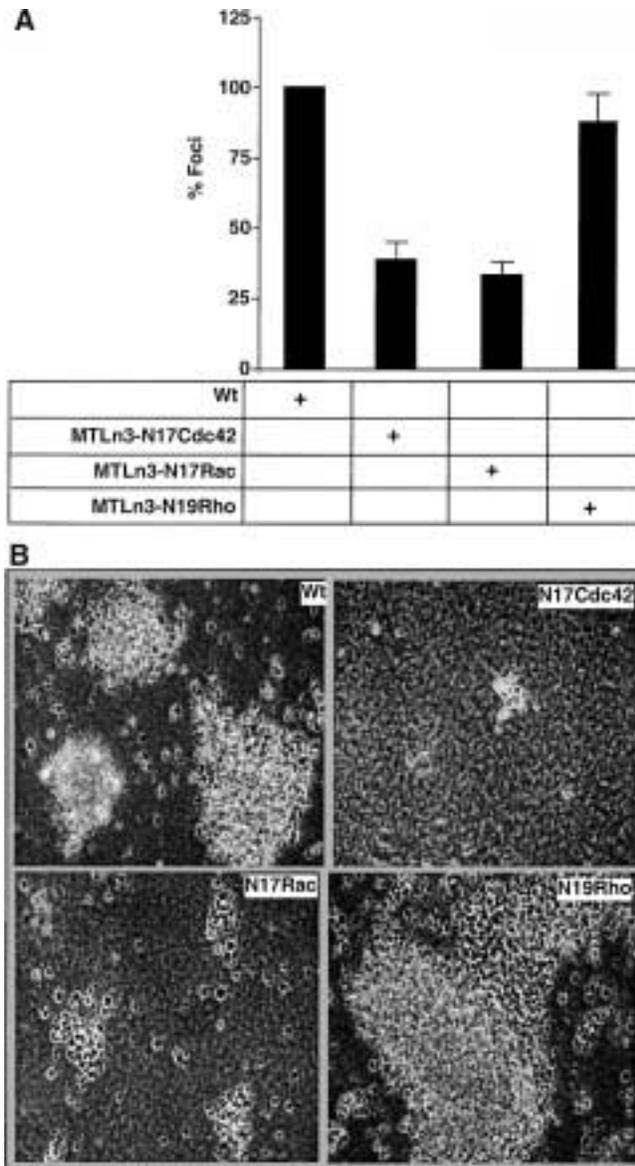


Fig. 5. Dominant inhibitory Rho family proteins reduce contact-independent growth. N17Rac1-, N19RhoA- and N17Cdc42-MTLn3 stable cells, or vector control MTLn3 lines, were grown in 6-well plates were assessed for growth on soft agar, as described in Materials and Methods. After 15 days, the plates were examined under a microscope at low magnification (4 \times or 6 \times) and the colonies were counted. (A) Experimental values represent the average number of foci in the 60 mm plates for each experimental condition; error bars represent the observed SEM for the 4 plates done in 3 separate experiments (B) Representative regions of plates.

Cell Spreading Assays

Previous studies have identified a role for Rho family proteins in Rat1 fibroblast spreading on fibronectin-coated surfaces (49). To determine the effects of dominant negative N19RhoA, N17Rac1 and N17Cdc42 expression on actin dynamics in MTLn3 cells, the rate of cell spreading on collagen was assessed at various time points and F-actin staining was assessed

simultaneously. F-actin staining was followed using rhodamine phalloidin in serum starved MTLn3 cells. F-actin containing structures were present in serum starved parental MTLn3 lines. Expression of N17Rac1 inhibited cell spreading of MTLn3 cells on collagen by 90%, compared to vector control cells at 45 min. This inhibition of cell protrusion and spreading by N17Rac1 was maintained at 140 min (Fig. 6A) and 310 min (Fig. 6B). MTLn3 cells expressing N19Rho or N17Cdc42 showed similar cell spreading as the parental controls (Fig. 6C). Similar trends were observed in 3 separate clones of each dominant negative Rho family protein. These results suggest an important role for Rac in MTLn3 cell protrusive activity and spreading on collagen.

