Binding of ZO-1 to α5β1 integrins regulates the mechanical properties of α5β1–fibronectin links

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ABSTRACT Fundamental processes in cell adhesion, motility, and rigidity adaptation are regulated by integrin-mediated adhesion to the extracellular matrix (ECM). The link between the ECM component fibronectin (fn) and integrin α5β1 forms a complex with ZO-1 in cells at the edge of migrating monolayers, regulating cell migration. However, how this complex affects the α5β1-fn link is unknown. Here we show that the α5β1/ZO-1 complex decreases the resistance to force of α5β1–fn adhesions located at the edge of migrating cell monolayers while also increasing α5β1 recruitment. Consistently with a molecular clutch model of adhesion, this effect of ZO-1 leads to a decrease in the density and intensity of adhesions in cells at the edge of migrating monolayers. Taken together, our results unveil a new mode of integrin regulation through modification of the mechanical properties of integrin–ECM links, which may be harnessed by cells to control adhesion and migration.

INTRODUCTION

General processes in development, wound healing, or cancer are driven by cell adhesion and migration, which are determined by the interaction between cells and the extracellular matrix (ECM; An et al., 2009). This interaction is largely mediated by integrins, and specific ECM–integrin links such as those formed by the ECM protein fibronectin (fn) and integrin α5β1 are involved in crucial cellular processes in signaling and mechanotransduction (Katsumi et al., 2004; Elosegui-Artola et al., 2014). Integrin-mediated functions are regulated by a myriad of integrin-binding adaptor proteins (Calderwood, 2004; Roca-Cusachs et al., 2012), which can affect both their activation and biochemical signaling (Ghatak et al., 2013; Hytönen and Wehrle-Haller, 2015) and their mechanical properties (Giobanasi et al., 2013; Goldmann et al., 2013; Das et al., 2014). In turn, the affinity and mechanical properties of integrin–ECM links (and specifically of α5β1–fn links) regulate mechanotransduction and the ability of cells to both transmit forces to the substrate and transduce them into downstream biochemical signals (Elosegui-Artola et al., 2014, 2016). Thus regulation of integrin mechanics by adaptor protein interactions emerges as a potential way to tune mechanotransduction.

An adaptor protein described to bind to α5β1 is tight junction protein ZO-1, which is generally localized to cell–cell adhesions (Fanning, 1998; González-Mariscal et al., 2008) but binds to the α5 subunit of α5β1 (Taliana et al., 2005; Tuomi et al., 2009). The formation of this complex affects cell motility (Tuomi et al., 2009) and is crucial for cytokinesis (Hämälistö et al., 2013). The formation of the α5β1/ZO-1 complex is mediated by protein kinase Cε-dependent phosphorylation of ZO-1. Once phosphorylated, ZO-1 then translocates to cell lamellipodia, but only in subconfluent cells. This interaction is believed to stabilize and polarize cells because if disrupted, directional persistence and migration velocity are modified in different cell types (Tuomi et al., 2009; Bazellières et al., 2015). Of interest, alterations in α5β1 and ZO-1 are related to malignant phenotypes (Roman et al., 2010; Ni et al., 2013; Paul et al., 2015). Thus ZO-1 acts as a regulator of α5β1 integrins, and we hypothesized that its effect could be mediated by changes in mechanical properties of the α5β1–fn link.

RESULTS AND DISCUSSION

Disruption of the α5β1/ZO-1 complex affects cell motility

In this study, we used the human mammary epithelial cell line MCF10A, the migration of which has been previously studied (Bazellières et al., 2015; Vincent et al., 2015). To check that ZO-1 was
translocating to the lamellipodia of cells at the edge of monolayers, we seeded a monolayer of cells on 12-kPa gels coated with 10 µg/ml of fn and stained them for activated β1 and ZO-1. As previously described (Tuomi et al., 2009), ZO-1 localized to cell–cell contacts in confluent cells and to the lamellipodia in cells at monolayer edges (Figure 1, A and B), that is, cells with a free edge without cell–cell contacts. Accordingly, colocalization with α5β1 was significantly increased only at lamellipodia (Figure 1C).

