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N-cadherin levels in endothelial cells are regulated by monolayer maturity and p120 availability

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Abstract

Endothelial cells (EC) express VE-cadherin and N-cadherin, and recent data suggest that VE-cadherin levels are dependent on N-cadherin expression. While investigating changes in N-cadherin levels during endothelial monolayer maturation, we found that VE-cadherin levels are maintained in EC despite a decrease in N-cadherin, suggesting that VE-cadherin levels may not depend on N-cadherin. Knockdown of N-cadherin did not affect VE-cadherin levels in EC with low endogenous N-cadherin expression. Surprisingly, however, knockdown of N-cadherin in EC with high endogenous N-cadherin expression increased VE-cadherin levels suggesting an inverse relationship between the two. This was further supported by a decrease in VE-cadherin following overexpression of N-cadherin. Experiments in which p120, a catenin that binds N- and VE-cadherin, was knocked down or overexpressed indicate that these two cadherins compete for p120. These data demonstrate that VE-cadherin levels are not directly related to N-cadherin levels but may be inversely related due to competition for p120.

Keywords

adherens junction; catenin; VE-cadherin; confluence

INTRODUCTION

Endothelial cells (EC) form a monolayer located at the inner lining of all blood vessels where they serve as a non-thrombogenic surface, contribute to the regulation of vascular tone, and act as a restrictive barrier between the vessel lumen and the underlying tissue. The integrity of the endothelial barrier is dependent on sufficient cell-cell adhesion, mediated in part by the cadherin-based adherens junction. Cadherins are a superfamily of cell adhesion molecules, two of which, VE-cadherin and N-cadherin, are expressed by endothelial cells. VE-cadherin is endothelial-specific, whereas N-cadherin is expressed in several cell types and is the predominant cadherin found in neuronal cells. The overall amino acid sequences of VE- and N-cadherin share 38% homology. The regions of most similarity lie in the cytoplasmic

domains, which share 47% homology and contain the binding domains for p120 and β -catenin/plakoglobin. P120 binds the juxtamembrane region of the cadherin. The interaction of p120 with this region of the cadherin is required for maintaining the levels of cadherins by preventing endocytosis (15;30;32), and post-translational modifications of p120 have been implicated in mediating changes in adhesion strength (34). Either β -catenin or γ -catenin (plakoglobin) binds the catenin-binding region of the cadherin and through an interaction with α -catenin regulates the association of the actin cytoskeleton with the cadherin-catenin complex, although recently this model of adherens junction linkage to the actin cytoskeleton has been questioned (7;33).

Although these two cadherins have similar structures and binding partners, i.e. p120, β -catenin, and plakoglobin, N- and VE-cadherin have been shown to have different functions in the endothelium. The current model suggests that VE-cadherin mediates homotypic endothelial cell-cell adhesion, while N-cadherin mediates heterotypic contacts between EC and vascular smooth muscle cells (VSMC) or pericytes on the abluminal surface of the EC (21). Both of these cadherins have been attributed functional roles in vasculogenesis consistent with their respective localizations. There have been two transgenic mouse models of VE-cadherin deficiency, both embryonic lethal due to vascular defects (4;11). The N-cadherin gene has also been specifically deleted in the endothelium in a transgenic mouse model and, like the VE-cadherin knockouts, was embryonic lethal due to vascular defects (22). Consistent with N-cadherin's presence at regions of heterotypic cell-cell contact, several studies have strongly asserted the importance of N-cadherin recruitment of mural cells to nascent vessels during angiogenesis, which is a key step in vessel maturation (9;27;29). However, Luo and Radice noted that embryonic death occurred in the N-cadherin conditional knockout (E10.5) chronologically before investment of mural cells, suggesting that N-cadherin serves additional functions in the endothelium prior to pericyte recruitment. Diminished VE-cadherin expression in these embryos led the authors to suggest that N-cadherin may be responsible for maintenance of VE-cadherin levels in the endothelium (22). SiRNA targeting N-cadherin in cultured HUVEC (human umbilical vein endothelial cells) produced similar results supporting their hypothesis.

Studies conducted in HUVEC have shown that the composition of the endothelial adherens junction changes as the junction matures. Plakoglobin expression and junctional localization were shown to increase as confluence progressed while those of β -catenin remained fairly constant. VE-cadherin mRNA, as assessed by northern blot, also remained fairly constant, although protein levels were not assessed (19). In investigating junctional maturation, we noted that N-cadherin levels decreased as the endothelial junction matured. Interestingly, VE-cadherin levels remained constant or increased slightly as the N-cadherin levels decreased. This led us to further investigate the relationship between VE- and N-cadherin in endothelial cell monolayers. Herein, we report that VE-cadherin levels are not dependent on N-cadherin expression; furthermore, at high levels of N-cadherin expression, there is an inverse relationship between the levels of these two cadherins. In addition, we demonstrate that N-cadherin levels are not maintained by contact of endothelial cells with vascular smooth muscle cells supporting the concept that N-cadherin function is required before vessel formation and maturation, which is consistent with findings in the endothelial specific N-cadherin deficient mouse model.

METHODS

Cell Culture

Bovine pulmonary artery endothelial cells (BPAEC; VEC Technologies) were grown in Minimum Essential Medium (Cellgro) supplemented with 10% fetal bovine serum (Atlanta Biologicals), 0.1 mg/ml heparin (Sigma), and Pen/Strep (Gibco), and were split 1:2 every third day. QBI-293A human embryonic kidney cells (Quantum Biotechnology) for production of

adenovirus were grown in Dulbecco's Modified Eagle's Medium (Cellgro) supplemented with 10% fetal bovine serum, and were split 1:10 every third day. Human lung microvascular cells (HLMEC; Lonza) were plated on 20 µg/ml Vitrogen (Cohesion Technologies) and grown in EGM2-MV endothelial growth medium (Lonza) supplemented with 5% fetal bovine serum. Human pulmonary artery endothelial cells (HPAEC; Lonza) were plated on 0.2% gelatin and grown in EGM2 endothelial growth medium (Lonza) supplemented with 2% fetal bovine serum. Human umbilical vein endothelial cells (HUVEC; Cascade) were plated on 0.2% gelatin and grown in EGM2 endothelial growth medium (Lonza) supplemented with 10% fetal bovine serum. Bovine lung microvascular endothelial cells (BLMEC; Vec Technologies) were grown in Minimum Essential Medium (Cellgro) supplemented with 20% fetal bovine serum and Pen/Strep. Rat vascular smooth muscle cells (VSMC) were isolated and provided to us by the laboratory of Dr. H. Singer, and were then grown in the same media as HDMEC, with which they would be co-cultured.

Human dermal microvascular endothelial cells (HDMEC) were isolated from human foreskins. Human foreskins were collected in PBS-Pen/Strep, rinsed with ethanol and then digested with 1.2 U/ml Dispase II at 37°C for 1 hour. Following passage of tissue through a 100 µm cell strainer, a mixed population of cells was plated on fibronectin. The following day, cells were trypsinized and endothelial cells were selected from the suspension by incubating with magnetic beads coated with an antibody to CD31 (Dyna; 1e4 beads/ml cell suspension) at 4°C for 20 minutes. Non-specifically bound cells were removed by washing with 0.5% BSA in HBSS-, and endothelial cells were plated and passaged once. Cells were then assessed for VE-cadherin and lectin binding to document that they were endothelial cells. Cells were then frozen and stored in liquid nitrogen. Upon resuspension HDMECs were grown in EGM2-MV endothelial growth medium (Lonza) supplemented with 5% fetal bovine serum.

HDMEC/VSMC co-culture

We used a Transwell (Corning) co-culture system as described by Fillinger et al. (8) and Isakson et al. (14). VSMC were seeded onto 100 mm plates (no contact) or onto the underside of a 75 mm diameter 0.4 µm pore cell culture insert (contact). After 4 hours, the insert was flipped and placed in a 100 mm plate, and media was added to both top and bottom chambers. The following day, HDMEC were seeded onto 3 cell culture inserts. One was placed into an empty 100mm plate, one was placed in the 100 mm plate containing VSMC, and the third was the insert with VSMC growing on the opposite side. Fresh media was added to the top and bottom chambers of each plate. RNA was isolated (from HDMEC only) 96 hours after seeding HDMEC, and VE-cadherin, N-cadherin, and GAPDH mRNA levels were determined by qRT-PCR.

