Clathrin mediates integrin endocytosis for focal adhesion disassembly in migrating cells

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Focal adhesion disassembly is regulated by microtubules (MTs) through an unknown mechanism that involves dynamin. To test whether endocytosis may be involved, we interfered with the function of clathrin or its adaptors autosomal recessive hypercholesteremia (ARH) and Dab2 (Disabled-2) and found that both treatments prevented MT-induced focal adhesion disassembly. Surface labeling experiments showed that integrin was endocytosed in an extracellular matrix-, clathrin-, and ARH- and Dab2-dependent manner before entering Rab5 endosomes. Clathrin colocalized with a subset of focal adhesions in an ARH- and Dab2-dependent fashion. Direct imaging showed that clathrin rapidly accumulated on focal adhesions during MT-stimulated disassembly and departed from focal adhesions with integrin upon their disassembly. In migrating cells, depletion of clathrin or Dab2 and ARH inhibited focal adhesion disassembly and decreased the rate of migration. These results show that focal adhesion disassembly occurs through a targeted mechanism involving MTs, clathrin, and specific clathrin adaptors and that direct endocytosis of integrins from focal adhesions mediates their disassembly in migrating cells.

Introduction

Directional cell migration is a fundamental process required for embryonic development, inflammation, wound healing, cancer metastasis, and atherosclerosis (Lauffenburger and Horwitz, 1996; Ridley et al., 2003). A key aspect of directional migration of well-adherent cells is the establishment of transient attachments to the ECM through integrin clusters that form plaques known as focal adhesions. Focal adhesions establish a connection between the ECM and the actin cytoskeleton and serve as points of traction for the cell. The contraction of focal adhesion–associated actin stress fibers is thought to propel the cell body forward.

As the cell migrates, integrin clustering induces the formation of small focal adhesions (also referred to as focal contacts) at the front of the cell. Some of these nascent focal adhesions mature into larger focal adhesions, whereas others are rapidly turned over. Whether nascent focal adhesions disassemble or become mature focal adhesions depends on Rho-regulated myosin contractility (Rottner et al., 1999; Webb et al., 2004; Gupton and Waterman-Storer, 2006). Mature focal adhesions are selectively disassembled in the cell body so that few remain in the tail (Abercrombie, 1980; Smilenov et al., 1999). The disassembly of focal adhesions is important to allow for tail retraction, and integrin detachment from the ECM is rate limiting for cell migration in several cases (Hendey et al., 1992; Palecek et al., 1997).

In contrast to well-established mechanisms for focal adhesion formation (for reviews see Sastry and Burridge, 2000; Webb et al., 2002), the mechanisms for focal adhesion disassembly are not well understood. Focal adhesions in the tail of the cell may be disassembled or left on the substratum in processes that are regulated by calpain (Palecek et al., 1998) and Rho (Worthylake et al., 2001). Microtubules (MTs) also contribute to focal adhesion disassembly by delivering a relaxing factor whose nature is unknown (Kaverina et al., 1999). In none of these cases is it clear how focal adhesion disassembly is regulated.
Figure 1. **Clathrin is required for MT-induced focal adhesion disassembly.** (a) Western blot of clathrin heavy chain (HC) siRNA– or GAPDH siRNA–treated NIH3T3 fibroblasts showing levels of clathrin heavy chain, FAK, pY397 FAK, and GAPDH. (b) Clathrin heavy chain immunofluorescence and transferrin.
spatially regulated to target some focal adhesions for disassembly while others remain intact.

The fate of integrins after focal adhesion disassembly is also unknown. Experiments have suggested that a proportion of integrins from the tail are left behind on the substratum (Palecek et al., 1996). Other studies have suggested that integrins travel through vesicular intermediates and endomembrane compartments (Lawson and Maxfield, 1995; Palecek et al., 1996; Pierini et al., 2000). In these experiments, integrin trafficking was correlated with cell migration, but the relationship between focal adhesion disassembly and the fate of the integrin was not clearly established. Nonetheless, a prevailing idea is that the formation and disassembly of focal adhesions during cell migration are coupled to the recycling of integrins through endocytic processes. This idea is supported by evidence that general integrin recycling can contribute to cell migration (Caswell and Norman, 2006; Pellinen and Ivaska, 2006; Nishimura and Kaibuchi, 2007) and that integrins are endocytosed into Rab-labeled endocytic compartments during growth factor stimulation of cells (Roberts et al., 2001; Pellinen et al., 2006).

Focal adhesion disassembly occurs in a common cytoplasm along with focal adhesion formation, and there are few systems in which disassembly can be studied independently of assembly. We developed an assay that kinetically separates focal adhesion disassembly from assembly based on our finding that MT regrowth after nocodazole washout induced synchronous disassembly of focal adhesions. MT-induced focal adhesion disassembly was dependent on FAK and dynamin but was independent of active Rho and Rho-regulated contractility (Ezratty et al., 2005). MTs and dynamin are also involved in the turnover of podosomes, adhesive structures in macrophages, and other hematopoietic cells that are related to focal adhesions (Bruzzone et al., 2005; Kopp et al., 2006). Dynamin’s canonical role in the scission of vesicles during endocytosis raises the possibility that endocytosis may be involved in focal adhesion and podosome disassembly (Burridge, 2005).

In this study, we examine whether endocytosis of integrins regulates focal adhesion disassembly by exploring the role of clathrin in MT-induced focal adhesion disassembly. We find that clathrin is involved in focal adhesion disassembly and that clathrin accumulates at focal adhesions and departs with integrin during focal adhesion disassembly. We also show that the clathrin adaptors, Dab2 (Disabled-2) and autosomal recessive hypercholesteremia (ARH) target clathrin to focal adhesions and participate in the disassembly process. Our study supports a new mechanism for focal adhesion disassembly involving MT-stimulated endocytosis of integrins at focal adhesions.