To study the effects of the α5β1/ZO-1 complex, we impaired its formation by using a combination of small interfering RNA targeting ZO-1 (siZO1) and independently by transfecting a dominant-negative ZO-1 plasmid (S168A) that impairs binding of endogenous ZO-1 to α5 (Tuomi et al., 2009; Hämäläistö et al., 2013). As a control, we used cells transfected with a nontargeting siRNA (siCT). ZO-1 concentrations decreased to ~25% in siZO1 and were only slightly affected in S168A-transfected cells (Figure 1, D and E). Whereas it still localized to cell–cell junctions, ZO-1 had reduced expression in siZO1 cells both at the center and at the edge of monolayers (Figure 1A). Consistent with its reported inability to bind α5, S168A did not localize to lamellipodia or colocalize with α5β1 in any case (Figure 1, A–C). Confirming the dominant-negative effect of S168A, total ZO-1 in S168A cells was also unable to localize to lamellipodia, whereas it localized normally, as expected, to cell–cell junctions (Supplemental Figure S1).

Previous studies demonstrated a role of the α5β1/ZO-1 complex in cell migration in different cancer cell lines (Tuomi et al., 2009). To verify this in our system, we seeded cells on the same 12-kPa gels. We then tracked the cells for 8 h and assessed their directional persistence as the ratio between their effective traveled distance (radial distance from the starting point) and their total traveled distance (sum of the total path). As previously shown, siZO1 and S168A cells were significantly less directional than siCT (Figure 1, F–H).

Thus, in agreement of previous work (Tuomi et al., 2009), our data confirmed that ZO-1 localizes to the lamellipodia of monolayer edge cells, and depleting ZO-1 or preventing its association with α5 affects cell migration.

To investigate whether ZO-1 binding to α5β1 affected the α5β1–fn link, we used a previously described setup based on
sured adhesion strength was specific to fn and mediated by \( \alpha_5\beta_1 \) integrins. We note that despite certain sample-to-sample variability, we did not observe significant differences between conditions at monolayer centers, further suggesting that ZO-1 had a differential effect on monolayer edges.

Formation of the \( \alpha_5\beta_1/ZO-1 \) complex increases \( \alpha_5\beta_1–fn \) recruitment

The measured effects in adhesive strength could be mediated by changes in the resistance to force of integrin–fn links or by changes in integrin recruitment. To assess potential changes in integrin recruitment, we incubated nonfluorescent, FN7-10–coated silica beads with cells (Elosegui-Artola et al., 2014), fixed the cells, and stained them for ZO-1 and activated \( \beta_1 \) (Figure 3A and total \( \beta_1 \) (Supplemental Figure S2). All conditions showed recruitment of magnetic tweezers (Elosegui-Artola et al., 2014). We first coated superparamagnetic beads with FN7-10, a fn fragment (Coussen et al., 2002; Elosegui-Artola et al., 2014) that binds mechanically to cells primarily through \( \alpha_5\beta_1 \) (Roca-Cusachs et al., 2009). We allowed beads to attach to cell monolayers for 35 min and then pulled on them using the magnetic tweezers with a force of 0.5 nN until beads detached both at the subconfluent edge and at the center of monolayers within the same sample (Figure 2, A and B).

In siCT cells, bead detachment times significantly decreased at monolayer edges with respect to the monolayer center (Figure 2C). This difference was lost both in siZO1 cells and in cells transfected with S168A, showing that this differential regulation is mediated by ZO-1 and its ability to bind \( \alpha_5 \). As negative controls, beads detached immediately when coated with either biotinylated bovine serum albumin (bBSA) or FN7-10 in cells preincubated with an \( \alpha_5\beta_1 \) blocking antibody (A\( \alpha_5\beta_1 \)) (Figure 2C). This indicates that the measured adhesion strength was specific to fn and mediated by \( \alpha_5\beta_1 \) integrins. We note that despite certain sample-to-sample variability, we did not observe significant differences between conditions at monolayer centers, further suggesting that ZO-1 had a differential effect on monolayer edges.

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bead detachment under force is regulated by the mechanical properties of the links and not by their recruitment (or affinity), which would lead to the opposite result. Further confirming this direct link to the properties of integrin-fn links, neither bead detachment times nor integrin recruitment were affected by impairing cell contractility with blebbistatin (Supplemental Figure S3). Thus our results are consistent with ZO-1 affecting α5β1-fn links specifically at monolayer edges by increasing both their off-rates (i.e., reducing their lifetime under force) and their on-rates (thereby increasing recruitment).