RNA Interference

siRNA was purchased from either Dharmacon (Human N-cadherin SmartPools + OnTarget Plus chemically modified individual sequences, negative control siRNA) or Ambion (pre-designed siRNA sequences for human N-cadherin: siRNA ID #10506 (N06; GGCCAAACAACUUUAAUU) and siRNA ID #10325 (N25; GGCUCUGGUGAAAUCGCA); custom siRNA design for the following target sequences: bovine N-cadherin GGAAUCACGAGAAAUAGAA, bovine p120 GCCAGAGAUCAGAUAAAGAA). SiRNA was delivered into cells (N-cadherin: 0.7 µg siRNA per 375,000 cells, p120: 1.0 µg siRNA per 375,000 cells) via electroporation using Ambion siPort electroporation buffer and the BioRad GenePulser Xcell Electroporation System. Knockdown was evaluated by Western blot.

Quantitative Real Time PCR

Copy number was calculated using the standard curve method described by Leong et al. (20). Briefly, two sets of primers were used. Primers positioned to generate slightly larger DNA

fragments were used to produce standards. Spectrometer readings were used to determine concentrations of standards in $\mu\text{g}/\mu\text{l}$. These concentrations were converted to number of copies, and serial dilutions were used to produce standards of known copy number. These standards were run simultaneously with cDNA samples using the primer sets positioned to generate smaller DNA fragments (i.e., primers that would recognize the standards as well as the cDNA samples), and copy number of samples was extrapolated by plotting threshold cycle vs. log copy number of the standards. Primers were as follows (5' – 3'): human VE-cadherin – forward: GGCAAGATCAAGTCAAGCGTG; human VE-cadherin – reverse: ACGTCTCCTGTCTCTGCATCG (16); N-cadherin – forward: GCCCTCAAGTGTTACCTCAA; N-cadherin – reverse: AGCCGAGTGATGGTCCAATTT (35). Primers for production of standards were as follows (5' – 3'): human VE-cadherin – forward: CCTCACTTCCCCATCATGTA; human VE-cadherin – reverse: CCAGCCTCTCAATGGCGAAC; N-cadherin – forward: TACTTGATATTAATGACAAT; N-cadherin – reverse: GCTGAGCAAAATCACCATTA. Reverse transcription was performed using the iScript cDNA Synthesis Kit (Bio-Rad). Quantitative real time PCR was carried out using the Bio-Rad iCycler system and SYBR green mastermix (Bio-Rad).

Adenovirus

Generation of adenoviral recombinants was performed as described by He et al. (12). The pAdTrack-CMV, p-Shuttle-CMV shuttle vector, and pAdEasy-1 vector were kindly provided by Dr. B Vogelstein. Recombinant adenovirus was amplified in QBI-293A cells and purified using CsCl_2 gradients. All infections were accompanied by a control GFP infection with MOI at or above greatest MOI used in the experimental groups. The p120-GFP cDNA construct was obtained from Dr. Keith Burridge. Myc-epitope tagged chimeric IL-2R/VE-cadherin constructs have been described previously (30), (3). IL-2R-VE-cad_{cyto} contains the extracellular and transmembrane portions of the IL-2 receptor and the intact cytoplasmic domain of VE-cadherin. IL-2R-VE-cad_{AAA} contains the extracellular and transmembrane portions of the IL-2 receptor and the VE-cadherin cytoplasmic domain with the AAA mutation at amino acids 562–564, and thus is incapable of binding p120.

Immunoprecipitation

Cells were lysed in Cytoskeleton Preserving Buffer (10 mM PIPES 100 mM NaCl, 300 mM sucrose, 3 mM MgCl_2 , 0.5% Igepal, 1 mM PMSF, 5 $\mu\text{g}/\text{ml}$ leupeptin, 2 $\mu\text{g}/\text{ml}$ aprotinin, 1 mM Na_3VO_4 , 1 mM EDTA), and lysates were precleared by centrifugation. Supernatants were incubated with primary antibody (p120 – Santa Cruz Biotechnology) for 1 hour at 4°C, then Protein A agarose beads (Invitrogen) were added and incubation was continued at 4°C for 1 more hour. Beads were washed 3 times in buffer; immunoprecipitation procedure was repeated with the supernatant; Laemmli gel sample buffer was then added to the beads and boiled to remove bound proteins, and samples were analyzed by Western Blot.

Gel Electrophoresis and Western Blot

At designated time points following experimental procedures cells were lysed in Laemmli gel sample buffer, and lysates were run on SDS-PAGE and transferred to nitrocellulose membranes. Primary antibodies used were as follows: VE-cadherin, p120 (Santa Cruz); β -actin (Sigma); N-cadherin (BD Transduction Laboratories; ECM biosciences); Myc-epitope tag (Cell Signaling Technologies). Secondary antibodies were HRP-conjugated goat-anti-mouse, rabbit-anti-goat, or goat-anti-rabbit (Jackson ImmunoResearch). Super Signal West Pico and West Femto Chemiluminescent Substrates were purchased from Pierce Biotechnology. Bands were visualized using the Fuji LAS-3000 Image Station. Densitometric analysis was performed using Fujifilm Multigauge 3.0 software.

Immunofluorescence Microscopy

Cells seeded on 0.1% gelatin-coated coverglasses were fixed in 4% paraformaldehyde, permeabilized in TBS with 0.1% Triton X-100, and blocked in 5.5% goat, donkey, or fetal bovine serum in TBS. Cells were then incubated with primary antibodies to VE-cadherin (Santa Cruz Biotechnology) or N-cadherin (Zymed) in TBS and fluorescent secondary antibodies (Molecular Probes, Jackson ImmunoResearch) in 3% BSA in TBS.

RESULTS

N-cadherin levels decrease during endothelial monolayer maturation

We were interested in studying the relationship between VE-cadherin and N-cadherin in the endothelium and first sought to determine whether VE- and N-cadherin levels, similar to plakoglobin levels, changed as cells become confluent. Human dermal microvascular endothelial cells (HDMEC) were plated at a density of 8×10^4 cells/cm², and phase contrast images were taken to document the degree of confluence reached at 24, 48, 72, and 96 hours after plating (Figure 1A). Within 24 hours, the monolayer appeared nearly confluent, and throughout the following 72 hours we observed a further increase in cell density. RNA and protein were also isolated at these time points. Immunoblots revealed that VE-cadherin levels remained constant or slightly increased over this time course (Figure 1B, left) while N-cadherin levels decreased sharply between 24 and 48 hours after plating, continuing to decrease over the course of the experiment (Figure 1B, right). N-cadherin is present in many endothelial cells at the 24 hour time point and localized diffusely with some junctional localization, and a decrease in N-cadherin labeling intensity is seen at 96 hours (Figure 1C, right panels). In contrast, VE-cadherin labeling intensity increased at 96 hours compared to 24 hours (Figure 1C, left panels). Determination of N-cadherin mRNA levels using quantitative real-time PCR (Figure 1D, right) showed a decrease between 24 and 48 hours, implying that the decrease in N-cadherin protein was due to a decrease in message. VE-cadherin mRNA levels, however, remained constant throughout the 96 hour maturation period, in agreement with Lampugnani et al. (19).