## Results

### Clathrin is required for MT-induced focal adhesion disassembly

We first tested general inhibitors of clathrin-dependent endocytosis. Potassium depletion, which inhibits clathrin-mediated endocytosis, inhibited focal adhesion disassembly (unpublished data). To test specifically whether clathrin is required for focal adhesion disassembly, we reduced clathrin heavy chain expression using siRNA. Clathrin protein was diminished by >80% in cells treated with clathrin siRNA without affecting the levels of focal adhesion components FAK or vinculin (Fig. 1 a and not depicted). As previously reported (Moskowitz et al., 2005), transferrin uptake was inhibited in clathrin-depleted cells (Fig. 1 b). MT-induced focal adhesion disassembly was inhibited by clathrin depletion compared with untreated cells or control (glyceraldehyde 3-phosphate dehydrogenase [GAPDH]) depletion (Fig. 1 c and Fig. S1 a). Consistent with the inhibition of focal adhesion disassembly, cells depleted of clathrin still maintained stress fibers after MT regrowth, whereas cells depleted of GAPDH completely disassembled stress fibers (Fig. 1 c). Two clathrin siRNAs reduced clathrin levels and inhibited MT-induced focal adhesion disassembly, showing that there were no off-target effects (Fig. 1 d and Fig. S1 c). Quantification of focal adhesion disassembly using either a cell by cell assay (cells with <10 focal adhesions were scored as having disassembled focal adhesions) or quantitative measurements of focal adhesion number showed that clathrin depletion inhibited focal adhesion disassembly by 60–80% (Fig. 1 d and Fig. S1 b). Focal adhesion disassembly measured biochemically by monitoring levels of pY397 FAK, a FAK phosphorylation site which depends on focal adhesion (Ezratty et al., 2005; Yeo et al., 2006), showed that clathrin-depleted cells maintained pY397 FAK after MT regrowth compared with untreated cells (Ezratty et al., 2005) or control-depleted cells (Fig. 1 e). Depletion of clathrin heavy chain did not visually disrupt MTs, the rate of MT regrowth (not depicted), nor the proper targeting of MTs to focal adhesions, as assessed visually or with a quantitative assay (Fig. 1 f).
Long-term depletion of clathrin by siRNA could inhibit focal adhesion disassembly indirectly by altering protein-trafficking pathways. To test this possibility, we acutely disrupted clathrin function by expressing a cross-linkable FK506-binding protein 12 (FKBP)–clathrin light chain (LC) fusion protein that oligomerizes upon addition of the dimeric rapamycin derivative AP20187 (Clackson et al., 1998). This approach rapidly and specifically inhibits clathrin by disrupting assembly and disassembly of clathrin lattices and inhibits transferrin endocytosis (Moskovitz et al., 2003). Focal adhesion disassembly and transferrin uptake were inhibited in an AP20187-dependent fashion in cells expressing FKBP–clathrin LC (Fig. 1 g). Almost 80% of cells transfected with FKBP–clathrin LC and treated with AP20187 were inhibited in focal adhesion disassembly (Fig. 1 g). Combined with the clathrin depletion experiments, these data indicate that clathrin is required for most focal adhesion disassembly induced by MTs.

α5β1 integrin is internalized during MT-induced focal adhesion disassembly

The clathrin and dynamin (Ezratty et al., 2005) dependence of focal adhesion disassembly suggests that integrins are endocytosed during the process. To examine this possibility, we followed cell surface α5β1 integrin, the major ECM receptor engaged at focal adhesions formed on fibronectin in NIH3T3 fibroblasts. We labeled surface α5β1 integrin using fluorescent α5 integrin antibody and quantified surface α5 integrin levels by flow cytometry (see Materials and methods). Surface α5 integrin decreased almost 50% during MT regrowth after nocodazole washout, returning to almost the same level observed in cells before nocodazole treatment (Fig. 2 a and Fig. S2 a). The loss of surface integrin correlated well with the time course of focal adhesion disassembly. Surface levels of the collagen receptor subunit α1 integrin, which is not clustered in focal adhesions in cells adhering to fibronectin, remained unchanged during MT regrowth, showing that only integrin engaged by the ECM was affected by MT regrowth (Fig. 2 b and Fig. S2 a). Because total α5 integrin did not change during MT regrowth (Fig. S2 f), we conclude that ECM-bound α5 integrin is specifically endocytosed during focal adhesion disassembly. Consistent with this idea, clathrin depletion inhibited the loss of surface α5 integrin during MT regrowth relative to control-depleted cells (Fig. 2 c and Fig. S2, d and e). These results imply that only focal adhesion–associated integrin is internalized during MT-induced disassembly.

Bulk integrin endocytosed during growth factor stimulation transits through Rab5- and Rab11-positive compartments before being recycled (Roberts et al., 2001; Pellinen et al., 2006). To test whether internalized integrin transits through Rab5 endosomes during MT-induced focal adhesion disassembly, cells were transfected with constitutively active Rab5Q79L to enlarge early endosomes (Stenmark et al., 1994) and then focal adhesion disassembly was stimulated by MT regrowth. Expression of Rab5Q79L did not affect MT-stimulated focal adhesion disassembly (unpublished data). Before MT regrowth, only a small subset of the enlarged Rab5Q79L endosomes contained integrin (Fig. 2 d). However, after MT regrowth and focal adhesion disassembly, many of the Rab5Q79L endosomes contained β1 integrin (Fig. 2 d). The extent of colocalization of β1 integrin with Rab5-positive endosomes increased more than twofold after focal adhesion disassembly (Fig. 2 e). We also observed that after MT-induced focal adhesion disassembly, β1 integrin accumulated in a perinuclear compartment that localized with internalized transferrin and partially with Rab11, suggesting that the integrin internalized during MT regrowth traffics through an endocytic recycling compartment (Fig. 2 f). These data further support the idea that integrin travels through endocytic compartments after focal adhesion disassembly.

