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**Autocrine laminin-5 ligates \(\alpha_6\beta_4\) integrin and activates RAC and NF\(\kappa\)B to mediate anchorage-independent survival of mammary tumors**

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Abstract
Invasive carcinomas survive and evade apoptosis despite the absence of an exogenous basement membrane. How epithelial tumors acquire anchorage independence for survival remains poorly defined. Epithelial tumors often secrete abundant amounts of the extracellular matrix protein laminin 5 (LM-5) and frequently express α6β4 integrin. Here, we show that autocrine LM-5 mediates anchorage independent survival in breast tumors through ligation of a wild-type, but not a cytoplasmic tail–truncated α6β4 integrin. α6β4 integrin does not mediate tumor survival through activation of ERK or AKT. Instead, the cytoplasmic tail of β4 integrin is necessary for basal and epidermal growth factor–induced RAC activity, and RAC mediates tumor survival. Indeed, a constitutively active RAC sustains the viability of mammary tumors lacking functional β1 and β4 integrin through activation of NFκB, and overexpression of NFκB p65 mediates anchorage-independent survival of nonmalignant mammary epithelial cells. Therefore, epithelial tumors could survive in the absence of exogenous basement membrane through autocrine LM-5–α6β4 integrin–RAC–NFκB signaling.

Keywords
mammary epithelial cell, β4 integrin, apoptosis, GTPase, microenvironment

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Autocrine laminin-5 ligates α6β4 integrin and activates RAC and NFκB to mediate anchorage-independent survival of mammary tumors

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Invasive carcinomas survive and evade apoptosis despite the absence of an exogenous basement membrane. How epithelial tumors acquire anchorage independence for survival remains poorly defined. Epithelial tumors often secrete abundant amounts of the extracellular matrix protein laminin 5 (LM-5) and frequently express α6β4 integrin. Here, we show that autocrine LM-5 mediates anchorage-independent survival in breast tumors through activation of NFκB, and overexpression of NFκB p65 mediates anchorage-independent survival of nonmalignant mammary epithelial cells. Therefore, epithelial tumors could survive in the absence of exogenous basement membrane through autocrine LM-5–α6β4 integrin–RAC–NFκB signaling.

Introduction

Malignant transformation is linked to the migration of transformed epithelial cells across the endogenous basement membrane (BM) and their survival and proliferation in the surrounding interstitial collagen-rich stroma. Because normal mammary epithelial cells (MECs) require ligation of BM receptors to grow and survive (Weaver et al., 1997), invasive breast cancers must be able to resist apoptosis. Consistently, endogenous apoptosis rates decrease as MECs transition from ductal carcinoma in situ to invasive carcinoma (Gandhi et al., 1998), and immortalized mammary tumor cells frequently exhibit apoptosis resistance (Fernandez et al., 2002). However, how transformed MECs acquire apoptosis resistance remains poorly understood.

Breast tumors and immortalized mammary tumor cells frequently lose expression of α3β1 and α2β1 integrins, which are the BM receptors that support normal MEC growth, differentiation, and survival (Weaver et al., 1996). Mammary tumors also express high amounts of antiapoptotic proteins such as activated focal adhesion kinase (Cance et al., 2000), NFκB (Sovak et al., 1997), bcl-2 (Kalogeraki et al., 2002), and survivin (Tanaka et al., 2000), and often have elevated levels of active prosurvival kinases such as ERK, PI 3, and AKT (Fukazawa et al., 2002). Therefore, apoptosis-resistant breast tumors could arise through selection of MECs that express sufficient quantities of antiapoptotic molecules that permit growth and survival in the absence of adhesion (anoikis; Frisch and Ruoslahti, 1997).

Primary human breast tumors frequently synthesize and secrete ECM proteins such as fibronectin (Ioachim et al., 1986). Despite much evidence that fibronectin promotes survival of epithelial tumors (Fukazawa et al., 2002), activation of ERK or AKT. Instead, the cytoplasmic tail of β4 integrin is necessary for basal and epidermal growth factor-induced RAC activity, and RAC mediates tumor survival. Indeed, a constitutively active RAC sustains the viability of mammary tumors lacking functional β1 and β4 integrin through activation of NFκB, and overexpression of NFκB p65 mediates anchorage-independent survival of nonmalignant mammary epithelial cells. Therefore, epithelial tumors could survive in the absence of exogenous basement membrane through autocrine LM-5–α6β4 integrin–RAC–NFκB signaling.

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Key words: mammary epithelial cell; β4 integrin; apoptosis; GTPase; microenvironment

Abbreviations used in this paper: 2D, two-dimensional; 3D, three-dimensional; β4Δcyto, tailless β4 integrin; β4WT, wild-type β4 integrin; BM, basement membrane; LM-5, laminin 5; MEC, mammary epithelial cell; rBM, reconstituted BM.
2002) and laminin-5 (LM-5; Davis et al., 2001). Malignant MECs often up-regulate α5 and αv integrins (Koukoulis et al., 1993; Pena et al., 1994) and retain α6β4 integrin expression (Davis et al., 2001). Ligation of α5β1 integrin by fibronectin (Nista et al., 1997) and α6β4 integrin by LM-5 (Bachelder et al., 1999) supports cell growth and survival. Thus, apoptosis-resistant mammary tumors could arise through increased growth and survival of MECs that possess enhanced autocrine ECM–integrin signaling, although such a possibility has yet to be investigated.

