Cytoskeletal responsiveness to progestins is dependent on progesterone receptor A levels

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Abstract

Changes in the cell cytoskeleton occur in cell transformation and recent data suggest the involvement of ovarian hormones, which are implicated in cancer development and progression. In human breast and endometrial tumors, there is disrupted expression of progesterone receptor (PR) isoforms and predominance of one isoform, usually PRA. PRA predominance is an early event in carcinogenesis, and in cancers is associated with poor clinical features. Overexpression of PRA in vitro causes altered progestin regulation of cell morphology, suggesting that PRA overexpression may provoke deleterious changes in cell functioning. This study aimed to identify pathways of cytoskeleton regulation responsive to progestins and to determine whether these are perturbed when PRA is overexpressed to the levels seen in cancers. Progestin treatment of PR-positive breast cancer cells caused increased cell surface area whereas after induction of a stably integrated PRA construct, cells became rounded and the cell surface was decreased. The effect of PRA induction on cell rounding was reversed by the anti-progestin RU38486. Altered tropomyosin (Tm) isoforms were implicated in these morphological differences, as there was a PRA-mediated alteration in Tm5 isoform levels, and transfection of Tm5a mimicked progestin-mediated cell rounding in PRA-overexpressing cells. Ezrin was redistributed from the membrane to cytoplasmic locations in the presence of progestin, and discrete focal localization was evident in cells with PRA predominance. Progestin effects on the cytoskeleton in PRA-overexpressing cells provide evidence for novel endocrine regulation of aspects of actin microfilament composition, suggesting that changes in the cytoskeleton known to be associated with cancer development and progression may be regulated in part by altered PRA expression which develops early in carcinogenesis.

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Introduction

There is well-described epidemiological, experimental and clinical evidence that ovarian hormones play a major role in the development and progression of female hormone-dependent cancers. Women without ovaries have a very low risk of breast and endometrial cancer, whereas cumulative exposure to ovarian hormones, particularly estrogen, is associated with increased risk (MacMahon & Feinlieb 1960). There is also evidence that women exposed to exogenous progesterone have increased breast cancer risk (Ross et al. 2000) and a recent randomized trial showed increased breast cancer risk in healthy postmenopausal women on combined estrogen–progestin hormone replacement therapy (Writing Group for the Women’s Health Initiative Investigators 2002). This is supported by evidence that ablation of nuclear progesterone receptor (PR) decreases mammary carcinogenesis in mice (Lydon et al. 1999), and suggests that progesterone may play a role in breast cancer development.

The mechanisms of progesterone involvement in breast cancer development are unknown, but there are recent data which have provided evidence for...
disrupted progesterone signaling in human breast and endometrial tumors. The nuclear PR is expressed as two proteins: PRB and PRA (Kastner et al. 1990), which have different transcripational activities (Gronemeyer 1991, Tung et al. 1993, Vegeot et al. 1993, McDonnell et al. 1994, Hovland et al. 1998, McGowan & Clarke 1999). PRA and PRB are co-expressed, at similar levels, in normal breast (Mote et al. 2002) and endometrial epithelial cells (Mote et al. 1999), implicating both isoforms in the known effects of progesterone in these tissues. Progression to malignancy in breast and endometrium is accompanied by progressively unequal expression of PRA and PRB, and a significant proportion of carcinomas express a predominance of one isoform, usually PRA (Arnett-Mansfield et al. 2001, Mote et al. 2002). This is associated with poor clinical outcomes: in endometrial cancer, PR isoform predominance, usually of PRA, is significantly more common in tumors of higher grade, indicating an association between PR isoform predominance and poor prognosis (Arnett-Mansfield et al. 2001). The loss of co-ordinated PRA and PRB expression is an early event in carcinogenesis and is evident in early pre-malignant lesions (Mote et al. 2002). The available evidence supports the view that progesterone signaling in normal target epithelial cells requires the concerted activity of co-expressed PRA and PRB, and that disruption of PRA and PRB expression results in altered biological features which may play a role in hormone-dependent cancers.