As shown in Figure 1B, as HDMEC progressed to confluence, VE-cadherin levels were maintained despite a decrease in N-cadherin levels suggesting that VE-cadherin is not dependent on N-cadherin levels. To further investigate the relationship between N- and VE-cadherin, we performed experiments using N-cadherin siRNA similar to those previously performed in HUVEC (22). We first used the pool of sequences to produce successful knockdown of N-cadherin in HUVEC (siGENOME SmartPools from Dharmacon, Inc). This pool of siRNA (Nsmrt) was delivered into HUVEC by electroporation, and N- and VE-cadherin were assessed by Western blot 48 hours later. As shown in Figure 2, N-cadherin was successfully knocked down (Figure 2, top left), and VE-cadherin levels were not altered (Figure 2, top right). As RNA interference has become a more prevalent tool, methods to chemically modify siRNAs in order to minimize off-target effects have been developed, and these modified siRNAs have now become commercially available. Thus, to control for any possible off-target effects, we used additional siRNA sequences designed by two different companies. We first purchased the chemically modified pool of sequences targeting human N-cadherin from Dharmacon (OnTargetPlus SmartPools). We screened the four supplied sequences for efficiency of N-cadherin knockdown, and two of these sequences successfully depleted N-cadherin 48 hours after electroporation (N6D - Figure 2, middle panels and N9D – not shown). Again we did not see a change in VE-cadherin levels. Finally, we used a chemically modified (by Dharmacon) version of a sequence we had originally obtained from Ambion Inc. (N25mod). Similar to the other sequences we used, N25mod also did not decrease VE-cadherin levels but did decrease N-cadherin levels (Figure 2, bottom panels). Thus we have concluded that N-cadherin expression is not required for maintenance of VE-cadherin levels in HUVEC.

Heterogeneity of N-cadherin expression in endothelial cell types

Our lab has worked with several different endothelial cell types, and we have observed that endothelial cells from different species and from different vascular beds express varying levels of N-cadherin protein. As shown in Figure 3, confluent HLMEC (human lung microvascular endothelial cells), HUVEC, HPAEC (human pulmonary artery endothelial cells), and BPAEC (bovine pulmonary artery endothelial cells) display different N-cadherin expression levels. The extremely low expression of N-cadherin in HLMEC relative to the other EC types was associated with high VE-cadherin expression providing further evidence that VE-cadherin levels are not dependent on N-cadherin levels. In addition it appears that low N-cadherin expressors (HLMEC) had high VE-cadherin expression whereas high N-cadherin expressors (BPAEC) had low VE-cadherin expression; HUVEC and HPAEC fell between the two extremes. These data suggest that there may be an inverse relationship between N-cadherin and VE-cadherin at high expression levels of one of these cadherins. To determine whether the differences in cadherin expression were due to tissue of origin, we compared human and bovine microvascular endothelial cells from the same tissue of origin (HLMEC and BLMEC); BLMEC expressed N-cadherin abundantly, in contrast to HLMEC (data not shown) suggesting that bovine cells have a greater level of N-cadherin than human cells.

Changes in N-cadherin expression levels affect VE- cadherin levels, and vice versa

In order to determine whether the relatively low VE-cadherin expression in BPAEC was related to their high expression of N-cadherin, we depleted N-cadherin in these cells using siRNA. The decrease in N-cadherin resulted in an increase in VE-cadherin (Figure 4A) supporting the idea that VE-cadherin levels are not dependent on N-cadherin, and also implying that the endogenously high N-cadherin expression in these cells limits VE-cadherin levels. This result further supports the idea that there is actually an inverse relationship between N-cadherin and VE-cadherin levels, rather than a direct relationship. Next, we expressed either N-cadherin or VE-cadherin using adenovirus containing Flag epitope-tagged-cadherin (AdN-cad-flag or AdVE-cad-flag) to determine if each would cause a decrease in the level of the other cadherin. Cells overexpressing N-cadherin displayed decreased levels of VE-cadherin in comparison to GFP-infected control BPAEC (Figure 4B) whereas cells with increased expression of VE-cadherin showed a decrease in the level of N-cadherin (Figure 4C, top). A corresponding decrease in N-cadherin labeling was also observed by immunofluorescence microscopy (Figure 4C, bottom panels). These data demonstrate an inverse relationship between N- and VE-cadherin, such that when levels of one are high the levels of the other are decreased. Since p120, an intracellular binding partner of cadherins, is known to maintain VE-cadherin levels (15) by preventing endocytosis of VE-cadherin in endothelial cells (30), we hypothesized that VE- and N-cadherin compete for p120 association in order to be stabilized at the membrane.

VE- and N-cadherin levels are dependent on p120

Previous studies have found that p120 maintains N-cadherin levels in neuronal cells (6), but this has not been shown in endothelial cells. Thus, we first depleted p120 in BPAEC using siRNA targeted to bovine p120 (Figure 5A). As anticipated, VE-cadherin levels decreased (Figure 5A, center), consistent with our previous findings that binding of p120 is required for maintaining VE-cadherin levels at the membrane (15). N-cadherin levels decreased to a similar extent upon depletion of p120 (Figure 5A, right), showing that p120 is required for maintenance of N-cadherin levels in endothelial cells. We next expressed exogenous p120 in order to increase p120 availability. Infecting confluent monolayers of BPAEC with adenovirus containing p120-GFP resulted in an increase in the levels of VE-cadherin. Interestingly, no significant change in the levels of N-cadherin (Figure 5B) was observed with increased expression of p120 suggesting that, although N-cadherin is dependent on p120 association, basal levels of N-cadherin are not limited by p120 availability in the endothelium.

To determine whether the level of association between each cadherin and p120 was different, we performed two successive immunoprecipitations of p120 in BPAEC lysates to remove p120 and all its associated proteins from the lysate. Both cadherins were clearly present in the p120 immunoprecipitate (p120 IP, Figure 5C, left). We also assessed the level of VE-cadherin or N-cadherin not associated with p120 by immunoblotting the remaining lysate following each IP (Post-IP, Figure 5C center) and comparing this to the whole lysate (Pre-IP). Densitometric analysis was performed on the VE- and N-cadherin immunoblots, and values were normalized to GAPDH to account for any dilution of the samples that occurred during the experiment. These values were used to approximate the percentage of total VE- and N-cadherin not bound to p120 ($100 * [\text{VE- or N-cadherin in Post-IP lysate} + \text{VE- or N-cadherin in Post-IP lysate}] / \text{VE-cadherin or N-cadherin in Pre-IP}$). These values were then subtracted from 100%, revealing that $82.61 \pm 3.60\%$ (65.25% in IP1 + 17.36% in IP2) of the total VE-cadherin and $75.54 \pm 7.71\%$ (63.25% in IP1 + 12.28% in IP2) of the total N-cadherin co-precipitated with p120 (Figure 5C, right). Thus, the percentage of N-cadherin associated with p120 is similar to that of VE-cadherin in a confluent monolayer expressing both of these cadherins.

The experiments in Figures 5A, B, and C show that VE-cadherin and N-cadherin both associate with and are dependent on p120. We used a VE-cadherin mutant incapable of binding p120 in order to directly test whether competition for p120 was responsible for the decrease in N-cadherin upon VE-cadherin overexpression observed in Figure 4C. The extracellular and transmembrane portions of the IL-2 receptor were fused to VE-cadherin cytoplasmic tail either without (AdIL-2R-VE-cad_{cyto}) or with (AdIL-2R-VE-cad_{AAA}) a mutation in the juxtamembrane domain which renders this mutant of VE-cadherin incapable of interacting with p120 (30). Expression of the intact cytoplasmic tail of VE-cadherin decreased N-cadherin levels, whereas the p120-uncoupled mutant did not (Figure 5D, top panel). Thus, competition for p120 is responsible for the inverse relationship between VE- and N-cadherin protein levels.