**Adhesion formation is affected by the α5β1/ZO-1 complex**

According to our previously described molecular clutch model of adhesion formation (Elosegui-Artola et al., 2014), a concomitant increase in both on- and off-rates (likely induced by ZO-1) should decrease force loading in integrins, impairing mechanotransduction, subsequent force transmission, and formation of adhesions. Surprisingly, we did not observe changes in force transmission in the different conditions (Supplemental Figure S4), possibly because the potential decrease mediated by ZO-1 was compensated by increased levels of total and phosphorylated myosin in siCT cells compared with other conditions (Supplemental Figure S3). However, and consistent with model predictions, the number and intensity of paxillin-rich adhesions at monolayer edges decreased in siCT cells compared with siZO-1 cells and cells transfected with S168A (Figure 4, A and B). As expected, we did not observe differences in adhesion formation at the center of monolayers (Figure 4A). Of interest, even though their intensity and density were lower (Figure 4, B and C), adhesions in siCT cells were significantly longer than in other conditions (Figure 4D). This suggests that mechanical regulation of the α5β1-fn link by ZO-1 may readily explain adhesion formation but not later maturation processes, which may depend on other factors (Choi et al., 2008).

Taken together, our results reveal how tight junction protein ZO-1, an integrin regulator, affects the mechanics and dynamics of α5β1-fn adhesions, increasing their affinity and decreasing their ability to withstand forces. Although a detailed confirmation would require a complex in vitro analysis with isolated purified proteins, this effect suggests an increase in both binding and unbinding rates between fn and α5β1. This could explain the associated decrease in adhesion density and intensity caused by ZO-1 at monolayer edges. In turn, the faster binding dynamics caused by ZO-1 association may facilitate the migration of control versus ZO-1–depleted cells by facilitating adhesion remodeling. How ZO-1 association leads to the observed mechanical changes in

**Figure 4:** Disruption of the α5β1/ZO-1 complex leads to a higher number of smaller nascent adhesions. (A) Paxillin staining at the subconfluent edge and at the center of monolayers. (B) Quantification of adhesion intensity. (C) Quantification of adhesion density. (D) Quantification of adhesion length. Scale bar, 50 µm. Insets show the area marked with a red square. Images are representative from three experiments.
α5β1–fn links and the combination of mechanical and biochemical factors by which it affects behaviors such as cell migration remain to be elucidated. Further, those effects may depend on cell type and substrate coating and stiffness, which are affected, for instance, in cancer (Paszek et al., 2005; Plodinec et al., 2012). Whereas we found that ZO-1 depletion decreased myosin phosphorylation and did not affect cell–matrix force transmission, studies using other cell types, matrix coatings, and conditions found even (Fanning et al., 2012) or increased myosin phosphorylation (Tornavaca et al., 2015) and increased contractility (Bazellières et al., 2015; Choi et al., 2016). However, and in general terms, our results exemplify how adaptor proteins can regulate integrin function by affecting not only their activation or affinity for ECM ligands but also their mechanical properties under force.

Previous work showed how adaptor proteins such as talin, α-actinin, or vinculin mediate integrin activation and mechanotransduction, leading to increased adhesion strength and reinforcement (Mierke et al., 2008; Roca-Cusachs et al., 2009). Here we demonstrate an alternative and counterintuitive mechanism by which another adaptor protein (ZO-1) promotes activation but decreases mechanical resistance. Because such mechanical regulation is bound to affect downstream mechanosensing processes, this provides an interesting and novel way to regulate cell adhesion, mechanorespons, and function in general.

MATERIALS AND METHODS
MCF10A cell culture and transfection
MCF10A cells were grown as described previously (Bazellières et al., 2015) and tested negative for mycoplasma contamination. Cells were transfected using the Lipofectamine 3000 transfection kit (Invitrogen) following manufacturer’s instructions using either a pool of three siRNAs (Bazellières et al., 2015) or 5 ng of the plasmid (ZO-1-1685–A-FLAG; S168A), a kind gift from the laboratory of Johanna Ivaska (University of Turku, Finland; Tuomi et al., 2009). Five days after transfection, cells were trypsinized and used for experiments. S168A plasmid has a point mutation in serine 168 that impairs its binding to α5 and is tagged with the FLAG peptide for identification.