Clathrin accumulates at focal adhesions during MT-induced focal adhesion disassembly

To explore how clathrin might participate in focal adhesion disassembly, we examined the localization and dynamics of clathrin before and during MT-induced focal adhesion disassembly. By epifluorescence microscopy with antibodies against clathrin heavy chain, clathrin was observed in a perinuclear region corresponding to the Golgi as well as throughout the cytoplasm and the plasma membrane, as reported previously (Fig. 1 b). To visualize plasma membrane–associated clathrin, we used total internal reflection fluorescence (TIRF) microscopy, which only excites the ventral ~100 nm of the cell. Localization of endogenous clathrin heavy chain by TIRF microscopy revealed typical clathrin puncta on the ventral surface of the cell with some puncta clustered and colocalized with focal adhesions (Fig. 3 a). Clathrin puncta decorated 64% of focal adhesions (n = 250) and covered 30–35% of the total focal adhesion area (Fig. S3) but only 4.5% of the total ventral membrane area (not depicted). Interestingly, clathrin localization at focal adhesions increased slowly during the several hours of nocodazole treatment we used to completely break down MTs (Fig. S3).

The localization data showed that clathrin is present on a subset of focal adhesions, so we next explored clathrin dynamics in living cells to address two questions: (1) do focal adhesions that exhibit clathrin colocalization disassemble during MT regrowth, and (2) do focal adhesions that lack clathrin colocalization before MT regrowth acquire clathrin during their disassembly? For these experiments, we used a cell line stably expressing GFP–clathrin LC that had also been transiently transfected with RFP-FAK. A typical example of a focal adhesion that had accumulated clathrin before MT regrowth is shown in Fig. 3 b. The focal adhesion disassembled during MT regrowth, as judged by loss of FAK fluorescence over a 15-min interval, which is consistent with disassembly intervals from previous studies (Kaverina et al., 1999; Ezratty et al., 2005). In almost every case (22/24), focal adhesions that had accumulated clathrin over at least 20% of their area before MT regrowth disassembled when MTs were regrown. Clathrin puncta on these focal adhesions dynamically rearranged as the focal adhesion disassembled (Fig. 3 b). Despite the relatively high clathrin colocalization on these focal adhesions before MT regrowth, quantification showed that overall clathrin levels increased 2.3-fold on average during disassembly (Fig. 3, c and f).

To address the second question, we imaged focal adhesions that had low levels of clathrin colocalization before
MT regrowth. Virtually all of these focal adhesions showed a dramatic accumulation (17.4-fold; \( n = 35 \)) of clathrin puncta during the early periods of MT regrowth (Fig. 3 d and f). For example, the large focal adhesion shown in Fig. 3 d exhibited a 16.6-fold increase in clathrin colocalization during MT regrowth and a peak of clathrin fluorescence during its disassembly (Fig. 3 e). In contrast, clathrin did not accumulate at focal adhesions imaged in cells maintained in nocodazole (in which focal
to observe integrin loss from the densely labeled focal adhesions at early stages of disassembly, we focused on focal adhesions that were disassembling after intermediate times of MT regrowth and therefore contained patches of integrin (Fig. 4 a).

To catch individual clathrin events, we acquired images at a higher rate (every 20 s) than in our earlier videos of clathrin targeting to focal adhesions (Fig. 3). In a typical video acquired at this higher frame rate, a clathrin puncta can be seen to be recruited to a patch of $\beta_1$ integrin in a disassembling focal adhesion, colocalize with $\beta_1$ integrin, and then disappear simultaneously with the $\beta_1$ integrin (Fig. 4 b). The integrin appeared to concentrate in the clathrin puncta during its residence at the focal adhesion (frames 11:00–11:40). Fig. 4 c shows another example of codisappearance of $\beta_1$ integrin and clathrin puncta from the evanescent field during focal adhesion disassembly.

Of the clathrin puncta that were colocalized with $\beta_1$ integrin in disassembling focal adhesions, 57% were dynamic and disappeared from the evanescent field during the 10 min of imaging (Fig. 4 d). In more than two thirds of the cases, the departure of clathrin puncta was coincident with the loss of $\beta_1$ integrin (adhesions are not disassembling), showing that the rapid accumulation of clathrin was MT dependent (Fig. 3, g and h). These experiments show that clathrin accumulates at focal adhesions during their disassembly and reveal a previously unknown MT-dependent targeting of clathrin to focal adhesions.

**Clathrin endocytoses integrin during focal adhesion disassembly**

To determine whether clathrin mediated endocytosis of integrin during focal adhesion disassembly, we imaged clathrin dynamics in cells stably expressing a GFP–$\beta_1$ integrin chimera. We previously used this integrin chimera in a study of focal adhesion disassembly (Ezratty et al., 2005), and other experiments have shown that it is a faithful reporter for integrins at focal adhesions (Smilenov et al., 1999). Previous studies have used TIRF microscopy to identify the internalization of single clathrin-coated structures and their cargoes from the plasma membrane by virtue of their simultaneous disappearance from the evanescent field (Merrifield et al., 2002; Ehrlich et al., 2004; Puthenveedu and von Zastrow, 2006). Because it was difficult to observe integrin loss from the densely labeled focal adhesions at early stages of disassembly, we focused on focal adhesions that were disassembling after intermediate times of MT regrowth and therefore contained patches of integrin (Fig. 4 a).