We have been studying the role of MEC–ECM interactions and apoptosis resistance in the pathogenesis of breast cancer using the HMT-3522 human tumor progression model (Weaver et al., 1996). Analogous to breast tumor progression in vivo, the early passage, nonmalignant cells in this series (S-1) require ligation of α2β1 or α3β1 integrin for their growth and survival, whereas their tumorigenic progeny (T4-2) are anchorage independent (Wang et al., 1998). Instead of dying, in the absence of β1 integrin ligation, the T4-2s revert to form polarized tissue structures that resemble nonmalignant acini (Weaver et al., 1997). We demonstrated that α6β4 integrin can mediate apoptosis resistance to exogenous apoptotic stimuli in three-dimensional (3D) tissues irrespective of growth and malignancy status if MEC tissues are polarized (Weaver et al., 2002). Because invasive breast tumors typically lose polarized tissue architecture (acini, ductal) and tumors often metastasize as isolated cells, we asked whether α6β4 integrin might also support the survival of mammary tumor cells lacking polar tissue structure, and if so, how. Here, we report that nonpolarized malignant MECs grown as 3D structures survive via α6β4 integrin, provided they synthesize and secrete sufficient quantities of LM-5 and activate RAC and NFκB.

**Results**

α6β4 integrin mediates anchorage-independent survival of mammary tumors

α6β4 integrin drives tumor invasion and migration (Mercuero and Rabinovitz, 2001), and mediates apoptosis resistance in polarized MEC acini (Weaver et al., 2002), suggesting that tumors that express α6β4 integrin could metastasize and acquire multi-drug resistance, provided they are able to recapitulate tissue polarity. Because tumor invasion requires loss of tissue integrity, and survival of individual and isolated clusters of tumor cells dictates metastatic efficiency (Wong et al., 2001), we investigated whether α6β4 integrin could also support the growth and survival of isolated tumor cells or disorganized tumor cell clusters. We used S-1 (nonmalignant) and T4-2 (tumorigenic) MECs from the HMT-3522 human mammary tumor progression model and investigated whether anchorage independence of the T4-2s depends on α6β4 integrin.

We found that total and cell surface β1 and β4 integrin expression are higher in the T4-2s compared with S-1s, but that α2 integrin levels remain constant (Fig. 1 A and Fig. 3 A; Weaver et al., 1997). We also determined that although isolated S-1s require β1 integrin ligation for growth and sur-

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**Figure 1. α6β4 integrin mediates anchorage-independent survival of mammary tumors.** (A) FACS® analysis of cell surface integrin levels showing increased β4 integrin in the T4-2s compared with S-1s (S-1, 265 vs. T4-2, 430) and similar amounts of α2 integrin. (B) Apoptotic labeling indices were calculated using the TUNEL assay in S-1s and T4-2s grown in rBM for 96 h in the presence of function-blocking mAb against β1 integrin (AIB2) and/or β4 integrin (ASC-3). Results are the mean ± SEM of at least three separate experiments. (C) Phase-contrast (c), conventional immunofluorescence (EGFP; c') and confocal Immunofluorescence microscopy of α6 integrin (Texas red; c’’), β4 integrin (EGFP; c’’’), and α6 and β4 integrin overlay (yellow; c’), as indicated by white and black arrows showing uniform inducible expression of tailless EGFP β4 integrin (T4 β4Δcyto) in T4-2s and clustering of α6β4Δcyto integrin at membrane adhesion plaques. Bars: (c and c’) 50 μm; (c’’-c’’) 20 μm. (D) Relative cell adhesion levels calculated using a fluorescence assay of control T4-2s (T4-2), and T4β4Δcyto cells showing comparable adhesion to rBM and LM-5 in both cell types. (E) FACS® analysis showing similar levels of the LM integrins β1, β4, α3, and α6 in T4-2 and T4 β4Δcyto cells. (F) Cell viability was calculated using the Live/Dead assay for T4-2 or T4 β4Δcyto cells grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin. (G) Soft agar assay results demonstrating that expression of the tailless β4 integrin (T4 β4Δcyto) inhibits anchorage-independent growth of T4-2s, so that infected cells behave like S-1 nonmalignant cells (S-1). Results for B, D, F, and G are the mean ± SEM of three to four separate experiments. (B and F) *, P ≤ 0.05. (G) **, P ≤ 0.01.
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Figure 2. β4 integrin overexpression does not induce anchorage-independent growth and survival of nonmalignant MECs. (A) Immunoblot analysis of total RIPA lysates for S-1 controls (S-1) and S-1 cells overexpressing a full-length β4 integrin (S-1 4WT) demonstrating increased expression of total β4 integrin in S-1 4WT cells. (B) FACS analysis of membrane β1 integrin and β4 integrin in S-1s and S-1 4WT showing elevated β4 integrin expression in the infectants (S-1, 260 vs. S-1 4WT, 650) and no effect on β1 integrin. (C) Cell viability was calculated using the Live/Dead assay for S-1 and S-1 4WT grown in rBM for 96 h with and without a function-blocking mAb to β1 integrin. (D) Percent apoptosis was calculated by scoring the number of caspase 3–positive cells for S-1 and S-1 4WT grown in collagen I for 96 h with and without function-blocking mAb to β1 integrin. (E) Soft agar assay results demonstrating that whereas malignant T4-2s exhibit anchorage independent growth and survival S-1s do not, even if they overexpress β4 integrin (S-1 4WT). Results for C–E are the mean ± SEM of three to four separate experiments. * P ≤ 0.05.