Effects of Rho GTPases on the Tumorigenicity, Blood Burden and Lung Metastasis of MTLn3 Cells

These experiments showed that the three Rho family proteins performed distinguishable roles in regulating growth, morphology and motility in MTLn3 cells. The MTLn3 cells form tumors in nude mice (50,51), however the molecular mechanisms and the role of the Rho family proteins in MTLn3 growth *in vivo* were unknown. Implantation experiments were therefore conducted in nude mice as described in Materials and Methods on 3 separate occasions. The growth of MTLn3 tumors *in vivo* at 5 weeks was reduced by more than 95% by N17Rac1 or N17Cdc42 and by 30- to 40% with the N19RhoA expressing cells (Table 1). These studies suggest that Rac and Cdc42 may be more important for sustained *in vivo* growth of MTLn3 cells than Rho.

MTLn3 cells can metastasize to the lung after intravenous injection or implantation (35,50,51). When subcutaneous implantation methods were used, tumor cell density in the blood correlated with lung metastasis, suggesting that efficient intravasation into the blood stream perhaps affects the genesis of the metastatic phenotype (35). To examine the role of Rho family proteins in MTLn3 cell metastasis, blood burden assays were performed after implantation of the stable MTLn3 Rho family dominant inhibitory lines into nude mice as recently described (35). Blood burdens were assessed either 5 weeks after implantation (Table 1), or after tumors were allowed to grow to the same size, at which point blood burden assays was determined (Fig. 7B). The number of circulating cells observed with each of the dominant negative Rho protein lines was reduced in number compared with wild type (Table 1). Western blotting of the colonies evidenced the presence of sustained expression of the epitope tags within the peripheral blood colonies (data not shown). To begin to distinguish effects of the Rho proteins on tumor growth, from effects on intravasation, tumors were grown to the same size and blood burden reassessed (Fig. 7B). When the Rho dominant inhibitory lines were grown for