In vitro studies aimed at asking whether PRA overexpression seen in cancers is a consequence of the carcinogenic process, or plays a role in its development and/or progression have revealed that overexpression of PRA results in altered progestin regulation of cell morphology, suggesting that development of PRA overexpression may itself provoke deleterious changes in cell functioning. In cells overexpressing the PRA isoform but not in control cells, cells acquired rounded morphology and there was decreased adherence of cells to tissue culture flasks (McGowan & Clarke 1999). This suggests that the cell cytoskeleton is a target of progesterone action, particularly when PRA predominates, and, as cell rounding and loss of adhesion are hallmarks of invasive potential, that the endocrine regulation of the cytoskeleton may be implicated in hormone-dependent cancer.

The cytoskeletal system most critical in transducing information from the cellular environment is the actin–tropomyosin (Tm) microfilament system (Hall 1998), which is closely linked to morphological changes that occur within cells during development (Gunning et al. 1998b) and in the transformed phenotype (Leavitt et al. 1987a,b). Changes in the microfilament system have been linked with cell transformation and tumorigenicity (Leavitt et al. 1987a,b). Microfilament bundles or stress fibers are commonly reduced in size and number and show altered organization after transformation (Pollack et al. 1975). Disruption of stress fibers has been shown to correlate with other features of the transformed phenotype, including loss of contact inhibition, acquisition of anchorage-independent growth, and tumorigenicity (Edelman & Yahara 1976, David-Pfeuty & Singer 1980). Tumor promoters alter the cytoskeleton, demonstrating a relationship between the cytoskeleton and the normal mechanisms of cellular proliferation control (Takahashi et al. 1986, Pienta et al. 1989).

The mechanisms underlying the changes in the cytoskeleton that occur in cancer are not fully understood; however, in light of the progestin effects on the cytoskeleton in cells overexpressing PRA, it is likely that cross-talk between the cytoskeleton and endocrine signaling may be involved. The purpose of this study was to determine whether the actin–Tm microfilament system was regulated by progestins and to determine whether overexpression of PRA, which is an early event in carcinogenesis of hormone-dependent tissues, alters the progestin effects on these cytoskeletal components. The overall aim of these studies is to identify cellular signaling pathways which may lead to changes in cell morphology in response to hormones and contribute to poorer prognosis phenotypes in hormone-dependent cancers.

Materials and Methods

Plasmids and antibodies

Rat Tm5a and Tm5b cDNAs were excised from the β-actin expression vector (Qin & Gunning 1997) and cloned into the SalI site of the pHβApr-4(sig-) vector (pHβApr-4Tm5a and pHβApr-4Tm5b). Monoclonal antibodies against human PR (hPRα6 and 7) were isolated from...
hybridoma cultures (Clarke et al. 1987). The wsa-9d polyclonal antibody, produced in rabbit (Schevzov et al. 1997), detects Tms from exon 9d of the α-Tmα and β-Tm α genes, and has been described previously (Percival et al. 2000). Rabbit polyclonal anti-ezrin (Upstate Biotechnology, Waltham, MA, USA), anti-vinculin and anti-β-actin monoclonal antibodies (Sigma Biosciences, St Louis, MO, USA), and mouse monoclonal β-catenin antibody (BD Transduction Laboratories, Franklin Lakes, NJ, USA), were used for immunoblots and immunofluorescence.

Cell culture

The estrogen receptor (ER)-positive, PR-positive T-47D cells (Keydar et al. 1979) (E G Mason Research Institute, Worcester, MA, USA) were cultured in antibiotic-free phenol red-free RPMI 1640 medium and supplemented as described (Graham et al. 1995a). T-47Dp3’SS clones containing the p3’SS vector, encoding the lac repressor, and the T-47DhPRA cell lines containing both the p3’SS and inducible pOP13 hPRA plasmids were maintained in medium supplemented with hygromycin B (200 µg/ml) and G418 (200 µg/ml). T-47Dp3’SS, which are similar to wild-type T-47D cells and were used as controls, and T-47DhPRA cell lines have been described previously (McGowan & Clarke 1999). Cells were negative for mycoplasma contamination as determined using the Genprobe rapid detection system (Gen-Probe, Inc., San Diego, CA, USA). Cell stocks were passaged regularly to maintain exponential growth. For experiments, cells were seeded at a density of 2 × 10^4 cells/ml unless otherwise indicated. Twenty-four hours later (day – 1) the medium was supplemented with vehicle or with 10 mM isopropyl-β-thiogalactosidase (IPTG) (Gibco/BRL, Melbourne, Australia), to sequester the lac repressor and allow induction of PRA from pOP13 hPRA. On day 0 the medium was supplemented with the synthetic progestin ORG2058 (10 nM) (Organon Chemicals, Amersham Australia, Sydney, NSW, Australia) or vehicle only. Where indicated cells were also treated with the anti-progestin RU38486 (100 nM), or estradiol (10 nM), either alone or in combination with ORG2058 (10 nM).