Increased β4 integrin does not induce anchorage-independent survival of nonmalignant MECs

Because altering β4 integrin activity had such a profound effect on tumor growth and survival, we asked whether overexpression of a β4WT would be sufficient to confer anchorage independence to nonmalignant MECs (S-1 cells). S-1 cells were infected with β4WT, and selected pooled populations of MECs expressing elevated levels of total (Fig. 2 A) and membrane-localized α6β4 integrin (Fig. 2 B) were used for experiments. Control S-1s grown in 3D reconstituted BMs (rBMs) died rapidly when β1 integrin–BM interactions were inhibited (Fig. 2 C; Weaver et al., 1997). However, S-1s overexpressing β4WT remained viable despite the absence of β1 integrin–BM interactions (Fig. 2 C), indicating that increased activity of α6β4 integrin can sustain the growth and survival of nonmalignant MECs. Yet, S-1s overexpressing β4WT died if β1 integrin signaling was inhibited when viability experiments were conducted in 3D collagen I gels where exogenous LM (α6β4 integrin ligand) was absent (Fig. 2 D). Moreover, S-1 4WT MECS failed to grow in soft agar (Fig. 2 E, compare S-1 4WT with S-1 control with T4-2). Therefore, a signal functionally linked to the cytoplasmic tail of ligated β4 integrin supports the survival of isolated and nonpolarized clusters of nonmalignant MECs grown in 3D.

Autocrine LM-5 mediates β4 integrin–dependent survival of mammary tumors

Breast tumors synthesize and secrete abundant quantities of LM-5 (Davis et al., 2001). Because we found that α6β4 integrin supports anchorage independence of T4-2s but not of S-1s, we postulated that malignant transformation is either associated with constitutive activation of α6β4 integrin or is linked to increased synthesis and secretion of autocrine LM-5. Consistent with the latter prediction, T4-2s have elevated levels of total β4 integrin (Fig. 3 A) and plasma membrane β4 integrin (Fig. 1 A), and synthesize and secrete more LM-5 than S-1s in 3D ECM gels (Fig. 3 B). Moreover, T4-2s colonies in soft agar are surrounded by copious amounts of LM-5 (Fig. 3 D), and when LM-5–α6β4 integrin interactions and β1 integrin ligation are simultaneously blocked (using function-blocking mAbs) in tumors embedded within a collagen I gel (in the absence of an exogenous α6β4 integrin ligand), T4-2s die (Fig. 3 C). Consistently, although S-1s and S-1 4WT cells embedded within 3D collagen I gels die when β1 integrin ligation is inhibited, S-1 4WT, but not control S-1s, grow and survive significantly better in the presence of exogenous purified LM-5 (Fig. 3...
D). Thus, tumors grow and survive in the absence of an exogenous BM if they synthesize and secrete sufficient quantities of LM-5 and up-regulate and ligate α6β4 integrin.

α6β4 integrin does not require ERK or AKT to mediate tumor survival

How does LM-5 ligation of α6β4 integrin induce tumor survival? α6β4 integrin can mediate the survival of tumor cells through activation of PI 3-kinase and AKT (Bachelder et al., 1999). Ligation of α6β4 integrin also activates ERK via SHC-dependent activation of ras (Dans et al., 2001), and ERK supports anchorage-independent survival in epithelial cells (Howe et al., 2002). Although treatment with PD98059 effectively repressed ERK activity (Fig. 4 B), T4-2 survival remained unaffected, even when β1 integrin ligation was simultaneously blocked (Fig. 4 A). Likewise, treatment of T4-2s with LY294002 inhibited AKT activity (Fig. 4 B), yet failed to compromise β1 integrin-independent growth and survival (Fig. 4 A). Indeed, concomitant inhibition of ERK, PI 3-kinase, and β1 integrin activity had no appreciable effect on tumor viability (Fig. 4 A).

Because AKT is an oncogene that can mediate adhesion-dependent survival in tumors (Hill and Hemmings, 2002), we directly tested the importance of AKT activity to α6β4 integrin–dependent survival by expressing a dominant-negative K179M AKT. Despite stable expression of high levels of HA-tagged dominant-negative AKT (Fig. 4 D), T4-2s remained completely viable (Fig. 4 C) irrespective of their β1 integrin ligation status, even when ERK activity was also inhibited (Fig. 4 C). Conversely, stable expression of the same dominant-negative AKT (Fig. 4 F) modestly but significantly compromised the viability of isolated S-1s embedded within rBM (Fig. 4 E). Consistently, stable overexpression of high levels of a constitutively active HA-tagged myristoylated AKT (Fig. 4 H, MyrAkt) only partially rescued S-1 survival when β1 integrin ligation was blocked (Fig. 4 G), whereas overexpression and ligation of α6β4 integrin was significantly more effective (Fig. 2 C). Thus, α6β4 integrin must be able to support MEC survival through pathways that are distinct from AKT and ERK.

