

Abi-interactor-1 (Abi1) has a role in cardiovascular and placental development and is a binding partner of the $\alpha 4$ integrin

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Dynamic signals linking the actin cytoskeleton and cell adhesion receptors are essential for morphogenesis during development and normal tissue homeostasis. Abi1 is a central regulator of actin polymerization through interactions with multiple protein complexes. However, the in vivo role of Abi1 remains to be defined. The $\alpha 4$ integrin adhesion receptor is associated with enhanced protrusive activity and regulation of directional cell migration. Among integrin subunits, $\alpha 4$ exhibits unique properties in that it predominantly accumulates at the leading edge of migrating cells; however, the pathways that link the actin-regulatory machinery to $\alpha 4$ at the leading edge have remained elusive. We generated Abi1 KO mice and found that loss of Abi1 phenocopies KO of $\alpha 4$. Mice lacking Abi1 or $\alpha 4$ exhibit midgestational lethality with abnormalities in placental and cardiovascular development. Notably, purified Abi1 protein binds directly to the $\alpha 4$ cytoplasmic tail and endogenous Abi1 colocalizes with phosphorylated $\alpha 4$ at the leading edge of spreading cells. Moreover, Abi1-deficient cells expressing $\alpha 4$ have impaired cell spreading, which is rescued by WT Abi1 but not an Abi1 mutant lacking the $\alpha 4$ -binding site. These data reveal a direct link between the $\alpha 4$ integrin and actin polymerization and uncover a role for Abi1 in the regulation of morphogenesis in vivo. The Abi1- $\alpha 4$ interaction establishes a mechanistic paradigm for signaling between adhesion events and enhanced actin polymerization at the earliest stages of protrusion.

WAVE complex | lamellipodia | chorio-allantoic fusion

Intercellular and cell-matrix interactions are essential for many diverse biological processes such as morphogenesis, immune responses, and cancer metastasis. These interactions are dependent on the dynamic regulation of the actin cytoskeleton downstream of adhesion receptors. Prominent among these receptors are the heterodimeric β integrin transmembrane receptors that bind to ECM proteins as well as a subset of cell adhesion molecules. Ligand-bound integrins promote not only adhesion to matrix or adjacent cells, but also initiate downstream signaling pathways through the recruitment of adaptor proteins to the integrin cytoplasmic domains (1). Bidirectional signals linking the actin cytoskeleton to integrin complexes provide cells with spatial cues to initiate directional cell migration, promote adhesion, and generate polarity. The protein complexes required for transducing integrin engagement to the regulation of actin cytoskeletal dynamics are beginning to be elucidated. Although numerous binding partners of the integrin β -subunit cytoplasmic tails have been identified, few interacting proteins for the integrin α -subunits have been uncovered (1). Among these is paxillin, a focal adhesion adaptor protein that binds directly to the $\alpha 4$ integrin cytoplasmic tail (2). This interaction is regulated by phosphorylation of $\alpha 4$ at S988 (3), as signals that initiate protrusive activity promote phosphorylation of $\alpha 4$ at the leading edge, resulting in dissociation of paxillin, thereby allowing for directional migration (4). Thus, paxillin binding to $\alpha 4$ negatively regulates protrusive activity. The $\alpha 4$ integrin exhibits unique

properties among integrin subunits in that $\alpha 4$ predominantly accumulates at the leading edge of migrating cells, rather than at focal adhesions. Moreover, $\alpha 4$ expression is associated with protrusive activity and enhanced cell migration; however, the pathways that link $\alpha 4$ integrin to the actin-regulatory machinery at the leading edge have remained elusive. Here we identify Abi1 as a target of $\alpha 4$ integrin that positively regulates membrane protrusion by promoting actin polymerization at sites of integrin engagement.

The Abi family proteins, Abi1 and Abi2, were originally identified as binding partners and substrates of the Abl tyrosine kinase (5, 6). Subsequent in vitro studies have implicated Abi proteins in the regulation of cell motility, intercellular adhesion, endocytosis, cytokinesis, Golgi architecture, and transport carrier biogenesis, through their ability to interact with distinct protein complexes involved in the regulation of actin nucleation (7–14). Abi proteins localize to sites of actin polymerization at cell-cell contacts in epithelial adherens junctions (8, 9) and immune synapses (15). Abi proteins also localize to the leading edge of lamellipodia, and this localization correlates with positive regulation of directional migration (12, 16, 17). Regulation of protrusive activity by Abi proteins has been linked to the formation of protein complexes with the WASP/WAVE family of nucleation-promoting factors (18), leading to activation of the Arp2/3 complex (19). Abi1 has also been shown to interact with Diaphanous formins to promote actin polymerization-driven protrusions at the leading edge of motile cells and at sites of cell-cell contact (9, 10).