Transient transfections

T-47DhPRA cells plated on Lab-Tek II RS glass chamber slides (Nalge Nunc International, IL, USA) at a density of 2 × 10^4 cells/ml were co-transfected with pHBsp-Air-4Tm5a (1 µg), or pHBsp-Air-4Tm5b (1 µg) and pEGFP (500 ng), using the Fugene transfection protocol (Boehringer-Mannheim, Indianapolis, IN, USA) (GFP = green fluorescent protein). Cells were treated with IPTG (10 nM) or vehicle for 24 h before treating with ORG2058 (10 nM) or vehicle for a further 72 h. All experiments were performed in triplicate.

Immunoblot analysis

T-47D p3’SS and T-47DhPRA cells were harvested and stored as described previously (Alexander et al. 1989). Cell pellets were thawed on ice in PEMTG buffer (Graham et al. 1995a) containing 0·4 M KCl and protease inhibitors (0·5 mM phenylmethylsulfonyl fluoride, 1·4 µM pepstatin A, bacitracin (100 µg/ml), 25 mM benzamidine, 86 µM leupeptin and aprotenin (77 µg/ml)). Cell extracts were prepared and 25 µg protein transferred to nitrocellulose as described (Graham et al. 1995a and references therein). Protein concentration was determined by the method of Bradford (Bio-Rad, Richmond, CA, USA). Blots were incubated with antibodies at saturating concentrations and goat anti-mouse and anti-rabbit immunoglobulins linked to horse radish peroxidase at 1:5000 (Bio-Rad). Immunoreactivity was visualized using a chemiluminescent method (ECL; Amersham Pharmacia Biotech, Piscataway, NJ, USA). The relative intensity, in the linear range of the film, of the immunoreactive bands was calculated after densitometric scanning of X-ray films and analysis using Imagequant software (Molecular Dynamics Inc., Sunnyvale, CA, USA). Results are expressed as the percentage of immunoreactivity compared with control.

Determination of cellular morphologies

For determination of cell shape after Tm5a or Tm5b transfection, the proportion of transfected (GFP-positive) cells with round morphology was determined as a proportion of all transfected cells in 20 fields at ×400 magnification. For determination of cell surface area, fluorescence and brightfield images of T-47D cells, stained with β-catenin antibody to delineate cell boundaries, were captured through FTTC filters using a SPOTII cooled CCD digital camera (Diagnostic

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Instruments, Inc., MI, USA) and fluorescence was visualized using an Olympus BX50 microscope. The surface area of the cells was measured using ImagePro plus software (MediaCybernetics, Silversprings, Marland, IL, USA) and analyzed using SPSS statistical software (SPSS Inc., Chicago, IL, USA).

## Results

### PRA predominance results in altered cell size

Progestin treatment of breast cancer cells expressing PRA and PRB resulted in growth inhibition and in adoption of a phase-dark, flattened morphology compared with control (Fig. 1A; a, b). Cell surface was greater in progestin-treated cells compared with vehicle-treated cells (Fig. 1B; \(P < 0.001\), one-way ANOVA). PRB was predominant over PRA in these cells and the ratio of PRA to PRB was around 0.3 (not shown). Induction of PRA, such that the PRA/PRB ratio increased to 3, caused no change either in cell morphology or in cell surface compared with control cells (Fig. 1A, e; Fig. 1B; \(P = 0.26\), one-way ANOVA). However, upon treatment with progestin, PRA-overexpressing cells experienced a marked change in cell morphology. Cells were phase-bright and rounded (Fig. 1A; f) and the cell surface was significantly smaller in progestin-treated PRA-overexpressing cells compared with progestin-treated controls (Fig. 1B; \(P < 0.001\), one-way ANOVA). The cell rounding observed on progestin treatment of cells overexpressing PRA was observed in a range of T-47DhPRA clones with different total PR levels (not shown), demonstrating that the effect seen upon PRA induction was not due to an overall increase in PR levels. In support of this, previous studies have shown that PRA predominance resulted in progestin-mediated effects on gene expression which were related not to total PR levels but to the ratio of PRA to PRB (McGowan & Clarke 1999).