β4 integrin mediates tumor survival through regulation of RAC

The Rho GTPase RAC is frequently overexpressed in tumors of the breast (Fritz et al., 1999), permits MEC growth in soft agar (Bouzahzah et al., 2001), and protects MDCK cells from anoikis (Coniglio et al., 2001). Because neither inhibition of AKT nor ERK kinase compromised T4-2 survival (Fig. 4, A–D), we asked whether α6β4 integrin mediated T4-2 survival via activation of the Rho GTPase RAC. We assayed for RAC activity and determined that both basal and EGF-induced RAC activity significantly correlated with levels of expressed and ligated α6β4 integrin. For example, T4-2s that express high α6β4 integrin also have increased basal and EGF-induced RAC activity relative to S-1s (Fig. 5, A and B, compare specific activity of RAC in T4-2 with S-1; and Fig. 5, C and D, EGF-stimulated RAC activity). Moreover, ablating α6β4 integrin function in the T4-2s by expressing the dominant-negative β4Δcyto, reduced both basal and EGF-induced RAC activity to S-1s (Fig. 5, A and B; and Fig. 5 D) RAC activity, and overexpression of B4WT in the S-1s cells led to an increase in both basal (Fig. 5, A and B) and EGF-stimulated RAC activity (Fig. 5 C).

To determine whether α6β4 integrin mediated anchorage-independent survival in T4-2s through RAC, we examined the survival phenotype of S-1 and T4-2s that expressed constitutively active and dominant-negative RhoGTPase mutants. Pooled populations of T4-2s stably expressing dominant-negative EGFP-N17 RAC (Fig. 6 A) had reduced RAC activity (unpublished data) and required β1 integrin ligation for survival (Fig. 6 B, despite high levels of BM-ligated α6β4 integrin (Figs. 1–3). S-1s that expressed low levels of endogenous β4 integrin (Figs. 1–3) no longer depended on β1 integrin activity for survival and were able to
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Growth in soft agar (Fig. 6, F and G) if they stably expressed the constitutively active c-myc-V12 RAC (Fig. 6 E). Because T4-2s stably expressing a dominant-negative EGFP N19 Rho retained their apoptosis-resistant phenotype (Fig. 6, C and D), we suggest that LM-5–ligated β4 integrin mediates mammary survival through RAC.

αβ4 integrin mediates tumor survival via RAC-dependent activation of NFκB

Having established a link between αβ4 integrin, RAC, and survival, we next sought to delineate the mechanism whereby RAC mediates MEC survival. RAC can activate NFκB p65 (Bouzahzah et al., 2001), and we showed that β4 integrin induces apoptosis resistance in acini through NFκB (Weaver et al., 2002). Upon investigation, we found that S-1s expressing c-myc-V12RAC had high amounts of nuclear NFκB (Fig. 7, A and B) and that treating reverted T4-2 acini, which exhibit constitutively active NFκB (Fig. 7, C and D), we suggest that LM-5–ligated αβ4 integrin mediates mammmary survival through RAC.

β1 integrin. Data indicate that although active AKT does significantly enhance anchorage-independent survival of nonmalignant MECs, it does not completely rescue S-1 viability. (H) Immunoblot analysis of HA in S-1 controls (Vector) or S-1s expressing an HA-tagged constitutively active myristoylated AKT (MyrAKT). All apoptosis data are the mean ± SEM of at least three separate experiments. **, P ≤ 0.01; ***, P ≤ 0.001.

Figure 4. αβ4 integrin does not mediate anchorage-independent survival in mammary tumors via ERK or AKT. (A) Percent apoptosis calculated by scoring the number of caspase 3–positive T4-2s grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin and treatment with 50 µM of the PI 3-kinase inhibitor LY 294002 (LY), 20 µM of the MEK inhibitor PD 98059 (PD), or vehicle (DMSO; Vehicle). (B) Immunoblot analysis of total and activated ERK and AKT in T4-2s grown in rBM with or without LY 294002 and/or PD 98059 treatment, was as described in A. (C) Percent apoptosis was calculated as described in A for T4-2 Vector control (Vector) and T4-2s expressing a dominant-negative AKT (DNAkt) grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin and treatment with the MEK inhibitor PD 98059 (PD) as indicated. (D) Immunoblot analysis of HA expressed in control T4-2s (Vector) and T4-2s expressing a HA-tagged dominant-negative AKT (DNAkt). (E) Percent apoptosis calculated as in A for S-1 control (Vector) and S-1s expressing a dominant-negative AKT (DNAkt) grown in rBM for 96 h. Note that expression of the dominant-negative AKT decreased survival of S-1s yet failed to compromise the viability of the T4-2s, even when β1 integrin and ERK activity were inhibited. (F) Immunoblot analysis of HA in RIPA lysates of S-1 controls (Vector) and S-1s expressing an HA-tagged dominant-negative AKT (DNAkt). (G) Percent apoptosis was calculated as in A for S-1 vector control (Vector) and S-1s expressing a constitutively active myristoylated AKT (S1 MyrAkt) grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin and treatment with PD 98059 (PD) as indicated. (H) Immunoblot analysis of HA expressed in control S-1s (Vector) and S-1s expressing a HA-tagged constitutively active myristoylated AKT (MyrAKT). All apoptosis data are the mean ± SEM of at least three separate experiments. **, P ≤ 0.01; ***, P ≤ 0.001.
nization appreciably, and eventually killed the T4-2 revertants (unpublished data). Because expressing a dominant-negative N19 Rho did not compromise the viability of T4-2 revertants (Fig. 6 D), whereas N17 RAC did (Fig. 6 B), we conclude that the Toxin A phenotype was likely due to inhibition of RAC.