Regulation of N-cadherin mRNA levels

In the absence of mural cells (pericytes and/or VSMC), we showed that N-cadherin mRNA and protein expression decreased over time as HDMEC grew into a confluent monolayer (Figure 1). Co-culture of endothelial cells with mural cells has been shown in several studies to regulate endothelial cell behavior (17;18;23;24;26;28). In one of these studies, N-cadherin localization was also found to shift from EC-EC junctions to heterotypic junctions in the presence of pericytes (18). Therefore, we hypothesized that heterotypic contact of HDMEC with mural cells would prevent the drop in N-cadherin expression with confluence that we saw in Figure 1. In order to discern effects of soluble factors produced by mural cells from those of direct cell-cell contact, we used two Transwell co-culture systems (Figure 6A): one in which endothelial cells and VSMC were seeded on opposite sides of a $0.4 \mu\text{m}$ filter which allowed direct contact (8;14), and another in which the EC were seeded on the filter, and the VSMC were seeded on the bottom of the cell culture plate not allowing direct heterotypic contact. VE- and N-cadherin mRNA expression in HDMEC was assessed 96 hours after plating the HDMEC. Unexpectedly, N-cadherin mRNA levels not only failed to be maintained, but were reduced in both co-culture systems as compared to HDMEC in monoculture (Figure 6B). These data suggest that although N-cadherin is required for recruiting mural cells (10;27;29), heterotypic contact between these cells does not maintain N-cadherin mRNA expression as a nascent vessel matures. VE-cadherin mRNA expression was unchanged by co-culture (Figure 6B).

Since heterotypic contact with VSMC simulates vessel maturation, we hypothesized that activated EC would show increased N-cadherin expression as compared to quiescent EC. To test this hypothesis, we incubated confluent HDMEC (72 hours post-seeding) in either basal medium (EBM) or 5% serum-supplemented basal medium (5%) for 24 hours, and assessed N-

and VE-cadherin mRNA levels. As shown in Figure 6C, serum stimulation increased N-cadherin mRNA expression and had no effect on VE-cadherin mRNA expression. Thus, N-cadherin expression is negatively regulated at the level of message by EC monolayer maturation (Figure 1C, Figure 6B), and positively regulated by activating stimuli such as growth factors present in serum (Figure 6C).

DISCUSSION

Changes in the expression of adherens junction components during endothelial monolayer maturation have been previously reported: plakoglobin (γ -catenin) expression and localization were shown to increase as confluence progressed while those of β -catenin remained fairly constant (19). We investigated the relationship between N-cadherin and VE-cadherin levels in developing endothelial monolayers and made an interesting observation: as an endothelial monolayer matures, N-cadherin levels decrease while VE-cadherin levels remained the same or increased slightly. The finding that VE-cadherin remained elevated in the absence of N-cadherin was inconsistent with the conclusion of Luo and Radice - that VE-cadherin levels are dependent on the level of N-cadherin - thus, we performed further experiments to investigate the relationship between N- and VE-cadherin in endothelial monolayers.

Through the use of pooled siRNA sequences and several different individual siRNA sequences from two different commercial sources, we demonstrated that VE-cadherin levels are not decreased by a decrease in N-cadherin levels following treatment of HUVEC with N-cadherin siRNA. However, using the same pool of sequences purchased from Dharmacon, Inc., Luo et al. (22) did observe a decrease in VE-cadherin levels. Several factors could have contributed to this discrepancy. For example, we used electroporation rather than lipid-based transfection to deliver siRNA. Our laboratory has used both methods, and we have achieved greater efficiency with the electroporation method. Notably, this difference in methods involves delivery of siRNA to the cells at different degrees of confluence: Electroporation is performed on cells while in suspension, therefore our cells were plated after siRNA had been delivered, whereas lipid-based transfection was carried out by Luo et al. in 70% confluent cells. Different doses of siRNA could also be a factor. In our electroporation method, we used a specified mass of dsRNA per cell number (0.7 μ g per 3.75e5 cells) rather than a molar concentration because the electroporation is carried out in a very small volume (75 μ l). In reference to culture volume in which the cells are then plated, this concentration would translate to approximately 50 nM. Luo et al. used a concentration of 100 nM to knock down N-cadherin in their system. We cannot be sure how these two doses of siRNA compare because of the difference in delivery, however, higher doses of siRNA are believed to be associated with a greater likelihood of off-target effects.

The conclusion that VE-cadherin levels are not dependent on N-cadherin expression is further supported by the findings in Figure 3 - heterogeneous expression of these two cadherins in endothelial cells from different species and different vascular beds. The heterogeneity in N-cadherin expression that we have observed is consistent with the global heterogeneity in endothelial cells that has been previously documented (reviewed in (1;2)). Of note is that endothelial cells with low N-cadherin expression (HLMCEC) display similar or elevated VE-cadherin levels as compared to cells that have a greater level of N-cadherin. Also, BPAEC, which had the greatest amount of N-cadherin had the lowest amount of VE-cadherin. This would suggest that N- and VE-cadherin levels may be inversely related rather than directly related. This idea is supported by studies in Figure 4, showing that in addition to not being required for maintaining VE-cadherin levels, N-cadherin expression, at high levels, in fact negatively regulates VE-cadherin expression. The result shown in Figure 4A is especially notable. If VE-cadherin levels were indeed dependent on N-cadherin, one would expect a decrease in VE-cadherin upon treatment of high N-cadherin expressing cells with N-cadherin

siRNA. However, we saw the opposite in these cells. The increase in VE-cadherin observed following depletion of N-cadherin in BPAEC implies that VE-cadherin levels are actually limited in these cells by the large quantity of endogenous N-cadherin. Our overexpression studies are consistent with this idea. As shown in Figure 4B, exogenous expression of N-cadherin in BPAEC produces a significant decrease in VE-cadherin levels. Likewise, overexpression of VE-cadherin reduced N-cadherin levels (Figure 4C). These results imply an inverse relationship between levels of the two cadherins in endothelial cells, which suggests competition between the two for a protein that stabilizes each at the membrane.

When speaking of cadherin levels, it is crucial to discuss p120, an intracellular binding partner of cadherins that is required for maintenance of cadherin levels. P120 is a member of the Armadillo family of proteins and a component of the cadherin-based adherens junction. P120 functions in cell-cell adhesion, signal transduction, and regulation of gene transcription. Previous studies from our lab have demonstrated that p120 is essential for the maintenance of VE-cadherin levels and barrier function of the endothelium (15), and other studies have shown that this depletion of VE-cadherin following loss of p120 binding is due to clathrin-dependent endocytosis of VE-cadherin (31). These data have established p120 as a point of control for regulation of VE-cadherin levels in endothelial cells, and similar data exist in several adhesive cell types (reviewed in (32)). Note, however, that dependence on p120 for maintained expression of N-cadherin has not previously been shown in endothelial cells. Indeed, p120 is required for maintaining N-cadherin as siRNA targeting p120 resulted in decreases in both VE-cadherin and N-cadherin levels (Figure 5A) Furthermore, by overexpression of p120 in BPAEC monolayers, we demonstrated that increased availability of p120 allowed for increased VE-cadherin levels (Figure 5B). Similar results were obtained in HLMEC (data not shown). This suggests that cadherin levels in the endothelial cell are limited by the available amount of p120.

Consistent with previous reports (25), p120 co-immunoprecipitates both VE-cadherin and N-cadherin (Figure 5C). In addition, we quantified by densitometry the percentage of each cadherin associated with p120 by assessing the amount of N and VE-cadherin remaining in the lysate following immunoprecipitation of p120 ($100 - [100 * (\text{Post-IP1} + \text{Post-IP2}) / \text{Pre-IP}]$). The vast majority of each cadherin (approximately 83% of the VE-cadherin and 76% of the N-cadherin) was bound to p120. Taken together with our p120 siRNA data, we conclude that N-cadherin, like VE-cadherin, is dependent on p120 association for maintaining levels at the membrane. Therefore, N-cadherin is regulated at the mRNA level by monolayer maturity and at the protein level by p120 availability. Interestingly, when we increased p120 availability in BPAEC via overexpression of p120 in confluent monolayers, we did not see an increase in N-cadherin levels suggesting protein levels of N-cadherin were likely limited by the level of mRNA expression in confluent monolayers.