### Progestin specificity of cell morphological changes

The anti-progestin RU38486 decreased cell numbers similarly to progestin (data not shown), but unlike progestin had no effect on cell surface (Fig. 1A: compare panel b with c; Fig. 1B). RU38486 treatment abrogated the progestin-mediated increase in cell surface (Fig. 1A: compare panel b with d; Fig. 1B). Cells overexpressing PRA and treated with RU38486 had the same appearance as control RU38486-treated cells (Fig. 1A: compare panel g with c). However, combined treatment with progestin and RU38486 reversed the progestin-mediated cell rounding and decreased cell surface in PRA-overexpressing cells (Fig. 1A, compare f and h; Fig. 1B) supporting the conclusion that these effects were mediated via the PR.

To determine whether the changes in cell morphology were specific to progestins when PRA was overexpressed, or whether activation of a related nuclear receptor could also produce cell rounding, cells were treated with estrogen to activate the ER pathway. Cells numbers were increased after 72 h of treatment with estradiol, as expected in these cells, but there was no alteration in cell morphology (Fig. 2E and F) as was seen in parallel cultures treated with progestin with or without PRA induction (Fig. 2C and D). ER is expressed in the T-47DhPRA N5 clones and this was confirmed by immunoperoxidase staining of cells grown on coverslips (data not shown).

### Progestin regulation of focal adhesion complex proteins and \(\beta\)-catenin expression in T-47DhPRA cells

To determine whether the reduced surface area caused by progestin in PRA-overexpressing cells was due to altered levels of cell interaction proteins, levels of candidate proteins were measured by immunoblot in T-47D and T-47DhPRA cells. Progestin treatment resulted in the well-described decrease in levels of PRA and PRB proteins (Fig. 3A) and the increased level of PRA after induction with IPTG is evident (Fig. 3A, compare lane 5 and 7). The levels of \(\beta\)-catenin, ezrin (Fig. 3B and C), \(\beta\)-actin, \(\gamma\)-actin, paxillin, talin or focal adhesion kinase (not shown) were unaffected by progestin treatment, either in T-47D cells or in control T-47DhPRA cells (Fig. 3B and C, lanes 1–6) or in cells overexpressing PRA (Fig. 3B and C, lanes 7–8). Vinculin levels were increased by ORG2058 treatment in parental and T-47DhPRA cells (Fig. 3D, compare lanes 1 and 2, 3 and 4, 5 and 6) and this increase was unaffected by induction of PRA (Fig. 3D, compare lanes 7 and 8 with 5 and 6).
The progestin induction of vinculin expression was slow and necessitated treatment times of several days to be maximally detectable by immunoblot.

Figure 2 Estrogen effects on cell morphology in T-47DhPRA cells. T-47 DhPRA (clone N5) cells were seeded in 75 cm² flasks at a density of 5×10⁵ cells. Cells were treated with IPTG (10 mM) or vehicle for 24 h then treated with ORG2058 (10 nM), 17β-estradiol (10 nM) or their combination. Photographs show the morphology of the cells 72 h later. (A) vehicle, (B) IPTG (10 mM), (C) ORG2058 (10 nM), (D) IPTG (10 mM) and ORG2058 (10 nM), (E) 17β-estradiol (10 nM), (F) IPTG (10 mM) and 17β-estradiol (10 nM). Original magnification ×400.

Figure 1 Progestin and anti-progestin effects on cell morphology in T-47DhPRA cells. T-47DhPRA cells were seeded in 25 cm² flasks at a density of 2×10⁴ cells/ml. Cells were treated with IPTG (10 mM) or vehicle for 24 h then treated with ORG2058 (10 nM), RU38486 (100 nM) or their combination. (A) Photographs a–h show the morphology of the cells 72 h later. (a) Vehicle, (b) ORG2058 (10 nM), (c) RU38486 (100 nM), (d) RU38486 (100 nM) and ORG2058 (10 nM), (e) IPTG (10 mM), (f) IPTG (10 mM) and ORG2058 (10 nM), (g) IPTG (10 mM) and RU38486 (100 nM), (h) IPTG (10 mM) and RU38486 (100 nM) and ORG2058 (10 nM). Original magnification ×400. (B) The surface areas of cells were measured using ImagePro software. All experiments were performed in triplicate and the data expressed as means±S.E.M., *significantly different from control (P<0·001, ANOVA).
Taken together, these data show that although there is progestin regulation of some adhesion proteins, there is no differential effect in control and PRA-overexpressing cells, at least at the level of steady-state protein expression. This suggests that the differences in progestin effects on cell morphology in cells overexpressing PRA did not result primarily from altered protein levels of membrane components of cell interaction complexes.