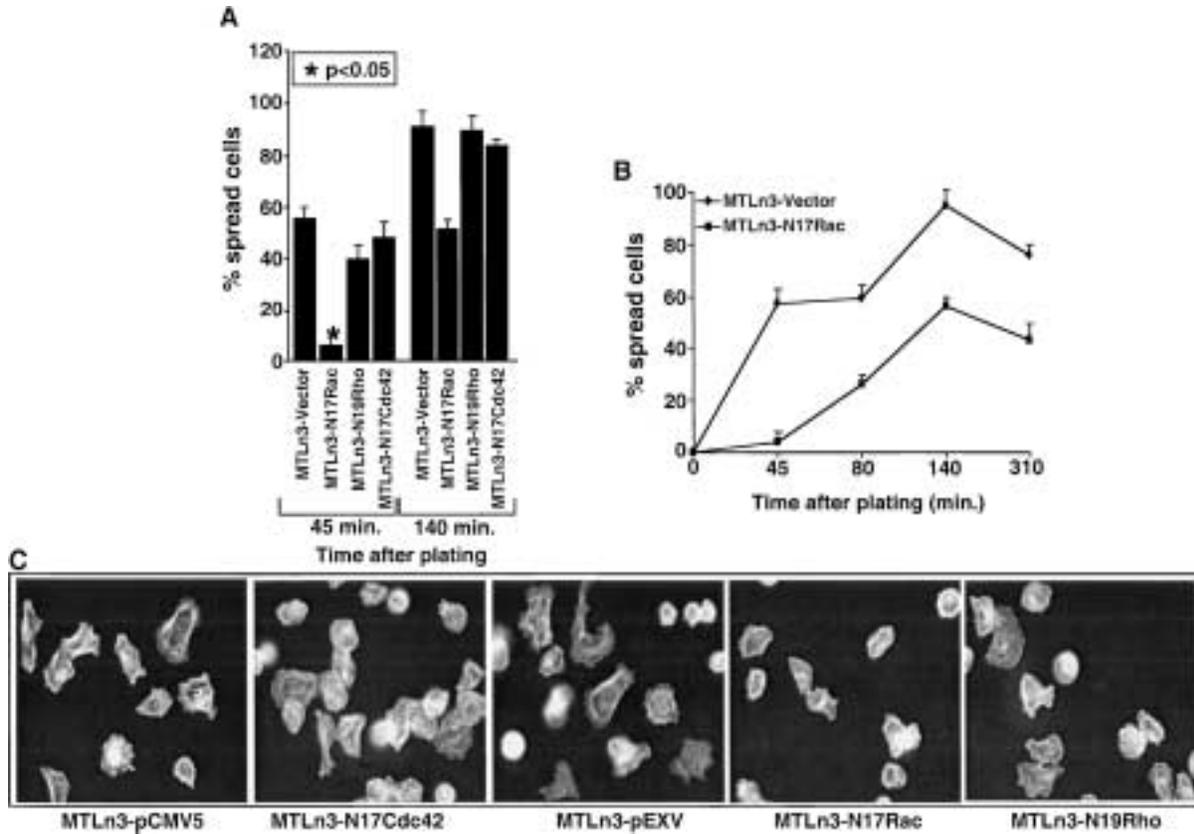


Fig. 6. Cell spreading. Parental and dominant inhibitory Rho protein-expressing MTLn3 cells were seeded onto 22 mm square collagen-coated glass cover slips and incubated in tissue culture plates. Cells were then fixed, permeabilized and stained with FITC-phalloidin at various time points. Spread cells were counted at 45 min and 140 min for all preparations (A) and also at 80 min, 140 min and 310 min for N17Rac1-expressing lines and their vector controls (B). The data are shown as the means \pm SEM for percentage positive cells in the 5 fields. Representative fields at 80 minutes (C).

8.5 weeks to the same size as vector control lines at 5 weeks, blood burden was detectable but reduced compared to vector control for each of the Rho proteins (Fig. 7B). These findings suggest that each of the Rho proteins contribute to intravasation.

A quantitation of lung metastasis was next performed in animals implanted with the MTLn3 vector control and Rho dominant negative cell lines. At 5 weeks post implantation, all MTLn3 control cells (pEXV, pCDNA3) had formed widespread lung

Table 1. Tumorigenicity and blood burden in nude mice. The inhibitory Rho family MTLn3 lines were assessed for growth 5 weeks after implantation into nude mice. The mean tumor size \pm SEM of at least 3 separate experiments is shown. Colony forming assays were performed on the peripheral blood of nude mice 5 weeks after implantation with MTLn3 lines encoding the dominant inhibitory Rho proteins or control lines.

	5 Weeks Growth		
	Tumor Size Diameter (cm) (Mean \pm SEM)	Cells in Blood (range)	Lung Metastasis (Mean \pm SEM)
MTLn3-pCMV5	1.36 \pm 0.13	78–80	26 \pm 4.74
MTLn3-N17Cdc42	0	0	0
MTLn3-pEXV	1.57 \pm 0.23	92–100	21 \pm 1.56
MTLn3-N17Rac	0	0	0
MTLn3-N19Rho	0.49 \pm 0.06	16–25	0