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We next determined whether the alternate clathrin adaptors numb, Dab2, and ARH were localized at focal adhesions. These proteins comprise the set of known clathrin adaptors that contain a phosphotyrosine-binding domain, which interacts with the nonphosphorylated NPXY endocytic motifs in the cytoplasmic domain of certain receptors, including β1 integrin subunits (Calderwood et al., 2003; Uhlik et al., 2005). A screen of these adaptors revealed that Dab2 puncta accumulated in focal adhesions, whereas numb, ARH, and AP-2 were largely excluded from focal adhesions (Fig. 5 a). Quantification showed that Dab2 colocalized with 35% of the total focal adhesion area in nocodazole-treated cells, whereas numb, ARH, and AP-2 co-localized with <6% of the total focal adhesion area (Fig. 5 b). Virtually all of the clathrin puncta at focal adhesions colocalized with Dab2 (Fig. 5 c), and 68% of focal adhesions (n = 100) showed colocalization of both clathrin and Dab2. We occasionally detected Dab2 puncta at focal adhesions that did not colocalize with clathrin (unpublished data). These results suggest that Dab2 may be the primary adaptor protein that targets clathrin to the focal adhesion. This idea would be consistent with evidence that Dab2 binds the cytoplasmic tail of integrins in vitro and coimmunoprecipitates with β1 integrin in vivo (Calderwood et al., 2003; Prunier and Howe, 2005).

Figure 4. Direct imaging of clathrin and β1 integrin during focal adhesion disassembly. (a) TIRF imaging of GFP-β1 integrin during MT-induced focal adhesion disassembly. Time is in minutes:seconds after nocodazole washout. The boxed region is shown magnified at various time points in the panels on the right. [b and c] Montages from TIRF imaging of DsRed-clathrin and GFP-β1 integrin during focal adhesion disassembly. Time is in minutes:seconds. The boxed areas are depicted in the panels on the right. In the merged panels, arrowheads point to clathrin/β1 integrin puncta that colocalize and then simultaneously disappear. (d) Quantification of the behavior of β1 integrin-associated clathrin puncta. The histogram represents data from 30 focal adhesions in four different videos, in which 50 individual clathrin puncta were analyzed during focal adhesion (FA) disassembly. Error bars are SEM. Bars: (a) 10 µm; (b and c) 2 µm.
Dab2 and ARH are involved in clathrin targeting to focal adhesions and focal adhesion disassembly

We used siRNA knockdown to test whether Dab2, ARH, or numb was required for MT-induced focal adhesion disassembly. Two different specific siRNA duplexes that reduced proteins levels at least 80% were used for each adaptor molecule (Fig. 5 e and not depicted). Individual depletion of Dab2 or ARH partially inhibited focal adhesion disassembly, whereas depletion of numb had almost no effect compared with the GAPDH control (Fig. 5 d). We also stably depleted Dab2 in cell lines by expressing a short hairpin directed against Dab2 (shDab2). In two cell lines in which the levels of Dab2 were reduced by >80% compared with controls (Fig. S5 a), focal adhesion disassembly was inhibited only by ~20% (Fig. 5 d). ARH is a distant relative of Dab2, and both proteins have been shown to play a functionally redundant role in low density lipoprotein receptor internalization (Keyel et al., 2006). Given this potential functional redundancy, we used siRNA to deplete ARH in the shDab2 cell lines. Focal adhesion disassembly was inhibited in >40% of the cells depleted of both Dab2 and ARH (Fig. 5, d and f). In contrast, focal adhesion disassembly was inhibited in ~20% of cells depleted of either adaptor protein alone. Inhibition of focal adhesion disassembly by simultaneous ARH and Dab2 knockdown but not by Dab2 knockdown alone was also evident with the biochemical marker pY397 FAK (Fig. 5 e). Simultaneous depletion of ARH and Dab2 also inhibited the loss of cell surface α5 integrin during MT regrowth (Fig. 5 g). These data suggest that Dab2 and ARH function additively during MT-induced focal adhesion disassembly and integrin endocytosis.

To test whether these adaptor proteins were involved in recruiting clathrin to the focal adhesion, we localized clathrin in cells depleted of these proteins. Individual knockdown of either adaptor protein alone did not alter the localization of clathrin to focal adhesions (Fig. S5 b and not depicted). Strikingly, ARH redistributed to focal adhesions upon depletion of Dab2, which may explain why cells depleted of Dab2 have persistent focal adhesion–associated clathrin (Fig. S5 c). In cells simultaneously depleted of Dab2 and ARH, clathrin localization to focal adhesions was reduced by >50% (Fig. 5, h and i). Clathrin levels remained unchanged by Dab2 and ARH depletion (Fig. 5 e). Collectively, these results indicate that ARH and Dab2 contribute to the targeting of clathrin to focal adhesions and focal adhesion disassembly.