The progestin effects on cell surface were observed in clonal T-47DhPRA cell lines with varying PR levels and PRA to PRB ratios. The N3 and E3 clones have abundant PR expression but very different PRA to PRB ratios, which were respectively 1 and 0.3 (Fig. 3E). N4 cells have lower total PR levels with a PRA to PRB ratio close to 1 (Fig. 3E). Induction of PRA with IPTG and progestin treatment in each of these clonal cell lines resulted in the decreased cell surface shown in Fig. 1. The data in Fig. 3E also illustrate the finding that PRB predominance does not result in the cell surface changes seen upon progestin treatment of PRA-overexpressing cells. A number of T-47DhPRA cell lines, illustrated by the E3 clone in Fig. 3E, have a significant predominance of PRB, yet progestin treatment of cell lines with PRB predominance does not result in cell rounding (not shown), whereas progestin treatment after induction of PRA in the same cells causes a clear reduction in cell surface. This is true also for cells with total PRB levels within the range of total PRA levels seen in the cells with induced PRA.

β-actin and γ-actin were abundantly expressed in T-47D and T-47DhPRA cells, and no change in the expression levels of either cytoskeletal protein was detectable by immunoblot on progestin treatment of control cells or in cells overexpressing PRA (data not shown).

### Regulation by progestin of the actin-Tm family of proteins

The effects of progestin on the expression of the integral proteins of the microfilament system were determined by immunoblot analysis. Tm isoforms are thought to differentially regulate the biochemical and physical properties of microfilaments: specific Tm isoforms associate in particular microfilaments, which may have distinct cytoskeletal localization and function, and thus control specific features of cell shape (Gunning et al. 1998a).

The high molecular weight Tm1 encoded by the β-gene was not present in T-47D cells, nor were the Tm2 and Tm6 isoforms, encoded by the...
cells (Fig. 5C and E). Cells with overexpression
of PRA in the absence of progestin treatment had a similar appearance to control cells (Fig. 5, compare B with A). Progestin treatment of PRA-overexpressing cells caused a decrease in fibrous cytoplasmic bundles and a distinctly rounded morphology (Fig. 5D and F). Consistent with the rounded morphology, redistribution of Tm from the cytoplasm to the periphery of cells was noted (Fig. 5F, arrows).

Effect of exogenously transfected Tm5a and Tm5b on cell shape in T-47D cells

The different distribution of Tms in progestin-treated cells, depending on the relative levels of PRA and PRB, suggested that the actin–Tm microfilaments may be involved in the cell rounding observed in PRA-overexpressing cells. This possibility was supported by the dependence on PRA levels of the ratios of Tm5a and Tm5b as shown in Fig. 4. To determine whether the alteration in the ratio of Tm5a and Tm5b was implicated in the PRA-mediated changes in cell shape, expression vectors encoding Tm5a or Tm5b were transfected into T-47DhPRA cells. T-47DhPRA cells treated with IPTG to induce PRA, or with vehicle and/or ORG2058, were used as controls and transfected cells were identified by GFP fluorescence. The shape of transfected cells was scored and the proportion of round cells determined. Cells overexpressing PRA and treated with progestin showed an increased proportion of round cells, in line with the decreased surface area shown in Fig. 1, compared with progestin-treated control cells (Fig. 6: compare IPTG+ORG2058+ with IPTG−ORG2058−; P<0·0001, one-way ANOVA). There was no statistical difference in the proportion of round cells in control (IPTG−ORG2058−), progestin-treated (IPTG−ORG2058+) or PRA-overexpressing cells in the absence of ligand (IPTG+ORG2058−). Transfection with Tm5a, in the absence of PRA overexpression, resulted in the same proportion of round cells as observed in the progestin-treated PRA-overexpressing cells (Fig. 6: compare Tm5a with IPTG+ORG2058+; P=0·41 one-way ANOVA), demonstrating that expression of Tm5a resulted in an effect on cell rounding that was similar to that provoked by progestin treatment in PRA-overexpressing cells. The proportion of round cells was significantly greater in Tm5a-transfected