Navarro et al. (25) observed greater p120 band intensity in the VE-cadherin IP compared to the N-cadherin IP. They also observed diffuse surface localization of N-cadherin in contrast to localization of VE-cadherin at the endothelial cell-cell junction (25). Their interpretation of these data was that p120 bound more abundantly to VE-cadherin than to N-cadherin due to a greater affinity of p120 for VE-cadherin, and that consequently N-cadherin would be excluded from EC-EC contacts. Although we did not directly test binding affinity, we do establish herein that similar percentages of each cadherin, which constitute the vast majority of each cadherin, co-precipitate with p120 demonstrating that the level of association between each cadherin and p120 is similar. Indeed, under control conditions approximately 80% of each cadherin is associated with p120 (Figure 5C). The levels of these two cadherins also showed similar dependence on p120 expression as decreased p120 availability results in a loss of both cadherins at the level of total protein (Figure 5A). When levels of one cadherin were elevated we observed a decrease in total protein of the other cadherin (Figure 4B), presumably due to endocytosis in

the absence of p120. When a p120-uncoupled VE-cadherin mutant was expressed in BPAEC, N-cadherin did not decrease (Figure 5D), thus competition for p120 was the mechanism by which excess levels of one cadherin resulted in downregulation of the other. This result plus the decreases in total N-cadherin seen in Figure 4C (top) and Figure 5A suggest that if VE-cadherin sequesters a greater amount of the available p120 in EC due to it having a greater affinity, then N-cadherin would be endocytosed and degraded. This decrease in N-cadherin level could result in a redistribution of N-cadherin away from the EC-EC junctions as observed by Navarro et al. (23). Indeed, we observed a decrease in junctional N-cadherin staining (Figure 4C, bottom panels) in conjunction with the decrease in total N-cadherin protein (Figure 4C, top) upon overexpression of VE-cadherin in BPAEC.

Increased expression of N-cadherin has been implicated as part of the angiogenic phenotype, playing a role in the recruitment of pericytes/smooth muscle cells to newly forming vessels (12–14). Serum factors are known to play a role in inducing angiogenesis, thus we treated confluent monolayers with serum to determine whether this activation of the EC would induce N-cadherin expression. Indeed, activation of confluent cells by addition of serum increased N-cadherin mRNA expression (Figure 6C). This expression pattern indicates that N-cadherin is important in disrupted endothelium, e.g. during angiogenesis, which is consistent with the functional roles that have been attributed to N-cadherin in mural cell recruitment (9;27;29) and in vascular development as suggested by the endothelial specific N-cadherin knockout (22). Our data further supports the idea that N-cadherin is required during development for mural cell recruitment but is not required for vessel maturation. This is supported by the finding that N-cadherin expression decreases with formation of a more mature junction (Figure 1) and that co-culture further decreased N-cadherin mRNA expression, in both direct contact and non-contact systems (Figure 6A, B). Somewhat surprising was the finding that N-cadherin decreased to a greater extent in the co-culture system than it did in confluent monocultured HDMEC (Figure 6A, B; Figure 1). However, a change in gene expression of a protein is consistent with studies showing that paracrine signaling between endothelial cells and mural cells is essential for vessel stabilization (reviewed in (9)), and regulates several functions in both cell types including gene expression (5;17), survival (5;17), endothelial monolayer permeability (18;23;24), migration (13;28), and proliferation (26).

Our data taken together with these functional data support the following model: N-cadherin is upregulated in disrupted endothelial cells, as it is essential for the maturation of newly forming vessels. As the endothelial monolayer begins to mature and is stabilized by mural cells, N-cadherin is no longer required and is consequently downregulated. Ultimately, downregulation of N-cadherin may result in increased p120 availability, enabling an increase in VE-cadherin levels and a mature, restrictive endothelial monolayer.

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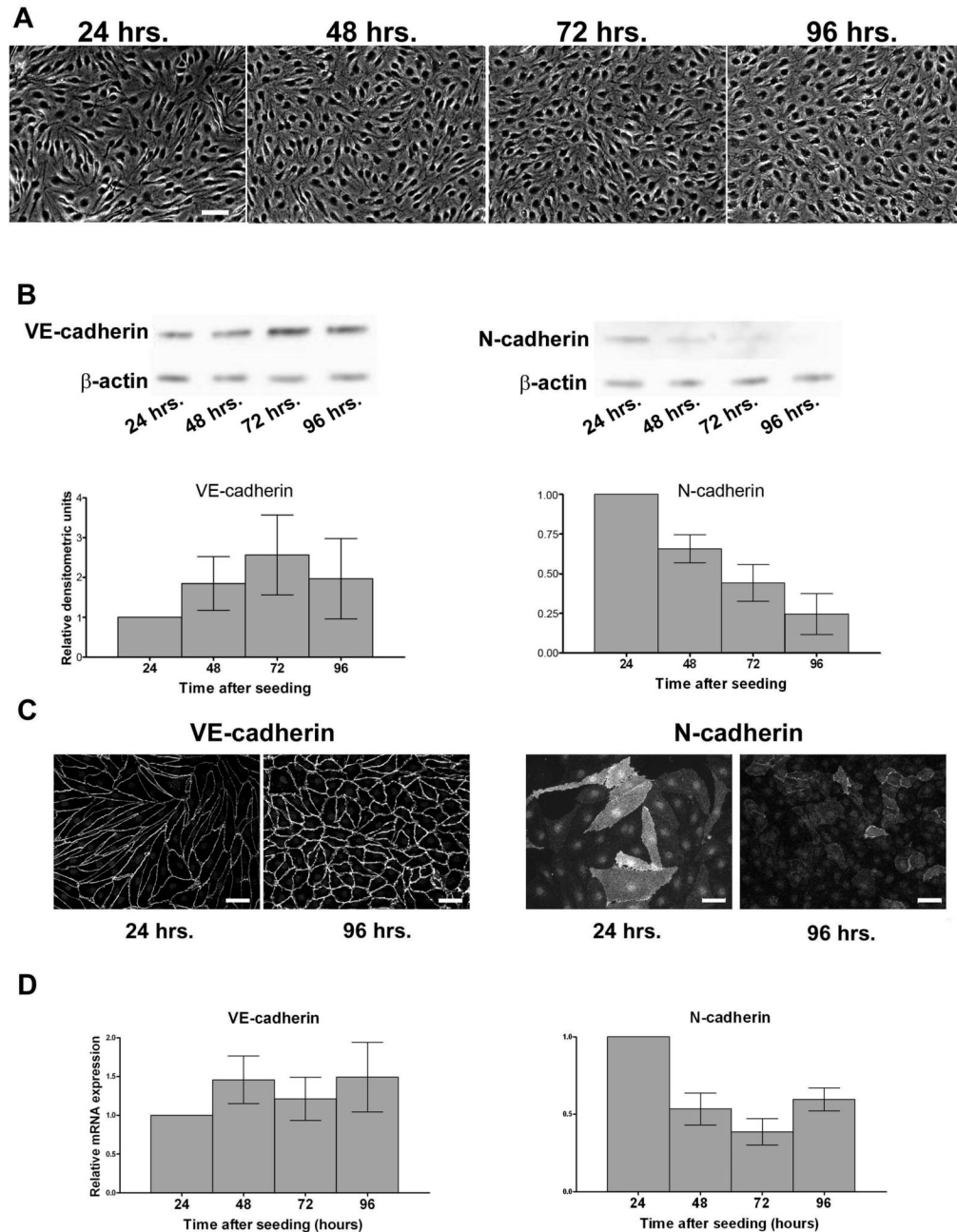


Figure 1. N-cadherin expression decreases as the endothelial monolayer matures

(A) Human dermal microvascular endothelial cells (HDMEC) were plated at a density of 8.0×10^4 cells/cm², and phase contrast images were taken every 24 hours to document degree of confluence. Bar = 50 μ m. (B) Top: cells were lysed at each time point, and proteins were separated by SDS-PAGE. Following transfer to nitrocellulose membranes, lysates were immunoblotted for VE-cadherin, N-cadherin, and β -actin as a loading control. Bottom: densitometric analysis of immunoblots. Values are normalized to β -actin and are expressed relative to the 24 hour time point (mean \pm SEM; n=6). N-cadherin: P=0.0011 by one-way ANOVA. (C) Localization of VE-cadherin and N-cadherin in HDMEC during growth to confluence: cells plated at 8.0×10^4 cells/cm² were fixed at the 24 or 96 hour time points, and

immunofluorescence microscopy was performed by incubating with antibodies to either VE-cadherin (left panels) or N-cadherin (right panels) followed by fluorescent-conjugated secondary antibodies. Bar = 50 μ m. (D) RNA was isolated at each time point, and quantitative real-time PCR was performed using primers to VE-cadherin, N-cadherin, and GAPDH, and copy number was determined using standard curves (20). Values for each time point are expressed relative to the 24 hour time point, and are normalized to GAPDH (mean \pm SEM; n=6). N-cadherin mRNA: P=0.0167 by one-way ANOVA.