Clathrin and its adaptors accumulate at a subset of focal adhesions in migrating cells

Next, we examined the localization of clathrin and its adaptors during cell migration, during which focal adhesion disassembly occurs in a physiological context. In unperturbed cells migrating into an in vitro wound, clathrin puncta were observed at a subset of focal adhesions (Fig. 6 a). The most prominent co-localization of clathrin was with large focal adhesions in the midregion of the cell; fewer clathrin puncta were found in focal adhesions close to the leading edge. We quantified the regional co-localization of clathrin with focal adhesions by dividing the cell into four regions evenly spaced along the leading edge–tail axis (Fig. 6 b). Clathrin colocalized with 30% of the focal adhesion area in zone 2 but only 8–12% of the focal adhesion area in other regions of migrating cells (Fig. 6 c). In contrast, only 2% of the total ventral plasma membrane area contained clathrin puncta, indicating that clathrin is specifically concentrated at certain focal adhesions.

Quantification of the co-localization of the adaptors at focal adhesions showed that Dab2, and to a lesser extent ARH, colocalized with a high percentage of focal adhesions in region 2 of migrating cells (Fig. 6, d and e). Interestingly, both Dab2 and ARH also showed relatively high levels of colocalization with focal adhesions in regions 1 and 3, in which clathrin colocalization was relatively low. The accumulation of Dab2 and ARH in focal adhesions was specific as numb did not show appreciable accumulation in any region of the cell (Fig. 6 e). These results demonstrate that clathrin and specific clathrin adaptors are localized in specific focal adhesions in migrating fibroblasts. Combined with an earlier study, which showed that disassembly of large focal adhesions occurred predominantly in the midregion of migrating cells (Smilenov et al., 1999), these results are consistent with the idea that these molecules are involved in focal adhesion disassembly during cell migration.
Clathrin and its adaptors contribute to focal adhesion disassembly in migrating cells

To determine whether clathrin and its adaptors are important for focal adhesion disassembly in a physiological context, we depleted clathrin, Dab2, and ARH using siRNA and tested the ability of these cells to migrate into a wounded monolayer. Clathrin and Dab2 have previously been implicated in cell migration (Hocevar et al., 2005; Nishimura and Kaibuchi, 2007; Chao and Kunz, 2009), and we confirmed that cells depleted of these proteins were impaired in directed migration into an in vitro wound (Fig. 7, a and b). Depletion of ARH, which has not previously been studied in cell migration, also inhibited direct migration (Fig. 7, c and d). This phenotype is characteristic of defective focal adhesion disassembly and had not been previously reported in other studies investigating the role of Dab2 and clathrin during cell migration. To determine whether depletion of clathrin was related to defects in focal adhesion disassembly, we analyzed focal adhesion disassembly in cells stably expressing GFP-FAK (Partridge and Marcantonio, 2006) and depleted of clathrin or ARH and Dab2. We found that overall focal adhesion disassembly was reduced by 60–70% in cells depleted of clathrin or ARH and Dab2 (Fig. 7, e and f). These results parallel our finding that depletion of both adaptor proteins was necessary to significantly inhibit MT-induced focal adhesion disassembly. Migrating cells depleted of clathrin or both clathrin adaptors also exhibited elongated tails (Fig. 7, c and d). This phenotype is characteristic of defective focal adhesion disassembly and had not been previously reported in other studies investigating the role of Dab2 and clathrin during cell migration. To determine whether inhibition of cell migration was related to defects in focal adhesion disassembly, we analyzed focal adhesion disassembly in migrating cells stably expressing GFP-FAK (Partridge and Marcantonio, 2006) and depleted of clathrin or ARH and Dab2. We found that overall focal adhesion disassembly was reduced by 60–70% in cells depleted of clathrin or ARH and Dab2 (Fig. 7, e and f). These results parallel our finding that depletion of both adaptor proteins was necessary to significantly inhibit MT-induced focal adhesion disassembly. Migrating cells depleted of clathrin or both clathrin adaptors also exhibited elongated tails (Fig. 7, c and d). This phenotype is characteristic of defective focal adhesion disassembly and had not been previously reported in other studies investigating the role of Dab2 and clathrin during cell migration. 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Figure 7. **Clathrin and clathrin adaptors are required for focal adhesion disassembly and polarized cell migration.** (a) Phase images from videos of NIH3T3 cells treated with the indicated siRNA and allowed to migrate into a wound. The same wound edge is shown at 0 and 6 h of migration. (b) Quantification of wound migration velocity of cells depleted of clathrin or clathrin adaptors. The histogram represents two to three independent experiments in which 20 wounds were analyzed per condition. *, $P \leq 0.0001$ for each condition compared with GAPDH by Student’s t test. (c) Immunofluorescence of NIH3T3 cells depleted of either clathrin or Dab2 and ARH and immunostained for vinculin. Cells were allowed to migrate into a wound for 9 h before processing for immunofluorescence. Arrows indicate examples of elongated tails. (d) Quantification of tail length in migrating cells depleted of clathrin or clathrin adaptors. The histogram represents two independent experiments in which at least 20 cells were analyzed per condition. *, $P \leq 0.005$ for each condition compared with mock by Student’s t test. (e) Fluorescence images from videos of migrating NIH3T3 cells stably expressing GFP-FAK and depleted of clathrin, Dab2, and ARH or mock treated (0-, 20-, or 30-min time points are shown). Small arrows point to focal adhesions that disassemble during 30 min of cell migration. Arrowheads point to focal adhesions that fail to disassemble during 30 min of cell migration. Large arrows indicate the direction of migration. (f) Quantitative analysis of focal adhesion (FA) disassembly in cells depleted of clathrin or ARH and Dab2. The histogram shows the percentage of focal adhesions that disassemble per cell during a 30-min interval. For each of the conditions analyzed, focal adhesion disassembly was quantified for at least 10 different cells (30–50 adhesions per cell) from five to eight separate videos. (b, d, and f) Error bars are SEM. Bars, 15 µm.
results are consistent with the idea that clathrin and its adaptors Dab2 and ARH play an important role in focal adhesion disassembly during directed cell migration.