Magnetic tweezers and bead-recruitment experiments
Magnetic tweezers experiments were carried out as previously described (Roca-Cusachs et al., 2009; Elosegui-Artola et al., 2014; Bazellières et al., 2015). Briefly, carboxylated 3-µm magnetic beads (Invitrogen) were coated with a mixture of biotinylated pentameric FN7-10 (a four-domain segment of fibronectin responsible for cell–substrate interactions. Fn-coated beads were then deposited on the coverslip. Polyacrylamide gels and traction measurements
Polyacrylamide gels of 12 kPa were prepared as described by Kandov et al. (2007) and incubated with 10 µg/ml fn (Sigma-Aldrich) overnight at 4°C. Gels were then sterilized with ultraviolet light and washed once with phosphate-buffered saline 1× for immediate use. Traction forces were computed using Fourier transform traction microscopy with finite gel thickness (Trepata et al., 2009) as previously described (Serra-Picamal et al., 2012). To calculate cell tractions in cell monolayers, we used a previously described system of polydimethylsiloxane stencils (Bazellières et al., 2015) to pattern cell monolayers on rectangle-shaped monolayers. We then allowed cells to spread for 4 h and calculated tractions as previously described (Bazellières et al., 2015).

Protein quantification
Protein expression levels were measured using Western blot as previously described (Elosegui-Artola et al., 2014; Bazellières et al., 2015). For the quantification of phosphomysin light chain 2 (pMLC) and myosin light chain 2 (MLC), the membrane was first probed for pMLC, stripped using Restore Western Blot Stripping Buffer (Thermo Fisher Scientific), and then reblocked and reprobed for MLC. Protein concentrations are reported normalized to glyceraldehyde-3-phosphate dehydrogenase (GAPDH) and relative to the control.

Antibodies
Primary antibodies used were anti–ZO-1 rabbit polyclonal (61-7300; Invitrogen), anti-GAPDH mouse monoclonal (6C5; sc-32233; Santa Cruz Biotechnology), anti–myosin light chain rabbit polyclonal (3672; Cell Signaling), anti–phosphomysin light chain rabbit polyclonal (3671; Cell Signaling), anti–FLAG rabbit polyclonal (F7425; Sigma-Aldrich), anti–activated β1 mouse monoclonal (12G10; ab30394; Abcam), anti–β1 mouse monoclonal (K20; IOTest CD29-FITC; Beckman Coulter), and anti-paxillin rabbit monoclonal (Y113; ab32084; Abcam) at 1:200 for immunostainings and 1:500 for Western blot. For Western blot, the secondary antibodies used were peroxidase-conjugated anti-mouse immunoglobulin G (Gg; 715-035-151; Jackson ImmunoResearch) and peroxidase-conjugated anti-rabbit IgG (Merck Millipore, AP132P) diluted 1:5000. For immunofluorescence, the secondary antibodies used were Alexa Fluor 488 anti-rabbit (A-21206; Invitrogen) and Alexa Fluor 555 anti-mouse (A-21422; Invitrogen) diluted 1:200. To block α5β1 integrin function, the antibody used was anti-α5β1 mouse monoclonal (10 µg/ml; JC55; MAB1969; Millipore).

Immunostaining
Immunostainings on glass and gels were performed as described previously (Elosegui-Artola et al., 2014; Bazellières et al., 2015). Fluorescence images were then acquired with a 60× objective. Adhesion intensity was determined by assessing the mean paxillin intensity on a whole cell normalized to the mean intensity of the cell cytoplasm background. Adhesion density was determined manually by assessing the number of adhesions in an 11-µm² circle divided by the area. Adhesion length was measured manually by tracing a line on top of it. To quantify integrin recruitment to beads, FN7-10–coated, 3-µm carboxylated silica beads (Kisker Biotech) were attached to cells, and protein recruitment (with respect to cytoplasmic levels) was calculated assessing the fluorescence intensity of beads (Ibeam), the cytoplasm (Icytoplasm), and image background (Ibackground) as

| a.u. = Ibeam – Ibackground |
| cytoplasm – Ibackground |

(1) correlation between ZO-1 and β1 intensity images was measured by calculating Pearson’s r by using ImageJ plug-in JACoP (Bolte and Cordelières, 2006).

Preparation of polyacrylamide gels and traction measurements
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Statistical analysis

All independent datasets were first checked for normality using the d’Agostino–Pearson K2 normality test. One-way or two-way (for time-lapse experiments) analysis of variance was performed for more than two comparisons. For one-to-one comparisons, we used a two-sided t test. For multiple comparisons, we used a Dunnett modified t test. If data sets were not normal, we used a Kruskal–Wallis test. All error bars shown are SEM.

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