Full-length β4 integrin but not a β4Δcyto permits nuclear translocation (Fig. 7, E and F) and activation (Fig. 7 G) of NFκB in T4-2s. Therefore, we predicted that if α6β4 integrin regulates NFκB activity via RAC, a constitutively active RAC (V12 RAC) should confer anchorage-independent survival to T4-2s expressing the β4Δcyto, and tumor viability should depend on NFκB activation. Consistently, T4-2s expressing both the β4Δcyto and a constitutively active RAC (β4Δcyto/V12 RAC) survived when β1 integrin–ECM interactions were blocked (Fig. 7 H), and tumor cells expressing both transgenes regained their ability to form colonies in soft agar (Fig. 7 I). Furthermore, anchorage-independent survival of the β4Δcyto/V12RAC-expressing T4-2s absolutely required NFκB activity (Fig. 8 B). Indeed, T4-2s expressing only the β4Δcyto died when β1 integrin function was blocked (Fig. 7 H and Fig. 1 F) and T4-2 β4Δcyto MECs failed to form colonies in soft agar (Fig. 7 I). Therefore, anchorage-independent growth and survival of T4-2s depends on a signaling pathway initiated through LM-5 ligation of α6β4 integrin that is transduced by RAC and that depends on NFκB activation.

NFκB activity is necessary and sufficient for anchorage-independent survival of MECs

NFκB is induced in the early stages of mammary involution and its activation is associated with enhanced MEC survival in culture (Clarkson et al., 2000). NFκB expression and activity are increased in mammary tumors (Sovak et al., 1997). We have shown that NFκB mediates resistance to chemotherapy, radiation treatment, and receptor-induced apoptosis (Baldwin, 2001; Weaver et al., 2002). To directly determine if α6β4 integrin-dependent activation of NFκB is essential for the survival of T4-2 revertants, we inhibited NFκB nuclear translocation and assayed for effects on β1 integrin–dependent survival. Incubation with the membrane-soluble peptide SN50 that specifically inhibits nuclear translocation of NFκB, but not the nonfunctional peptide SN50M, induced apoptosis in the control T4-2 and T4 β4Δcyto/V12RAC revertants, but had no effect on viability when β1 integrin was ligated (Fig. 8, A and B; unpublished data). Moreover, sequestering NFκB in the cytosol through expression of a mutant IkBα (IkBαM) also rendered the T4-2s anchorage dependent for their survival (Fig. 8 C). Therefore, our data indicate that LM-5 ligation of α6β4 integrin likely activates NFκB via a RAC-dependent pathway that acts upstream of IKKa/β kinases. If true, then we reasoned that constitutive activation of NFκB should render nonmalignant MECs anchorage independent for growth and survival. We addressed this possibility by assaying for integrin-dependent survival and anchorage-independent colony formation in S-1s that overexpressed NFκB. Consistently, we found that expressing an exogenous NFκB in S-1s led to constitutive nuclear localization of p65 (Fig. 8, D and E) and per-

Figure 6. RAC mediates anchorage-independent survival in MECs. (A) Phase-contrast (a) and immunofluorescence microscopy images (EGFP; a’ and a”) of low (a and a’) and high (a”) magnifications of EGFP-tagged N17 RAC showing uniform high expression of the stable, exogenously expressed dominant-negative RAC protein in T4-2s. Bars: (a and a’) 100 μm; and (a”) 20 μm. (B) Cell viability calculated by the Live/Dead assay in control T4-2s (T4-2) and T4-2s expressing the N17 RAC (T4 N17Rac) grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin. (C) Immunofluorescence microscopy images of EGFP (FITC) detected only in T4-2s stably expressing EGFP N19Rho. Bar, 20 μm. (D) Cell viability calculated by the Live/Dead assay in control T4-2s (T4-2) and T4-2s expressing the N19Rho (T4N19Rho) grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin. (E) Immunofluorescence microscopy images of c-myc (Texas red) detected only in S-1s stably expressing a myc-tagged V12 RAC. Bar, 20 μm. (F) Cell viability calculated by the Live/Dead assay for S-1 controls (S-1) and S-1s expressing a c-myc-tagged V12 constitutively active RAC (S-1 V12Rac) grown in rBM for 96 h with or without a function-blocking mAb to β1 integrin. (G) Soft agar assay results demonstrating that expression of exogenous V12RAC (S-1 V12Rac) renders nonmalignant S-1s (S-1) anchorage independent for growth and survival. Results shown in B, D, F, and G are mean ± SEM of three experiments. *, P ≤ 0.05; **, P ≤ 0.01.
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Mitigated S-1s to form viable colonies in soft agar (Fig. 8 G) and to grow and survive in the absence of β1 integrin ligation (Fig. 8 F). Thus, in the absence of ECM adhesion NFκB can sustain MEC survival.