Discussion

We have shown that MT-induced focal adhesion disassembly occurs via the direct endocytosis of integrin at focal adhesions mediated by clathrin. It has been suggested that general loss of contractility is the primary mechanism by which focal adhesions disassemble in response to MT targeting (Kaverina et al., 2002). However, our data indicate that MT-induced focal adhesion disassembly is mediated by a hierarchical assembly of endocytic components at the focal adhesion, which then induces disassembly by mediating endocytosis of integrins. TIRF imaging data support the direct endocytosis of integrin by clathrin because we observed the simultaneous disappearance of clathrin and integrin during disassembly, although we cannot rule out that some integrin is disassembled through other processes. Our current results, coupled with earlier results showing the dynamin dependence and Rho independence of focal adhesion disassembly (Ezratty et al., 2005), strengthen the idea that focal adhesion disassembly is mechanistically distinct from focal adhesion formation.

The factors involved in integrin endocytosis have been unexplored until recently. Our study points to clathrin and its adaptors Dab2 and ARH as key participants in integrin endocytosis. The weaker inhibition of focal adhesion disassembly and integrin endocytosis in Dab2/ARH compared with clathrin-depleted cells suggests that there may be additional adaptors involved in clathrin-mediated disassembly of focal adhesions. In fact, while our paper was under review, another paper reported that Dab2 and AP-2 were involved in focal adhesion disassembly and integrin endocytosis in human fibrosarcoma cells (Chao and Kunz, 2009). We did not observe AP-2 localization to focal adhesions or inhibition of disassembly with dominant-negative AP-2 constructs, suggesting that AP-2 is not generally involved in focal adhesion disassembly. Studies on HeLa cells have suggested that numb and Dab2 participate in integrin endocytosis, although in these studies, the internalized integrin was not likely to be engaged and in focal adhesions (Nishimura and Kaibuchi, 2007; Teckchandani et al., 2009). There is also evidence that non-ECM–engaged integrin can be endocytosed by a clathrin-independent, caveolin-dependent pathway (Fabbri et al., 2005; Echarri and Del Pozo, 2006). Together, these studies point to a degree of complexity in integrin endocytosis in which it is possible that different adaptors are used for different integrins or for the same integrin but in different states of engagement with the ECM.

The accumulation of clathrin and its adaptors at focal adhesions provides a way to temporally and spatially regulate focal adhesion disassembly during cell migration and suggests that focal adhesions are more heterogeneous structures than had previously been appreciated. We found that both clathrin and its adaptors Dab2 and ARH were required for focal adhesion disassembly and localized to focal adhesions in specific regions of migrating fibroblasts. Interestingly, Dab2 and ARH were observed at more focal adhesions than clathrin, suggesting that Dab2 and ARH may precede clathrin at focal adhesions or that clathrin may not persist for long before focal adhesion disassembly commences. Additional imaging studies will be needed to ascertain the dynamic relationship between Dab2, ARH, and clathrin at disassembling focal adhesions.

How might clathrin and its adaptors target focal adhesions for their disassembly? Previous studies have shown that more mature focal adhesions accumulate late focal adhesion proteins such as α-actinin (Laukaitis et al., 2001) and have higher levels of tyrosine phosphorylated focal adhesion components (Zamir et al., 1999). The accumulation of clathrin or its adaptors at focal adhesions may be caused by these or other maturation processes at focal adhesions. MTs may also contribute to the accumulation by rapidly directing clathrin and/or its adaptors to focal adhesions. Once clathrin and its adaptors accumulate at focal adhesions, they may stimulate endocytosis not only by recruiting clathrin but by competing with existing interactions between focal adhesion components and integrin. For example, Dab2 may compete with talin for NPXY-binding sites, and this could decrease integrin–ECM engagement by reversing integrin activation and/or releasing integrin from its primary connection to the actin cytoskeleton.

It is well established that MTs are not required for endocytosis per se, and it has been suggested that MTs likely deliver a relaxing factor to focal adhesions that may facilitate their disassembly (Kaverina et al., 2002). Both clathrin and Dab2 accumulated slowly (over hours) at focal adhesions in the absence of MTs, indicating that MTs are not absolutely required for their targeting to focal adhesions. However, a much more rapid accumulation (in minutes) of clathrin at focal adhesions occurred during MT-induced focal adhesion disassembly, suggesting that MTs may facilitate the targeting of clathrin to focal adhesions. The rapid accumulation of clathrin may reflect delivery of clathrin by MTs as clathrin puncta have been seen traveling along MTs near the plasma membrane (Rappoport et al., 2003). The possibility that Dab2 or ARH move along MTs has not yet been explored. Alternatively, MTs may regulate the accumulation of clathrin or its adaptors at focal adhesions indirectly by regulating binding sites for these proteins at the focal adhesion. It will be interesting to determine the mechanism by which MTs enhance clathrin and perhaps other endocytic protein targeting to focal adhesions during disassembly.

Materials and methods

Chemicals and cell culture

All chemicals were purchased from Sigma-Aldrich unless otherwise noted. NIH3T3 fibroblasts were cultured in DME and calf serum as previously described (Ezratty et al., 2005). HA1 cells (NIH3T3 fibroblasts engineered to express human α1 integrin) were cultured in DME, calf serum, and 300 µg/ml G418 as previously described (Briesewitz et al., 1993). For serum starvation, cells on acid-washed coverslips were grown to confluence (~2 d) and transferred to serum-free medium (DME and 10 mM Hepes, pH 7.4) for 48 h as previously described (Ezratty et al., 2003). shDab2 pools and clonal cell lines were derived as previously described (Prunier and Howe, 2005) except NIH3T3 fibroblasts were cotransfected with the shDab2 construct (provided by P. Howe, Lerner Research Institute, Cleveland, OH) and pcDNA3.1 Zeo and then selected using 500 µg/ml Zeocin (Invitrogen).