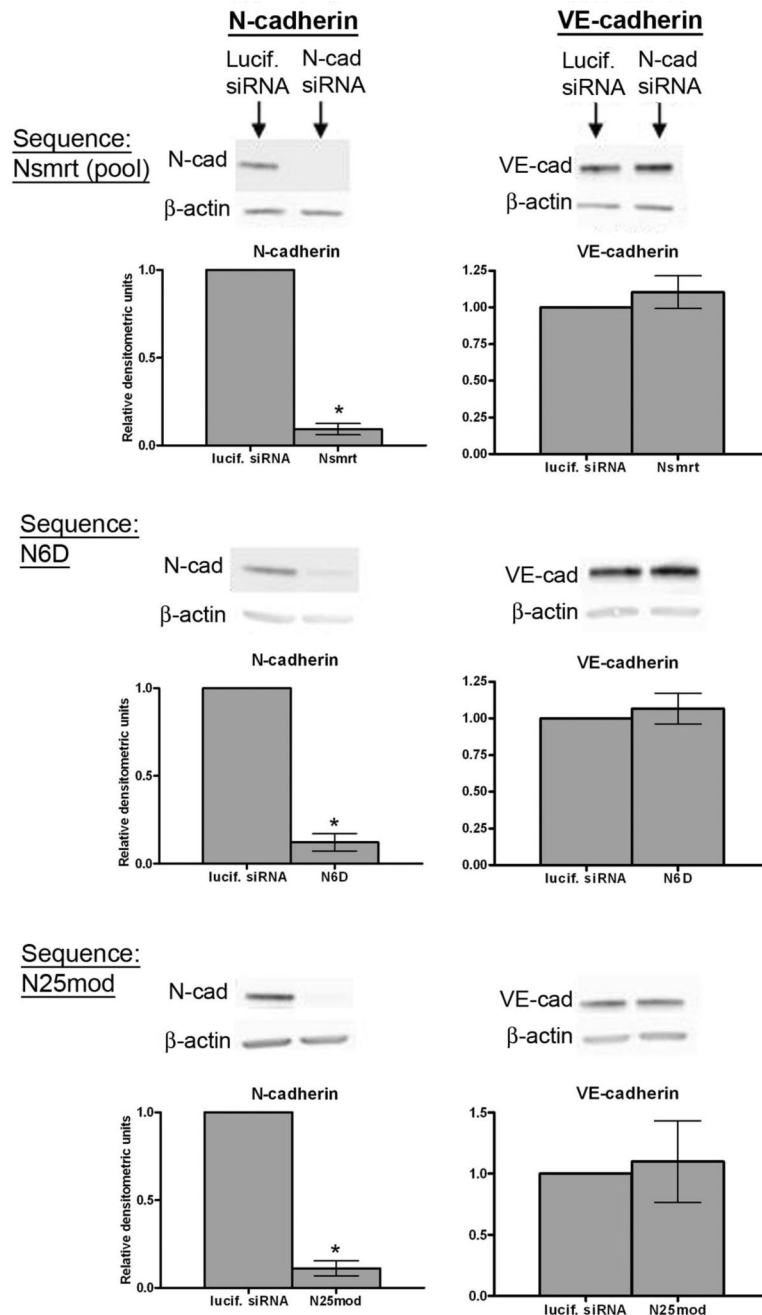


Figure 2. VE-cadherin levels in HUVEC are maintained in the absence of N-cadherin
 (A) SiRNA sequences targeting either luciferase (Lucif. siRNA) or N-cadherin (Nsmrt, N6D, or N25mod) were delivered into HUVEC via electroporation, and cells were plated at confluence. Immunoblot analysis for N-cadherin (left), VE-cadherin (right), and β -actin was carried out 48 hours after plating. Densitometric analysis of immunoblots is shown below. Values for N-cadherin and VE-cadherin are normalized to β -actin and are expressed relative to luciferase control (mean \pm SEM; minimum n=3, for N6D $P < 0.0001$, N25mod $P = 0.0024$, Nsmrt $P < 0.0001$). N6D: individual chemically modified sequence purchased from Dharmacon. Nsmrt: N-cadherin siGENOME Smartpools (unmodified) purchased from Dharmacon.

N25mod: individual sequence purchased from Ambion, subsequently chemically modified by Dharmacon.

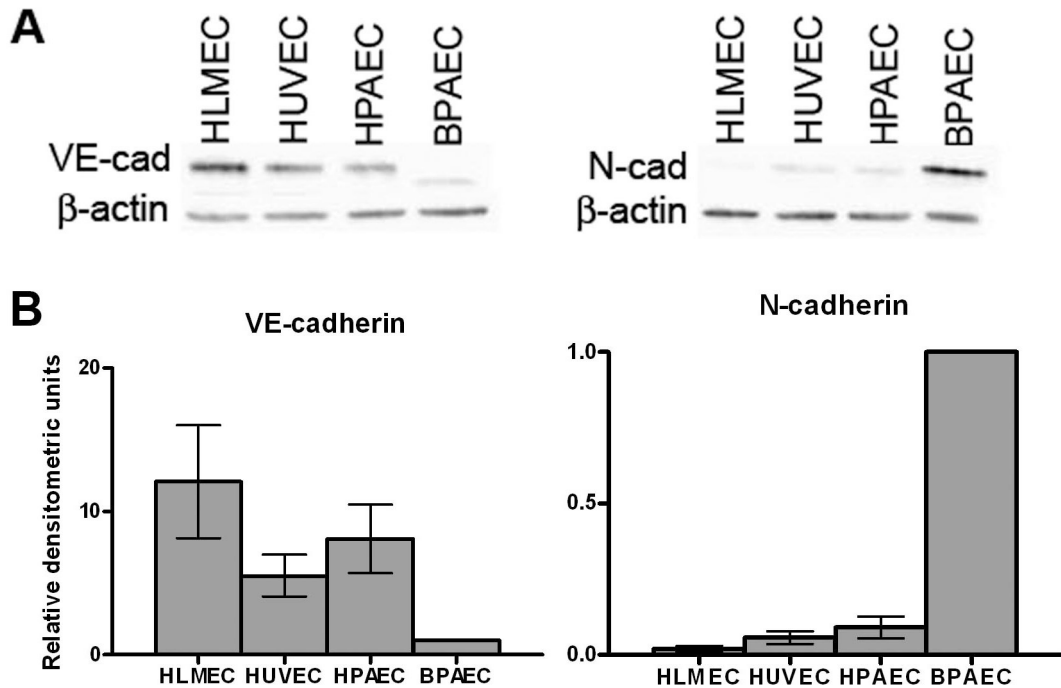


Figure 3. N-cadherin protein expression varies in endothelial cells from different sources

(A) Confluent monolayers of HLMEC, HUVEC, HPAEC, and BPAEC were lysed 72–96 hours after plating, and proteins were separated by SDS-PAGE. Following transfer to nitrocellulose membranes, lysates were immunoblotted for VE-cadherin (left) and N-cadherin (right). β -actin was used as a loading control. (B) Densitometric analysis of immunoblots shown in (A). Densitometric values were normalized to β -actin and values are expressed relative to BPAEC. (mean \pm SEM)