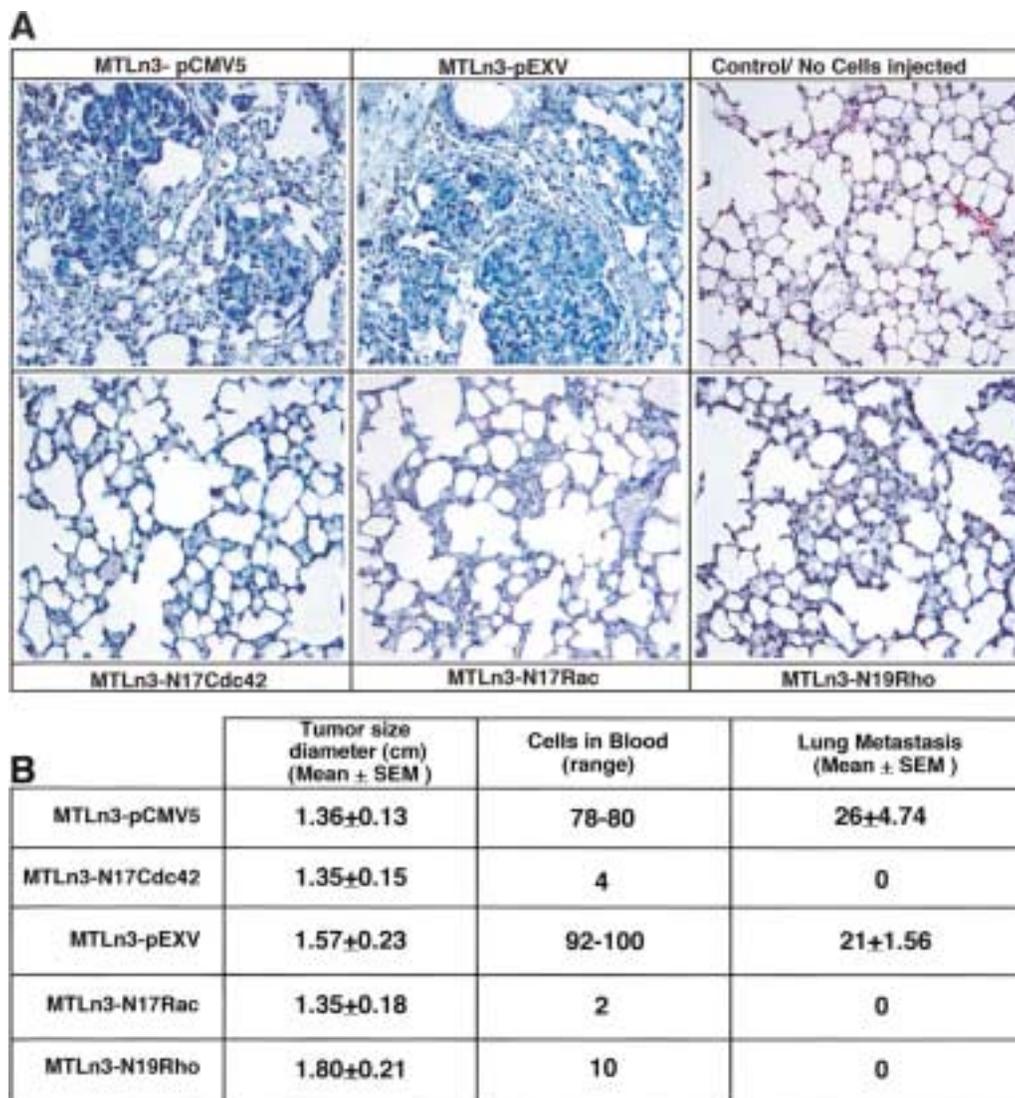


Fig. 7. Blood burden and lung metastasis in nude mice. MTLn3 Rho inhibitory tumors were grown to the same size in the tissues of mice and CFU and lung metastasis assessed. (A) Representative examples of the lung tumors stained with H&E are shown. (B) MTLn3 Rho inhibitory tumors were grown to the same size in the tissues of mice and blood burden (CFU) and lung metastasis were assessed, as described in Materials and Methods. Results are from at least 3 separate experiments under each condition.

metastatic foci (Fig. 7A). In contrast, none of the cell lines expressing dominant inhibitory Rho GTPases formed detectable macro foci, even though the N19RhoA lines were able to produce detectable tumor burdens in the peripheral blood.