**Discussion**

We used paired nonmalignant (S-1) and tumor cells (T4-2) from the HMT-3522 tumor progression model and 3D agarose, collagen I, and rBM assays to investigate whether anchorage-independent growth and survival of mammary tumors depends on autocrine LM-5 ligation of α6β4 integrin. We found that malignant transformation is associated with up-regulation of α6β4 integrin and increased LM-5 secretion, and that ligation of overexpressed full-length but not a β4Δcyto, in combination with autocrine LM-5, is necessary and sufficient to induce anchorage-independent growth and survival in MECs, even in the absence of a polar tissue structure. Our results are consistent with the idea that MECs that secrete sufficient quantities of LM-5 and retain α6β4 integrin become selected during malignant transformation because they are able to grow and survive in the absence of exogenous...
BM cues. Because apoptosis limits metastatic efficiency (Wong et al., 2001), and LM-ligated β4 integrin also supports epithelial migration and invasion (Russell et al., 2003) and mediates immune and multi-drug resistance (Weaver et al., 2002), our results could explain why metastatic breast tumors frequently express β4 integrin (Menard et al., 1994) and why patients that express both BM protein and α6β4 integrin have the worst prognosis (Tagliabue et al., 1998).

α6β4 integrin can support tumor survival through PI 3- and AKT kinase (Bachelder et al., 1999) and keratinocyte proliferation through ERK (Dans et al., 2001), and β4 integrin can activate NFκB through AKT and ERK (Bozinovski et al., 2002). Yet, we found that α6β4 integrin activates NFκB and mediates MEC survival through RAC, and not through AKT or ERK. One plausible explanation for the discrepancy between our results and those published by others is that we conducted our experiments in the context of 3D malleable gels. In contrast to cells grown as two-dimensional (2D) monolayers on rigid, planar substrates, cells embedded within 3D malleable gels more accurately recapitulate normal and malignant tissue organization and behavior (Jacks and Weinberg, 2002). For example, MECs grown to form tissue-like structures (acini) in 3D BM gels are able to differentiate and optimally synthesize β casein in response to lactogenic hormones (Roskelley et al., 1994). Likewise, salivary epithelial cells form acini that express cystatin only in the context of a 3D BM gel (Royce et al., 1993), and keratinocytes recapitulate epidermal differentiation, including filagrin expression, most efficiently when grown as 3D organotypic rafts (Javaherian et al., 1998). Furthermore, MMP1 significantly enhances tumor growth in 3D, but has no effect on cell proliferation in 2D (Hotary et al., 2003); and RAC is required for cyst polarity in MDCKs grown within 3D collagen gels, but has no effect on MDCK polarity when cells are grown on 2D planar, rigid membranes (O’Brien et al., 2001). Why cells behave differently when grown on a planar, rigid substrate versus a 3D malleable gel remains an open question. What is known is that fibroblasts do not assemble focal adhesions containing αvβ3 integrin and activated focal adhesion kinase in response to a 3D ECM, but do so when plated on top of a 2D matrix (Cukiernian et al., 2001). Moreover, MECs cultured on 2D planar substrates transiently activate MAP kinase in response to EGF, whereas MECs grown within 3D gels to form acini do not (Wang et al., 1998); and polarized mammary structures grown within 3D gels are recalcitrant to a diverse array of apoptotic stimuli, whereas MECs spread on a 2D planar substrate remain sensitive (Weaver et al., 2002; unpublished data). Thus the composition of integrin adhesions and integrin signaling function appear to be differentially regulated in 2D and 3D, implying that α6β4 integrin may regulate epithelial survival by different mechanisms in 2D and 3D.

The Rho GTPases, RAC and Rho, are overexpressed in tumors (Fritz et al., 1999), and RAC enhances tumor invasion in culture (Keely et al., 1997) and supports breast tumor metastasis in vivo (Bouzahzah et al., 2001). We found that in MECs, EGF stimulation of RAC depends almost entirely on β4 integrin. Because LM-5 and α6β4 integrin are so often retained in primary breast tumors (Tagliabue et al., 1998; Davis et al., 2001), our results offer a plausible explanation for why RAC and NFκB activity are frequently elevated in these same malignant tissues (Sovak et al., 1997; Fritz et al., 1999). Moreover, by establishing a functional link between RAC and NFκB in 3D tissues, our findings could explain how RAC supports mammary tumor growth in soft agar (Bouzahzah et al., 2001) and why RAC supports the viability of cells actively migrating into 3D collagen gels (Cho and Klemke, 2000). Finally, our data predict that α6β4 integrin could drive tumor metastasis through an alternative PI 3-kinase and Akt-independent mechanism (Mercurio and Rabinovitz, 2001).
Current theory maintains that anoikis is circumvented early during malignant transformation (Frisch and Ruoslahti, 1997) and that metastatic cells are selected thereafter from the invasive tumor population through pressures exerted by the tumor tissue microenvironment (Wouters et al., 2003). Yet, tumor metastasis can occur early during cancer (Wasserberg et al., 2002); metastatic cells have been found in the bone marrow of patients with early stage tumors (Menaard et al., 1994), and tumor cells do circulate in the blood of patients with benign disease (Hardingham et al., 2000). Metastatic tumors frequently express integrins such as αv, α5, α6, and β4, and often secrete ECM proteins including collagen IV, LM-5, and fibronectin (Davis et al., 2001; Ioachim et al., 2002). Tumor metastasis and extravasation are facilitated by integrin–ECM interactions (Clezard, 1998). Therefore, it is plausible that apoptosis-resistant metastatic tumors arise early during malignancy through selection of transformed cells that express ECM proteins and retain integrins that support migration, invasion, and survival. Because we show that malignant transformation is linked to autocrine LM-5, that LM-5 supports cell survival by inducing α6β4 integrin–RAC–NFκB signaling, and that LM-5–ligated α6β4 integrin and RAC support epithelial motility and invasion (Russell et al., 2003), our data underscore the feasibility of this concept.