Focal adhesion disassembly and cell migration experiments

Focal adhesion disassembly and cell migration experiments were performed as previously described (Ezratty et al., 2005). In brief, NIH3T3...
cells grown on glass coverslips were treated with 10 µM nocodazole for 3–4 h to completely depolymerize MTs. The drug was washed out with serum-free medium, and MTs were allowed to repolymerize for different intervals. Cells were either fixed in −20°C methanol for 10 min and rehydrated in TBS or fixed in 4% paraformaldehyde in PBS for 10 min, followed by permeabilization with 0.5% Triton X-100 in PBS for 5 min before immuno-fluorescence staining. Depending on the experiment, cells were stained with phalloidin (mouse monoclonal; 1:100 dilution in PBS-H), DAPI or Alexa Fluor 488 to label nuclei, cytoskeletal actin (monoclonal; 1:1000; Invitrogen), α-actinin (1:100), vinculin (1:100; Sigma-Aldrich), dynamin (hdy-1; 1:100; Millipore), clathrin [TD.1 [1:50; BD] or X22 [1:100; Abcam]), AP2-1 [1:100; α-adaptin subunit; Abcam], Dab2/p96 [mouse monoclonal; 1:100; BD], ARH (rabbit polyclonal; 1:800; provided by M. Farquhar, University of California, San Diego, La Jolla, CA), nNOS (1:100; Santa Cruz Biotechnology, Inc.), b1 integrin [1:100; BD], tyrosinated tubulin [rat monoclonal YL1/2; 1:10 dilution of culture supernatant], or Cy5/shodamine-phalloidin (1:200 dilution; Cytoskeleton, Inc.). Secondary antibodies were purchased from Jackson ImmunoResearch Laboratories, Inc. and were absorbed to minimize cross-reaction with other species.

Clathrin inhibition and RabG79L experiments
NIH3T3 cells were transfected with FKBP–clathrin LC–GFP (provided by T. Ryan, Cornell University, New York, NY; Moskowitz et al., 2003) or RabG79L (provided by J. Lippincott-Schwartz, National Institutes of Health, Bethesda, MD; Stenmark et al., 1994) with Lipofectamine Plus (Invitrogen) according to the manufacturer’s instructions. After 24–48 h of transfection, cells were treated with 10 µM nocodazole for 4 h, and then focal adhesion disassembly was stimulated via MT regrowth. For clathrin inhibition, 100 nM AP20187 (ARIAD) was added to FKBP–clathrin LC–GFP-transfected cells for 30 min before nocodazole washout and during MT regrowth. For transferin internalization, cells were incubated at 37°C with 10 µg/ml Alexa Fluor 546-labeled transferrin (Invitrogen) for 10 min. Remaining plasma membrane-bound transferrin was removed by incubating cells for 2 min with 0.5 M NaCl and 0.2 M of Na acetate buffer, pH 4.5, followed by washing with the same solution before processing for immunofluorescence.

siRNA knockdown experiments
RNA duplexes [21 nucleotides] were purchased from Invitrogen or Shangh hai GenePharma and resuspended in RNAase-free water at a concentration of 20 µM. siRNAs were designed based on previously published sequences (Hoevar et al., 2005; Moskowitz et al., 2005) as follows: clathrin-1, 5′-AACAUUGCCUCAUGCUAGUGTGT-3′; clathrin-2, 5′-AAUUGGCAUCUGUAAACUGGGT-3′; Dab2-1, 5′-CGUCUACUCCACAGAGUAATT-3′; Arh-1, 5′-CGTGCTGTAGTGGAGAAGTT-3′; Arh-2, 5′-GTTGCGACAGAATCTGGACG-3′; and nNOS, 5′-CUAGUAAGUGUGAUGUGAC-3′. The mouse GAPDH siRNA sequence was 5′-CGUGAAUUGUGAAGUGAC-3′ and was used as a negative control in all siRNA experiments. For some experiments, Dab2 and ARH were knocked down simultaneously by treating with both siRNAs. NIH3T3 fibroblasts were transfected with siRNA duplexes using Lipofectamine RNAiMax [Invitrogen] according to the manufacturer’s instructions for reverse transfection. Optimal knockdown for all proteins was obtained by two transfections spaced 48 h apart. Knockdown efficiency and effects on focal adhesion disassembly and transferrin uptake were analyzed 24–48 h after the last transfection.

Western blotting
Cells were lysed in lysis buffer (1% NP-40, 50 mM Tris, pH 7.4, 150 mM NaCl, 2 mM MgCl₂, 1 mM EDTA, and protease and phosphatase inhibitors). 2× Laemmli sample buffer was used to load lysates. Proteins were separated by SDS-PAGE on 7.5 or 10% gels. Lysate protein concentration was determined by Coomassie blue or Amido Black staining. Gels were transferred to nitrocellulose, blocked 1 h at room temperature in cold 4% paraformaldehyde/PCN, and the distribution and mean fluorescence intensity of each sample was quantified using a FACScan (BD).