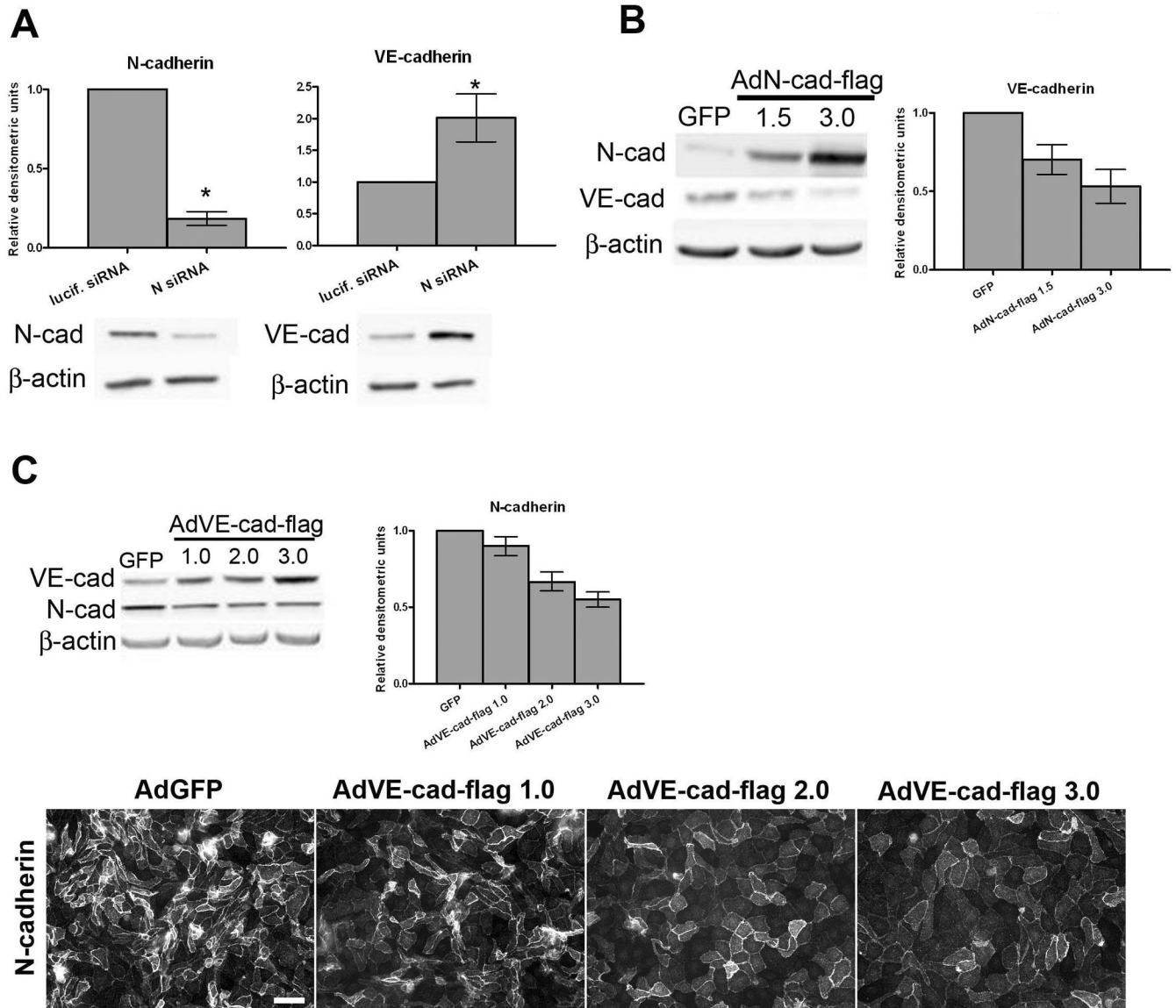


Figure 4. Relationship between VE-cadherin and N-cadherin protein levels in BPAEC
 (A) siRNA sequences targeting either luciferase or N-cadherin were delivered into BPAEC via electroporation, and cells were plated at confluence. Immunoblot analysis for N-cadherin, VE-cadherin, and β -actin was carried out 48 hours after plating. Densitometric analysis: Values for N-cadherin and VE-cadherin are normalized to β -actin and are expressed relative to luciferase control (mean \pm SEM, n=8; for N-cad P<0.0001, VE-cad P=0.0321 using single sample t-test. N siRNA: individual sequence targeting bovine N-cadherin purchased from Ambion.) * denotes significance (p<0.05) as determined using single sample t-test. (B) Confluent BPAEC monolayers were infected with either adenovirus containing GFP or one of two doses of flag epitope-tagged N-cadherin (AdN-cad-flag). After 48 hours, cells were lysed, and N-cadherin and VE-cadherin were detected in lysates by immunoblot analysis. β -actin was used as a loading control (bottom panels). (C) Top: Confluent BPAEC monolayers were infected with either GFP or one of three doses of VE-cadherin-flag adenovirus (AdVE-cad-flag) and immunoblotted for N-cadherin and VE-cadherin. Right – densitometric analysis:

Densitometric values for cadherins are normalized to β -actin expressed relative to control GFP infection (mean \pm SEM, n=3; for AdN-cad-flag, VE-cad P=0.0197 by one way ANOVA; for AdVE-cad-flag, N-cad P<0.0001 by one way ANOVA). Bottom: Following the 48 hour infection with AdVE-cad-flag, immunofluorescence microscopy was performed on BPAEC using an antibody to N-cadherin followed by a fluorescent-conjugated secondary antibody. Note the decrease in N-cadherin staining intensity with increasing dose of AdVE-cad-flag (left to right). Bar = 50 μ m.