Materials and methods

Materials

We used commercial EHS matrix (Matrigel; Collaborative Research) for the rBM assays; Nitrogen 100 (bovine skin collagen I; Celtrix Laboratories); 3 mg/ml and 10 μg/ml of affinity-purified LM-5 (Russell et al., 2003) for coating culture dishes; and 0.3% Collagen Solution AC-5 (ICN Biomedical Products) for the 3D collagen I assays. Primary antibodies were as follows: LM-5, rabbit sera pκκ1 and clone BM165 (Russell et al., 2003); α6 integrin, clone CoH3 (BD Biosciences); β1 integrin, clones AlIB2 (provided by C. Damsky, University of California, San Francisco, San Francisco, CA), and TS2/16 (ATCC); β4 integrin, rabbit sera, and clones 3E1, ASC-3, and ASC-8; and α2 integrin, clone 10G11 (all from Chemicon International); kBA/ MAD-3, clone 25, and NfκB p65, rabbit sera (Santa Cruz Biotechnology, Inc.) and clone 20 (BD Biosciences); cytokeratin 18, clone RCK106 (BD Biosciences); RAC1, clone 102 (BD Biosciences); c-myc, clone 9E10 (OncoResearch Research Products), and AKT and Phospho-ser473/474-AKT; ERK1, rabbit sera (BD Biosciences), phosphoERK1/2 (Th202/Yr204), rabbit sera (New England Biolabs, Inc.); activated caspase 3, rabbit sera (Cell Signaling), and HA.11, clone 16B12 (Babco). Secondary antibodies were as follows: horseradish peroxidase, and biotinylated mouse IgG (Vector Laboratories); FITC, and Texas red–conjugated secondary antibodies. Nuclei were counterstained with DAPI (model Fluoroskan Ascent Fl; LabSystems).

Cell culture

The HMT-3522 MECs were grown in 2D and embedded (0.5–0.8 mmg/ml) were blocked (1 h; 0.1% BSA), incubated (60 min, 37°C), washed (3× PBS), incubated with 4 μM calcein (20 min, RT), and quantified using a fluorescence plate reader (model Fluoroskan Ascent Fl; LabSystems).

Anchor-independent assay

Anchor-independent growth was assessed using a soft agar assay (Wang et al., 1998). In brief, 20,000 cells were plated in 1 ml DME/Ham’s F12 containing 0.7% agarose, overlaid with 1 ml of 1% agarose, and 40-μm colonies were scored positive after 21 d.

Function-blocking studies

To inhibit integrin function or LM-5 binding, cells were incubated with mAbs against β1 integrin, clone 110B2 (1:25–1:100 ascites/ml ECM); β4 integrin, clones ASC-3 or ASC-8 (4–16 μg IgG/ml ECM); LM-5, clone BM165 (10 μg IgG/ml ECM); or IgG isotype-matched control mAb (4–16 μg IgG/ml ECM) at the time of embedding. To inhibit NfκB nuclear translocation, the active inhibitor NfκB SN50 or the inactive analogue NfκB SN50M was added directly to the media.

Immunofluorescence analysis

Cells were directly fixed using 2–4% PFA or 100% methanol, and samples were incubated with primary mAbs, followed by either FITC- or Texas red–conjugated secondary antibodies. Nuclei were counterstained with DAPI (Sigma-Aldrich). Cells were either visualized using a scanning confocal laser (model 2000-MP; Bio-Rad Laboratories) attached to a fluorescence microscope (model Eclipse TE-300 [Nikon] or model MDIRBE [Leica]). Confocal images were recorded at 120× and conventional images were recorded at 40–60×.

Apoptosis assay

Apoptosis was assayed by the Live/Dead Assay (Molecular Probes) or by detection of internucleosomal DNA fragmentation in fixed cells using an in situ TUNEL assay (Boehringer) or via immunodetection of activated caspase 3. Percent death was calculated as cells positive for ethidium bromide expressed as a percentage of the total number of live cells scored positive by calcein staining (FITC). The apoptotic labeling index was calculated as the percentage of total cells positive for FITC-labeled 3′OH DNA ends, and percent apoptosis was determined as the percentage of total cells positive for activated caspase 3. The minimum number of cells scored was 200–400 per experimental condition. Cell death by apoptosis was confirmed by showing that DNA cleavage or caspase 3 activity could be inhibited by prior treatment with the caspase inhibitors YVAD CHO or DEVD-CHO (1 μM, BIOMOL Research Laboratories, Inc.).