Discussion

The current studies determined the role of specific Rho GTPase family members in mammary epithelial cell growth, guided migration and tumor metastasis (Table 2). and thereby identified the collaborative interactions between these family members for the full metastatic phenotype *in vivo*. The role of the Rho family proteins and their hierarchy of interaction in

breast epithelial tumor cell function are not well understood. In addition, a systematic analysis of each of the diverse roles of Rho proteins had not been performed previously, especially in relation to understanding the fundamental biology of mammary tumor metastasis *in vivo*. Rac was the dominant regulator of cell spreading and both Rac and Cdc42 were important in MTLn3 cellular growth. The Rho family proteins regulate diverse signaling pathways controlling a large number of essential functions, including the organization of the actin cytoskeletal, cell migration, gene transcription and cellular growth. In the current studies inhibition of either Rac, Rho or Cdc42 signaling blocked intravasation and lung metastasis suggesting that each of these

Table 2. Summary of the role of Rho GTPases derived from analyses of the dominant inhibitory MTLn3 lines. The ability to regulate function completely (>90%, triple arrow), partially (40–70%, double arrow), modestly (10–30% single arrow) or not detected (none).

	cdc42N17	RacN17	RhoN19
Focal contacts and tyrosine phosphorylation	Disruption ↓	↑	Disruption ↓
Protrusion/Spreading	↓	↓↓↓	↓
EGF-guided chemotaxis	↓ 50%	↓ 50%	↓ 80%
Cellular proliferation	↓ 60%	↓ 60%	↓ 20%
Primary tumor growth	↓↓↓	↓↓↓	↓ 60%
Colony formation (CFU) (cells in blood)	↓↓↓	↓↓↓	↓↓
Lung macro-metastasis	none	none	none
Foci formation (Soft agar)	↓ 70%	↓ 75%	↓ 10%

GTPases independently contribute to the final metastatic phenotype *in vivo*.

Several Rho family members have been shown to be essential for Ras transformation (1). Previous studies have identified specific roles for Rho family proteins in cultured cells, although cell type-specific differences have been observed. Different models have been proposed for the relationship between Cdc42, Rac and Rho. Hierarchical interactions have been identified between Rho family members, providing evidence for a “cascade” model (52), regulating the cytoskeleton in Swiss 3T3 fibroblasts, with Cdc42 activating Rac, which in turn activates Rho (7). In contrast with this model, an “antagonistic model” has been proposed in which Cdc42/Rac and Rho function in an antagonistic manner. For example Rac can downregulate Rho activity (53) and inactivation of Rho in neuroblastoma cells leads to neurite extension which is inhibited by either Cdc42-N17 or Rac1N17 (54). A “convergent” pathway model was proposed to define interactions between Rho proteins in several other cell types (49), for example, during integrin-mediated signaling (49,55,56). In fibroblasts, several studies suggested that Ras, Rac, and Rho function in a cascade relationship of interdependency (7), while other studies suggested “convergent” (49,57) or perhaps “independent” pathways to explain the relationships between these proteins in other cell types. Finally, the Rho GTPases have important transcriptional functions in which Rac, Rho and Cdc42 act largely independently of each other (1). Although the current studies suggest that Rac, Rho and Cdc42 each contribute independently to the metastatic phenotype *in vivo*, further studies are required to distinguish whether the Rho GTPases function in a “cascade” or parallel pathway in mammary epithelial cells.

The role of Rho family members in defining mammary tumor cell morphology was assessed by

immunofluorescence of the MTLn3 cytostructural proteins in the stable dominant negative lines. Rho and Cdc42 contributed to the co-localization of pTyr and vinculin without altering the total abundance of vinculin in the cell. pTyr was observed in a more dispersed pattern in cells expressing dominant negative RhoA (Fig. 2D vs. 2A). The loss of pTyr and vinculin co-staining in the N19RhoA MTLn3 cells is consistent with observations that quiescent cells with low Rho activity adhere and spread on ECM, but do not form well developed focal adhesions (58,59). Focal contacts are one form of cell adhesion in cultured cells and contain anchor and cytoskeletal molecules including vinculin, paxillin and talin (36,38), together with signal transduction molecules such as focal adhesion kinase (FAK) (39).