α-gene, detected. Tm3 and the alternatively spliced isoforms, Tm5a and Tm5b (Percival et al. 2000) were detected in T-47D cells. Tm3 was abundant in parental T-47D and in T-47DhPRA cells and its levels were unchanged by progestin treatment or by overexpression of PRA (Fig. 4A). In parental T-47DN and in T-47DhPRAN3 cells, ORG2058 treatment altered the relative expression of Tm5a and Tm5b, preferentially increasing the Tm5b isoform at the expense of the Tm5a isoform (Fig. 4A; compare lanes 1 and 2, and 3 and 4, and 5 and 6). Overexpression of PRA in T-47DhPRA cells diminished the progestin-induced increase in the level of Tm5b and this resulted in altered relative levels of Tm5b/Tm5a (Fig. 4A, compare lanes 7 and 8). These changes in Tm5 levels required prolonged exposure to progestins, as they were not evident at 24 h after treatment (not shown) whereas they were detectable 72 h after treatment. The progestin effects on relative Tm5 isoform levels was seen in all T-47DhPRA clones tested (hPRA inducible clones with different total PR levels: N3, B8 and E3 and their parental N, B and E cells lacking PRA inducibility are shown in Fig. 4), although the magnitude of the Tm5b/Tm5a ratio was different in each cell line. Progestin treatment caused an augmentation in the Tm5b/Tm5a ratio in the T-47DhPRA B8 and E3 clones, as well as the N3 clones shown in Fig. 4A, and in each cell line PRA induction resulted in an abrogation of the progestin effect on the Tm5b/Tm5a ratio, which in E3 and B8 cells was reduced to close to control levels (Fig. 4B and C). No effect of IPTG treatment was observed in the parental lines lacking hPRA inducibility (Fig. 4).

Effect of progestin on distribution of actin–Tm microfilaments in T-47DhPRA cells

Tm was diffusely expressed in the cytoplasm of control cells and was detected in poorly organized stress fibers. Strongly staining foci or actin-dense protrusions were observed at the periphery of some cells (Fig. 5A). Adhesion belts were not visible and cells displayed a granular appearance. Progestin treatment resulted in increased cell surface, as previously shown in Fig. 1 and fibrous cytoplasmic bundles (Fig. 5E, arrow). No distinct cellular borders were observed between individual cells (Fig. 5G and E). Cells with overexpression


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Figure 4 Progesterin effects on the expression of TM-5a and TM-5b in T-47D and T-47DhPRA cells. T-47D (-N, -B and -E) parental clones containing the p3′SS vector encoding the lac repressor and T-47DhPRA (-N3, -B8 and -E3) clones containing both the p3′SS and inducible pOP13 hPRA plasmids were treated with IPTG (10 mM) or vehicle. After 24 h cells were treated with ORG2058 (10 nM) or vehicle, harvested 72 h later and total cytosol protein prepared. Tm isoform expression was detected using a polyclonal antibody which detects all α-TM isoforms on immunoblots (25 µg cytosol protein/lane). Tm5b/Tm5a ratio was calculated densitometrically and presented as histograms. (A) Upper panel, immunoblot of -N and -N3 cells, lower panel, histogram of Tm5b/Tm5a ratio. (B and C) Histogram of Tm5b/Tm5a ratio in cell lines, -B and -B8 and -E and -E3.
cells than controls (Fig. 6: compare Tm5a with IPTG -- ORG2058 -- ; P<0.002, one-way ANOVA). Transfection of Tm5b did not have an effect on the proportion of round cells (Fig. 6: compare Tm5b with IPTG -- ORG2058 -- ; P=0.98, one-way ANOVA).

**Distribution of ezrin in T-47DhPRA cells**

The closely related ezrin–radixin–moesin (ERM) proteins function as general cross-linkers between actin filaments and the cell membrane and are implicated in regulation of cell morphology, adhesion and motility (Tsukita & Yonemura 1999).