cDNA constructs

Full-length β4 prkK-5 (provided by F. Giancotti, Memorial Sloan-Kettering Cancer Center, New York, NY) was used directly. The 2,710-bp EcoRI-BglII fragment from the β4 prkK-5 construct was ligated with the EcoRI-BamHI vector fragment of pEGFP-N2, and an EcoRI–NotI fragment containing the β4 integrin EGF fusion was subcloned into an EcoRI–NotI vector fragment of Hermes HRS puro-GUS (provided by H. Blau, Stanford Medical Center, Stanford, CA). Myc-tagged V12Rac1 (provided by A. Hall, University College, London, UK) was cloned as an EcoRI fragment into LZRS-ires-blasticidin, and N17Rac1 and N19RhoA (provided by E. Butcher, Stanford Medical Center) were cloned into the EGF fusion vector EGFP-C1 (CLONTECH Laboratories, Inc.), and excised and recloned into LZRS-ires-blasticidin by PCR using the EcoRI tagged primer GTpaseF, 5′-kbAa and 65 cloned into plZRS’ (provided by F. Khavari, Stanford Medical Center) were used directly. The BglII-BamHI fragment containing HA-tagged dominant-negative AKT (K179M) and the HindIII–EcoRI fragment containing the myristoylated HA-tagged AKT (provided by P. Tischler, Tufts University, Boston, MA) were subcloned into plZRS.

Gene expression manipulations

Anaphosphoretic retrovirus was produced in either modified 293 cells or in Phoenix amphotropic cells (provided by G. Nolan, Stanford Medical Center), and MECs were spin infected and selected using blastidin. MECs were transfectioned with full-length β4 prkK-5 and pcDNA3.1 plasmid vector DNA or vector plasmid alone using LipofectAMINE (GIBCO BRL), and selected using G418. S-1 β4 prK-5–transfected cells were enriched for increased membrane localized β4 integrin through differential adhesion to LM-5, and increased β4 integrin levels were verified by FACS® analysis.

Electrophoretic mobility shift assay

To prepare nuclear extracts, cells were washed (1× PBS) and homogenized in nuclear isolation buffer (10 mM Heps, pH 7.9, 10 mM KCl, 1

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mM EDTA, 1 mM EGTA, 1 mM DTT, and 1 mM Pefabloc SC) with an addition of Igepal to 0.5%. After incubation (10 min, 4°C), nuclei were isolated by centrifugation (1 min, 14,000 rpm, 4°C) and nuclear extracts were prepared by homogenization and incubation in nuclear extraction buffer (20 mM Hepes, pH 7.9, 420 mM KCl, 1.5 mM MgCl\(_2\), 20% glycerol, 0.5 mM DTT, 1 mM Pefabloc SC, and 10 µg/ml leupeptin), followed by centrifugation (15 min, 14,000 rpm, 4°C). Equal amounts of nuclear protein were used in the EMSA reaction with the NFκB consensus oligonucleotide sequence 5′-AGT TGA GGC GAC TTT CCC AGG C-3′. 32P-Labeled oligonucleotide (150,000 cpm) was incubated with 5 µg of nuclear extract and gel shift binding buffer (10 min, RT; Promega Gel Shift Assay System). p65 rabbit antisera were added after the binding reaction, and the mixture was reincubated (20 min, RT). Specificity of binding was tested using competition experiments in which 10-fold molar excess of nonlabeled oligonucleotide sequence was added to a binding reaction. Complexes were resolved in 4.5% polyacrylamide gels (TE buffer: 90 mM Tris, 90 mM boric acid, 10 mM EDTA, pH 8.0).

**Flow cytometry**

Cells were isolated, nonspecific binding was blocked (60 min Dulbecco’s PBS, 0.1% BSA) and incubated with saturating concentrations of primary mAb (1 h), washed three times with Dulbecco’s PBS, and labeled with FITC-conjugated goat immunoglobulin. Stained cells were washed three times with Dulbecco’s PBS and immediately analyzed on a FACScan (Becton Dickinson). All manipulations were conducted at 4°C.

**Immunoblot analysis**

Cells were lysed (RIPA buffer; 50 mM Tris-HCl, pH 7.4, 150 mM sodium chloride, 1% NP-40, 0.5% deoxycholate, 0.2% SDS containing 20 mM sodium fluoride, and 1 mM sodium orthovanadate, and a cocktail of protease inhibitors), and equal amounts of protein were separated on reducing SDSPAGE gels, immunoblotted, and detected with a ECL-Plus system (Amersham Biosciences). To assay for differences in total secreted LM-5, LM-PAK1 cDNA was immunoprecipitated from conditioned media, and protein from equal cells was separated on SDS-polyacrylamide gels, immunoblotted, and detected as above.

**RAC activation**

Cells were treated with vehicle or 20 ng/ml EGF and incubated for indicated times, washed (2× PBS), and extracted (G protein buffer; 25 mM Hepes, pH 7.5, 150 mM NaCl, 1% Igepal CA-630, 10 mM MgCl\(_2\), 1 mM EDTA, 10% glycerol, 1 mM Pefabloc SC, 10 µg/ml leupeptin, 10 µg/ml aprotinin, 1 mM sodium orthovanadate, and 1 mM sodium fluoride; 5–10 min). Lysates were centrifuged (10 min, 14,000 rpm), and supernatants were mixed with GST-PBD and incubated with glutathione-Sepharose beads (Amersham Biosciences) to assay for differences in total secreted LM-5, LM-PAK1 cDNA was immunoprecipitated from conditioned media, and protein from equal cells was separated on SDS-polyacrylamide gels, immunoblotted, and detected as above.

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**References**


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