The assembly and tyrosine phosphorylation of FCs depends upon actomyosin contractility and is regulated by cytoplasmic factors, including Rho, or microtubular integrity (37,40,42–45,60). FCs exhibit characteristically high levels of tyrosine phosphorylation (36). Vinculin, a major structural component of FCs, co-localizes at cell-matrix adhesions with pTyr in both primary human fibroblasts (61) and in the MTLn3 cells. Rho-induced FC formation requires a functional cellular contractile apparatus. Inhibiting intracellular contractile forces promotes disassembly and prevents FC formation even though cells remain adherent to the substratum (44,62). Rho is required for complete phosphorylation of FAK induced by matrix adhesion (49), although FAK is phosphorylated at multiple sites and some Rho-independent FAK phosphorylation also occurs upon integrin aggregation (63). The observation that inhibition of Rac signaling did not interfere with pTyr and vinculin co-localization is consistent with findings that FAK phosphorylation and paxillin tyrosine phosphorylation are Rho-dependent and Rac-independent in Rat1 cell fibroblasts (49).

The reduction in vinculin staining at FCs in the N17Cdc42 lines suggests that Cdc42 activity contributes to FC formation in MTLn3 cells as previously described in fibroblasts.

The current studies demonstrated a critical role for Rac1 in MTLn3 cell spreading and protrusion. These findings of diminished cell spreading in MTLn3 cells expressing dominant negative Rac1 are consistent with studies of spreading on fibronectin which was shown to be selectively Rac-dependent in T lymphocytes (64) and in fibroblasts (49). The directional migration of Rat1 cells towards PDGF-BB, LPA or fibronectin was blocked by N17Rac1 and not by dominant negative mutants of RhoA or Cdc42 (11,13). Inhibition of spreading by the dominant inhibitory Rac1 was associated with a modest increase in the number of FCs which were PTyr containing. These findings are consistent with observations in NR6 fibroblasts in which FC disassembly was associated with EGF-enhanced migration into an acellular area (46). Rac1 is activated by a variety of tyrosine kinase receptors including platelet-derived growth factor (PDGF), EGF and insulin (65). The adhesion and spreading of T cells provides a more streamlined shape reducing shear imposed upon them by vascular flow and may have analogous functions for mammary epithelial cells. The current studies in MTLn3 mammary adenocarcinoma cells, together with previous studies in Swiss 3T3 and endothelial cells, suggest that the role of Rac1 in cell spreading is well conserved between cell types.

Guided-cell migration in response to a chemotactic gradient involves processes that are distinct from random protrusions and motility (66,67). During guided chemotaxis, focal adhesion disassembly occurs both at the trailing and the leading edge. MTLn3 cells demonstrate classical amoeboid chemotaxis on a planar surface in response to EGF (24). Herein, dominant inhibitory Rac1 and Cdc42 reduced EGF-induced guided chemotaxis 50%, and N19RhoA by 90%. Rho function is necessary to establish classical focal adhesion allowing stress fibers to anchor to the extracellular surface (59). Smaller adhesion complexes at the periphery of the cell are induced by activation of Rac and Cdc42 (7). As the relative abundance of the N19RhoA and N17Rac1 in the stable lines was similar by western blotting, and Rac preferentially regulated MTLn3 cellular spreading, these findings suggest that individual GTPases play distinct roles in spreading compared with guided chemotaxis.