Ezrin was the principal ERM protein detected in T-47D cells and low levels only of radixin and moesin were detectable immunohistochemically (data not shown), so ezrin detection formed the focus of ERM protein analysis in this study. No change in ezrin protein levels was observed on immunoblot analysis (Fig. 3) in progestin-treated T-47D cells with or without PRA overexpression, and immunohistochemical analysis to detect ezrin was carried out to document cell distribution of this molecule in progestin-treated cells. In the absence of progestin, ezrin protein staining was evident in densely staining foci at the periphery of cells in control and PRA-overexpressing cells (Fig. 7A and B). Some cytoplasmic foci were also present. Intracellular localization was confirmed after confocal microscope analysis of the fluorescently labeled sections (not shown). Progestin treatment
resulted in a redistribution of ezrin in control cells, being detected as a marked loss of focal staining at the cell periphery and an increase in cytoplasmic staining (Fig. 7C). Progestin treatment of cells overexpressing PRA resulted in ezrin distribution that was different from that seen either in progestin-treated cells or in control cells. Ezrin was located in discrete foci within the cytoplasm (Fig. 7D). Taken together, the results show that ezrin distribution is under progestin control in T-47D cells and that overexpression of PRA is associated with cytoplasmic focal localization. Given the role of ERM proteins in anchoring stress fibers to the membrane (Tsukita & Yonemura 1999), the sequestration of ezrin into large cytoplasmic foci only in cells overexpressing PRA suggests that their rounded cell morphology could result in part from the changed localization of ezrin in these cells.

Discussion

Ovarian hormones are involved in cancer development and progression and there is evidence that disruption in progesterone signaling, through altered expression of PR isoforms, is an early event in cancer development and leads to aberrant regulation of cell morphology. This study has demonstrated that altering the cellular ratio of PRA and PRB, to mimic the increased relative levels of PRA observed in cancers in vivo, markedly changed the nature of progestin regulation of cell morphology.

To determine the mechanisms through which progestin effects on cell morphology were mediated, levels and distribution were investigated initially of membrane-associated and cytoplasmic adhesion components. Overall, steady-state levels of membrane adhesion proteins, β-catenin and E-cadherin (not shown) either were unchanged by progestin treatment, or, in the case of the focal adhesion protein vinculin, were similarly regulated by progestin treatment in control and PRA-overexpressing cells. Integrin levels were similarly regulated by progestin in control and PRA-overexpressing cells (E M McGowan, S Saad, L J Bendall, K F Bradstock and C L Clarke, unpublished observations). In cell lines with similar levels of PRA and PRB, there is evidence of increased cell spreading and focal adhesions after progestin treatment (Lin et al. 2000, 2001) and this study provides support for progestin regulation of various membrane-associated and cytoplasmic adhesion components. However, as the effects of progestin on membrane components of cell adhesions were the same in cells overexpressing PRA, the differences in progestin effects on cell morphology in cells overexpressing PRA are unlikely to reside primarily at the membrane. The data presented in this study did not exhaustively examine cell distribution and activation states of all of these components, and it may be that there are PRA-specific effects on phosphorylation or cell distribution of some membrane-associated and cytoplasmic adhesion components. This possibility will be explored in future investigations.

Progestin regulation of cell morphology was shown to involve actin microfilaments and evidence for this was provided by the effects of progestin on Tm isoforms. Relative levels of Tm5a and Tm5b were hormonally regulated: in progestin-treated
control cells Tm5b was elevated at the expense of Tm5a whereas in PRA-overexpressing cells Tm5a and Tm5b were expressed at similar levels. Selective degradation of the Tm isoforms could account for this observation. However, as these Tm isoforms are alternatively spliced products of the same gene (Goodwin et al. 1991), these data suggest that PRA overexpression decreases the progestin-mediated elevation in Tm5b and moreover suggest the previously unsuspected progestin sensitivity of the Tm splicing process. The regulation of Tm splice site selection is not understood although certain factors that appear to mediate alternate splicing in Tm genes have been described (Wieczorek 1988), and alternate Tm splicing has been shown to occur in response to nerve growth factor treatment (Weinberger et al. 1993) or in response to serum factors during myogenesis (Wieczorek 1988).