Cubing optimized for coumarin, fluorescein/GFP, rhodamine, and Cy5 fluorescence. Images were captured with a camera (Kodak KAF 1400-chip MicroMax; Princeton Instruments) using MetaMorph software (MDS Analytical Technologies). For TIRF microscopy, preparations were observed using a 60× NA 1.45 Apochromat objective on a microscope (TE2000-U; Nikon) equipped with a TIRF illuminator and fiber optic-coupled laser illumination. Illumination was performed with three separate laser lines (Ar [488 nm], 110 mW; and HaNa [543 and 568 nm] and HeNe; 40 mW). Imaging of fluorescein/GFP, Cy3/Alexa Fluor 546, and Cy5 fluorescence (Chroma Technology Corp.). Images were captured with a camera (OrcaIIR; Hamamatsu Photonics) using MetaMorph software. For live cell imaging of clathrin and FAK, stable NIH3T3 cell lines expressing GFP–clathrin LC (provided by C. Waterman-Storer, National Institutes of Health) were generated as previously described (Merrifield et al., 2002) and transiently transfected with RFP–FAK (Partridge and Marcatronico, 2006) for 24–48 h before TIRF imaging. Images of RFP–FAK and GFP–clathrin were acquired every 60 s. For live cell imaging of integrin and clathrin, stable NIH3T3 cell lines expressing low levels of a GFP–b1 integrin chimera (Smilenov et al., 1999) were transiently transfected with DiRed–clathrin LC (Merrifield et al., 2002) for 24–48 h. Images of GFP–b1 integrin and clathrin were acquired every 20–60 s. Imaging was performed at 37°C in recording media (Gomes and Gundersen, 2006) and started within 15 min of MT regrowth.

For imaging cell migration, monolayers of cells transfected with siRNAs were wounded and imaged in recording media supplemented with 2% calf serum. Multiple fields were imaged every 30 min for 9–12 h using an XY stage (Prior) and a 10× NA 0.46 Plan-Fluor objective (Nikon). Photobleach contrast images were collected with a camera (CoolSNAP HQ; Roper Industries) controlled by MetaMorph. For analysis of focal adhesion dynamics, monolayers of NIH3T3 cells stably expressing GFP–FAK (Partridge and Marcatronico, 2006) and transfected with siRNAs were imaged as described above, except that multiple fields were imaged every 10 min for 2–3 h.

Quantitative immunofluorescence and image processing
The MetaMorph measure colocalization function was used to quantify colocalization of focal adhesions and clathrin or adaptor proteins and colocalization of integrin and RabG79L endosomes. Each source image was equally thresholded to obtain just focal adhesion, clathrin, or adaptor protein staining before measuring the extent of colocalization. For colocalization experiments in migrating cells, the length of the wound edge cells was measured from tail to front, and the perimeter was determined by manually tracing an outline of the cell. This entire area was then divided into four equal regions along the front to back axis. A region tool was used to select a specific region of interest (labeled as quadrants 1–4) for the measurement of clathrin or adaptor/focal adhesion (detected with FAK-Y397 or vinculin) colocalization. For analysis of focal adhesion disassembly during cell migration, individual focal adhesions were used as a negative control thresholded and monitored for their disappearance or maintenance within a 30-min interval during cell migration. For plots of clathrin disassembly during focal adhesion disassembly, the region tool was used to outline a given focal adhesion, and its area was determined in sequential images during MT-induced focal adhesion disassembly. The measure colocalization function was then used to determine the area of overlap between the focal adhesion and clathrin. To measure the size or number of focal adhesions, fluorescent images of individual cells were thresholded, and the measurement function in MetaMorph was used to calculate the area and number of objects. To analyze the length of tails in migrating cells, a line was drawn from the end of the tail to the back of the nucleus, and the total length of the line was determined. For all images shown, the brightness and contrast were first optimized and appropriately scaled in MetaMorph before importing into Photoshop (Adobe) for the preparation of figures.

Cell surface labeling and flow cytometry
Cells were harvested by trypsinization at time points after MT regrowth and washed in cold PCN (1× PBS, 0.5% calf serum, and 0.1% NaNO₃). Approximately 4 × 10⁵ cells per sample were stained at 4°C without fixation. For the detection of α5 integrin in NIH3T3 and HA1 fibroblasts, PE-conjugated rat α-mouse CD49e monoclonal antibody (BD) was used at a dilution of 1:1,500. For the detection of α1 integrin in HA1 cells, mouse α-human TS2/7 monoclonal antibody (gift from T. Springer, Harvard University, Cambridge, MA) was used at a dilution of 1:200, followed by 1:500 dilution of FITC α-mouse IgG (Jackson ImmunoResearch Laboratories, Inc.). After washes to remove unbound antibody, samples were fixed in cold 4% paraformaldehyde/PCN, and the distribution and mean fluorescence intensity of each sample was quantified using a FACScan (BD).
Statistical analysis
Where indicated, data were analyzed using the Student’s t test.

Online supplemental material
Fig. S1 shows that clathrin knockdown prevents focal adhesion disassembly and transferrin internalization compared with mock-transfected cells. Fig. S2 shows the total surface s5 or s1 integrin levels before and after MT-mediated focal adhesion disassembly. Fig. S3 shows that clathrin accumulates at focal adhesions during nocodazole treatment. Fig. S4 shows that the expression of Eps15-295 inhibits transferrin uptake but not focal adhesion disassembly. Fig. S5 shows the localization of clathrin and clathrin adaptor proteins in cells depleted of Dab2. Online supplemental material is available at http://www.jcb.org/cgi/content/full/jcb.200904054/DC1.

E.E. Marcantonio would like to dedicate this paper to the memory of Dr. Tucker Collins.

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