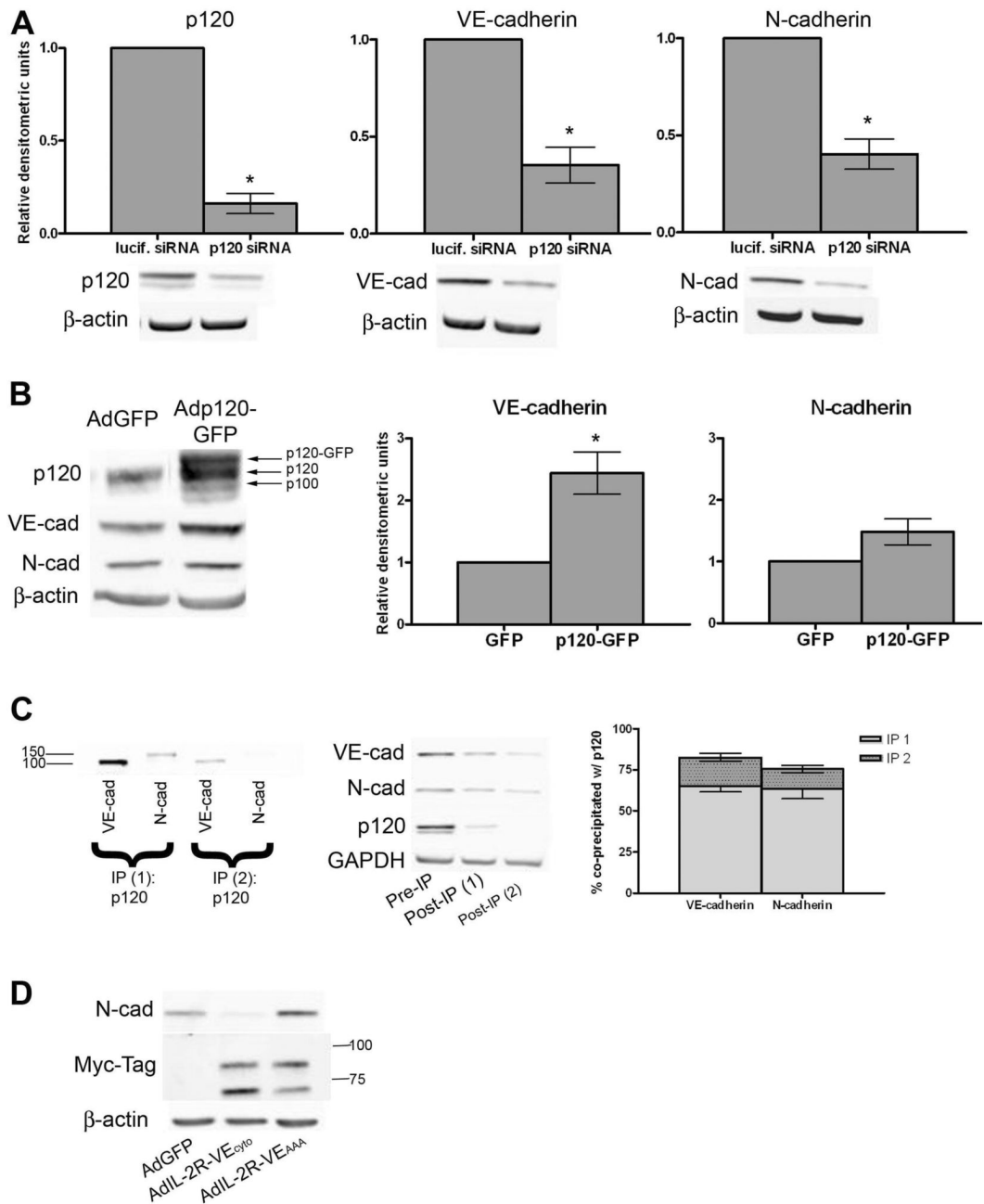


Figure 5. Regulation of cadherin levels by p120

(A) siRNA targeted to either luciferase (lucif. siRNA) or p120 was delivered via electroporation into BPAEC. After 48 hours, cells were lysed and immunoblot analysis was carried out for p120 (left), VE-cadherin (center), and N-cadherin (right). β -actin was used as a loading control (not shown). Densitometric values are normalized to β -actin and are expressed relative to luciferase siRNA (mean \pm SEM, n=3; for p120 P=0.0043, VE-cad P=0.0199, N-cad P=0.0166). (B) Confluent monolayers of BPAEC were infected with either GFP adenovirus or p120-GFP adenovirus. 48 hours after infection, cells were lysed and immunoblot analysis was carried out for p120, VE-cadherin, N-cadherin, and β -actin as loading control. Densitometric values (right) for cadherins are normalized to β -actin and are expressed relative to GFP infection (mean \pm

SEM, n=5; for VE-cadherin $P=0.0132$). * denotes significance ($p<0.05$) as determined using single sample t-test. (C) Similar proportions of VE-cadherin and N-cadherin associate with p120 under control conditions. Lysates of BPAEC were subject to two consecutive cycles of immunoprecipitation for p120. Left: immunoblot of each p120 immunoprecipitate for VE-cadherin (left lanes) and N-cadherin (right lanes). Center: samples taken from lysate before the first immunoprecipitation (Pre-IP) and from supernatants following each immunoprecipitation (Post-IP 1 & 2) were immunoblotted for VE-cadherin, N-cadherin, p120, and GAPDH. Right: graph represents percent of total VE-cadherin and N-cadherin precipitated from lysate in each p120 immunoprecipitation step, as analyzed by densitometry. Densitometric values for each cadherin were normalized to GAPDH to account for any dilution of the lysate and are displayed as percent of total (mean \pm SEM, n=4). (D) BPAEC monolayers were infected with adenovirus containing either GFP (AdGFP), the extracellular and transmembrane portions of the IL-2 receptor fused to the cytoplasmic domain of VE-cadherin with a Myc-epitope tag (AdIL-2R-VE-cad_{cyto}), or the extracellular and transmembrane portions of the IL-2 receptor fused to a p120-uncoupled cytoplasmic domain of VE-cadherin with a Myc-epitope tag (AdIL-2R-VE-cad_{AAA}). 48 hours after infection, cells were lysed and immunoblot analysis was carried out for N-cadherin, the Myc-epitope tag to detect the fusion proteins, and β -actin as a loading control. Blots representative of 4 separate experiments.

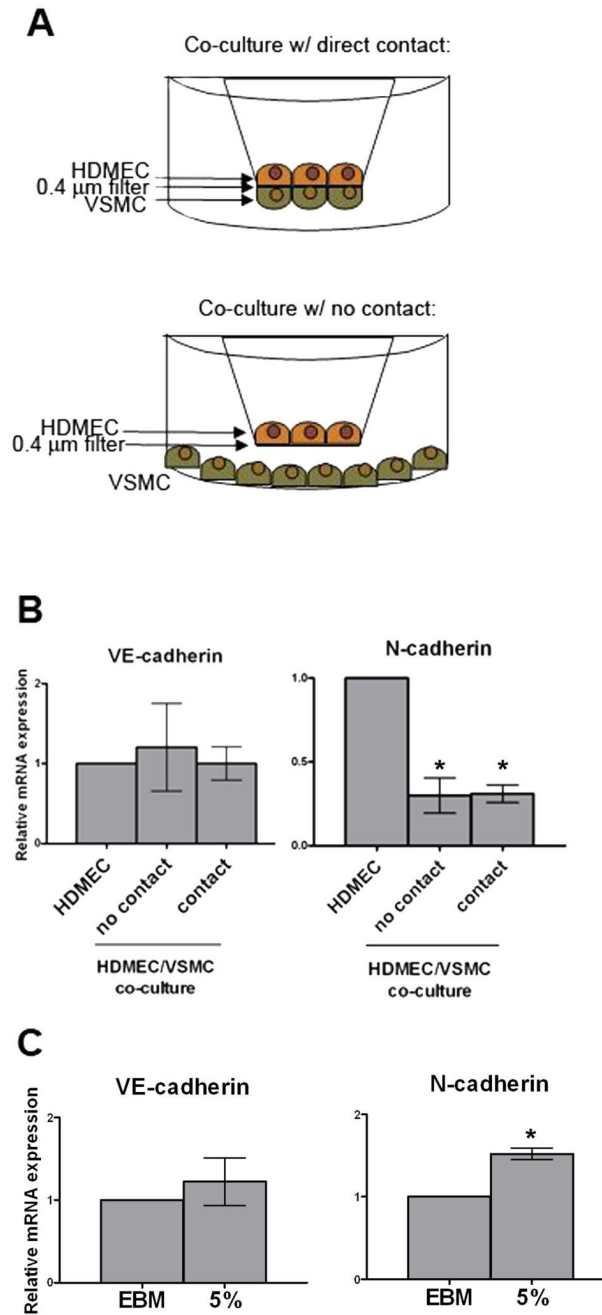


Figure 6. Modulation of N-cadherin mRNA expression by mural cells, serum treatment
 (A) Co-culture systems (8): Top – direct contact. Rat vascular smooth muscle cells (VSMC) were seeded on the bottom surface of a 0.4µm pore Transwell cell culture insert. 24 hours later, HDMEC were seeded on the top side of the wells, and were allowed to grow for 96 hours. Bottom – no contact. Rat vascular smooth muscle cells (VSMC) were seeded on 100mm plate. 24 hours later, HDMEC were seeded on a 0.4µm pore Transwell cell culture insert which was placed in the 100mm plate, and were allowed to grow for 96 hours. HDMEC (not shown) – cells were seeded on a 0.4µm pore Transwell cell culture insert which was placed in an empty 100mm plate and were allowed to grow for 96 hours. (B) RNA was isolated from HDMEC. Graphs show VE-cadherin and N-cadherin mRNA expression. Values are normalized to

GAPDH, and are expressed relative to HDMEC (mean \pm SEM, n=4; * denotes significance ($p < 0.05$) as determined using single sample t-test; for N-cad no contact $P = 0.0068$, N-cad contact $P = 0.0010$). (C) HDMEC were seeded at confluence in growth medium. After 72 hours media was changed to either basal medium (EBM) or basal medium supplemented with 5% FBS (5%). After 24 hours RNA was isolated. Graphs show VE-cadherin and N-cadherin mRNA expression. Values are normalized to GAPDH and are expressed relative to EBM treated cells (mean \pm SEM, n=3; * denotes significance ($p < 0.05$) as determined using single sample t-test; for N-cad $P = 0.0182$).