Transfection of Tm5a and Tm5b isoforms showed that disruption of the levels of these proteins alone was sufficient to mimic the progestin-mediated effects in PRA-overexpressing cells. Introduction of Tm5a provoked cell rounding in most of the transfected cells, in the absence of any hormonal treatment. That cell rounding was also seen upon progestin treatment of cells overexpressing PRA, but not control cells, implicates the Tm5 isoforms in the cell rounding observed in PRA-overexpressing cells and this is further supported as mentioned above by the differential effects of progestin on Tm5 isoform levels seen on immunoblot analysis in PRA-overexpressing vs wild-type cells. The mechanisms underlying the hormonal regulation demonstrated in this study are unclear at present, but progestin regulation of α-Tm5 gene splice site selection, taken together with the known role of Tms in actin microfilament regulation (Gunning et al. 1998a), provides support for direct progestin regulation of actin filament dynamics in hormone-dependent cells.

The involvement of actin microfilaments in the cell morphology changes seen upon PRA overexpression was supported also by the altered localization of ezrin in PRA-overexpressing cells. While no changes in ezrin levels were observed on immunoblot analysis, it was clear that progestin treatment altered ezrin localization and moreover that PRA overexpression was associated with a different effect to that seen in control cells. Whereas progestin treatment in control cells caused a reduction in membrane staining of ezrin and increased cytoplasmic granular staining, ezrin localization in PRA-overexpressing cells was in distinct cytoplasmic foci. Current evidence suggests that ezrin anchoring to the membrane via CD44 is required for stress fiber attachment (Vaheri et al. 1997); the sequestration of ezrin in cytoplasmic foci in PRA-overexpressing cells suggests that this protein is unavailable for interaction with the cytoskeleton and is consistent with the relative lack of stress fiber-associated microfilaments in these cells. In contrast to the ezrin, levels of the focal contact protein vinculin were increased in PRA-overexpressing cells despite the decrease in cell surface area which may imply a decrease in focal adhesions. The implication is that vinculin is no longer attached to adhesion sites and that different cellular mechanisms control its targeting and expression when PRA is overexpressed. This study suggests therefore that multiple PRA-sensitive mechanisms appear to exist in order to regulate the expression and localization of cytoskeletal-associated proteins.

The mechanisms of progestin regulation of the cytoskeletal components investigated in this study are not known. Progestin is likely to be acting indirectly on membrane proteins such as ezrin, which were redistributed upon progestin treatment, without alteration in its cellular levels. Progestin effects on vinculin involved increased levels of the protein after prolonged treatment, which may imply transcriptional regulation of vinculin by indirect means. As discussed above, the mechanism of progestin regulation of Tm splice site selection is still to be determined although the delayed appearance of the progestin effect on Tm5 isoform levels suggests that it is likely to be indirect. Although delineation of the mechanisms of PR activity on the cytoskeletal pathways investigated here were beyond the scope of the current study, it is possible that PR modulation of previously described cell signaling pathways in the cytoplasm will be involved. PR, through interaction with SH3 domain of Src kinases, is able to activate MAP kinase and exert rapid regulation of cytoplasmic signaling pathways (Boonyaratanakornkit et al. 2001) although the delayed appearance of the effects documented in this study may argue against the involvement of a rapid MAP kinase-mediated mechanism. Nevertheless, the results of this study
support a cytoplasmic action for PR, and delineation of the mechanisms and the relative contributions of nuclear and cytoplasmic PR action in regulation of the cytoskeleton remains the focus of ongoing studies.

Altered regulation of the cytoskeleton upon PRA overexpression is consistent with the changed PRA expression demonstrated in cancers. Progression to malignancy in breast and endometrium is accompanied by disrupted PRA and PRB expression, and a significant proportion of carcinomas express a predominance of one isoform, usually PRA (Graham et al. 1995b, Arnett-Mansfield et al. 2001, Mote et al. 2002). This is associated with poor clinical outcomes: in endometrial cancer, PRA predominance is significantly more common in tumors of higher grade, indicating an association between PR isoform predominance and poor prognosis (Arnett-Mansfield et al. 2001). This study suggests that PRA predominance may confer progesterin responsiveness on specific components of the cytoskeleton and shows for the first time that the PR isoforms have distinct effects on the cytoskeleton by regulating multiple signaling pathways, some of which may be associated with changes in the cytoskeleton known to be associated with cancer development and progression.

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