Analysis of Rho family proteins in anchorage-independent growth demonstrated that Cdc42 and Rac function contributes to mammary adenocarcinoma cellular proliferation and growth in soft agar. Although Rho played a dominant role in guided chemotaxis, the contribution of Rho to cellular proliferation and contact-independent growth of the MTLn3 cells assessed in soft agar assays was relatively modest. This is similar to the situation in fibroblasts

(47,48,68–71). The induction of anchorage independence by oncogenic Ras however requires Rho, Rac and Cdc42 (47,48,71) and inhibition of Rho as well as Rac and Cdc42 can inhibit G₁ phase progression in other cell types (72). The PAK kinases may serve as Rac and Cdc42 effectors involved in integrin-mediated ERK activation and integrin-mediated adhesion is required for efficient coupling of Rac1 to PAK1 (73). Overexpression of activated Rac and Cdc42 can bypass this need and induce anchorage-independent ERK activity (69,74) and expression of the cyclin A gene, a key regulator of S phase progression (68–70). The enhancement of G₁ phase progression and cellular growth by Rac appears to be strongly related to the induction of the cyclin D1 gene. Activating mutations of Rac promote DNA synthesis and activate the cyclin D1 promoter directly (28,75) and inhibition of Rac or Cdc42 block cyclin D1 expression and contact independent growth induced by oncogenic Neu (25). The modest effect of blocking Rho on MTLn3 growth in soft agar, compared with Rac1N17 and Cdc42, may reflect the relatively greater importance of Rac/Cdc42 in contact-independent growth of mammary adenocarcinoma cells.

The process of metastasis to the lungs involves several independent processes including migration, invasion of tissues, intravasation and growth. Inhibition of Rac, or Cdc42 blocked growth of MTLn3 cells in nude mice significantly better than N19RhoA at 5 weeks. When tumors were grown to a similar size, each of the dominant inhibitory Rho family members significantly reduced the number of colonies detectable in the peripheral blood by more than 90%. Invasion of tissues may involve induction of proteases through Rho family protein-dependent mechanisms including induction of cytokines such as IL-1 α which thereby activates collagenase-1 gene expression (76).

It is clear from a number of studies that Rho family proteins play important but distinguishable roles in cell growth and invasion of cellular matrix (1). Although Rho proteins had been postulated to regulate cellular metastasis, the requirement for Rho proteins in regulating breast cancer metastasis *in vivo* using a heterologous system had not previously been determined. Dominant negative mutants of Rho family proteins block serum-induced DNA synthesis in Swiss 3T3 cells (72). Rat1 transformation induced by activating mutants of ErbB-2 is also blocked by dominant inhibitory Rho proteins (25). Despite the importance of the EGF receptor and its related family members to mammary epithelial cell growth, the role of Rho proteins in EGF receptor-dependent growth is poorly understood, particularly in the mammary epithelium. MTLn3 cells proliferate in response to EGF or serum (77), and inhibition of either Cdc42 or Rac, but not Rho activity, significantly reduced cellular proliferation (Fig. 5) and growth in soft agar (Fig. 6). An activating mutant of Rac1, but not Cdc42, enhanced cell growth in the absence of serum in Rat1 cells (48)

further suggesting that each RhoA protein family member regulates growth in a distinct manner. Our studies using MTLn3 cells suggest an important role for Rac and Cdc42 in MTLn3 cellular proliferation and growth in soft agar.

Taken together, the *in vitro* studies in MTLn3 cells showed that Rho family GTPases are differentially employed in signaling cell morphology changes, motile functions and aspects of growth that correlate with tumorigenesis and metastasis. Primary mammary tumor size was reduced at 5 weeks by each of the dominant inhibitory Rho family proteins. When tumors were grown to the same size, the reduced intravasation assessed by blood burden, suggested that entry into, and survival in, the peripheral blood is regulated by each of the Rho GTPases examined and is not simply a function of reduced tumor mass. These findings are consistent with recent studies in which RhoC was shown to play an important role in the metastatic behaviour of melanoma cells (78). The events governing entry into the peripheral blood and survival may include resistance to shear forces, deformability, apoptosis in the absence of substratum and intravasation at the primary site (35). While the role of the Rho family proteins in these events remains to be determined, the N17Rac1 MTLn3 cells were found to have enhanced sensitivity to osmolar stress (RGP, BB, unpublished). In conclusion these experiments establish, for the first time, a critical role for Rho proteins in the regulation of mammary tumor growth and blood burden.

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