Genetic studies have shown that Abi family proteins are required for the regulation of cytoskeletal dynamics in vivo and morphogenetic abnormalities are observed in flies and mice lacking Abi proteins. We showed that homozygous deletion of mouse Abi2 results in abnormal phenotypes in the eye and brain—tissues with high Abi2 expression (8). *Drosophila* embryos lacking both maternal and zygotic *abi* exhibit embryonic lethality and collapse of axonal patterning (20). In contrast to Abi2, Abi1 exhibits widespread expression (8). To investigate Abi1 function during morphogenetic events in vivo, we disrupted the *Abi1* gene in mice. Abi1-deficient mice exhibit defects in placental and cardiovascular development, leading to midgestational embryonic lethality. These phenotypes mirror those of mice deficient for the $\alpha 4$ integrin and its counter receptor VCAM1 (21–24). Integrin $\alpha 4\beta 1$ binding to VCAM1 is essential not only for placental and

Author contributions: C.R., M.H.G., and A.M.P. designed research; C.R. and J.H. performed research; M.H.G. and J.H. contributed new reagents/analytic tools; C.R., M.H.G., and A.M.P. analyzed data; and C.R. and A.M.P. wrote the paper.

The authors declare no conflict of interest.

*This Direct Submission article had a prearranged editor.

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This article contains supporting information online at www.pnas.org/lookup/suppl/doi:10.1073/pnas.1012316108/-DCSupplemental.

cardiac development but also for extravasation of circulating lymphocytes into inflamed tissue by binding of $\alpha 4\beta 1$ -expressing lymphocytes to VCAM1 expressed on activated endothelia (25). Binding of $\alpha 4\beta 1$ to VCAM1 also regulates tumor angiogenesis and homing of hematopoietic stem and progenitor cells (26, 27). Thus, blocking the interactions of $\alpha 4$ with downstream targets may be an effective therapy for the treatment of autoimmune diseases, pathological angiogenesis, and other disorders (25). Our findings reveal that *Abi1* interacts with the $\alpha 4$ cytoplasmic tail and may mediate specific functions associated with $\alpha 4$ -dependent processes in normal and pathological conditions.

Results

Role for *Abi1* in Chorio-Allantoic Fusion. To examine the role of the *Abi1* adaptor molecule in vivo, as well as to identify *Abi1*-dependent signaling pathways, we used homologous recombination to generate *Abi1*-deficient mice (Fig. 1 and Fig. S1). Loss of *Abi1* resulted in embryonic lethality between embryonic day (E) 8.5 and E10.5, depending on the genetic background (Fig. S2). Mouse embryonic fibroblast (MEF) cells derived from these embryos showed absence of *Abi1* protein expression, indicating that these embryos carry a null mutation (Fig. S3). At E7.5 to E8.0, there was no significant difference in appearance between WT and *Abi1*^{-/-} embryos; however, as development proceeded, *Abi1*^{-/-} embryos showed developmental delays and growth retardation. The onset of phenotypic abnormalities occurred at approximately E8.5 for *Abi1* KO mice in the C57BL/6 strain, and E9.5 for *Abi1* loss in the CD1 strain. Notably, *Abi1*^{-/-} mice bred on the C57BL/6 background showed failure of chorio-allantoic fusion (61 of 111; 55%), which was not associated with growth retardation (Fig. 1). The allantoises of these *Abi1*^{-/-} embryos extended through the exocoelomic cavity, achieving a size com-

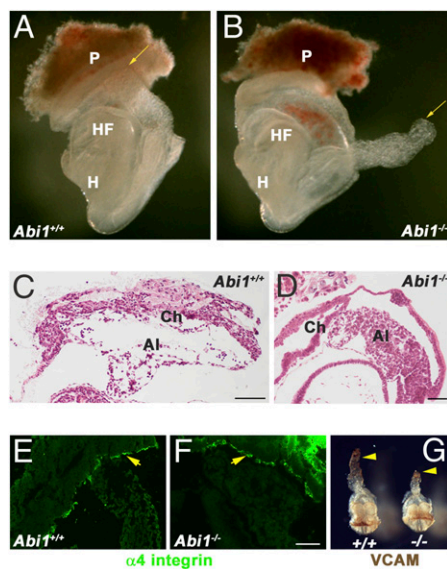


Fig. 1. Defective chorio-allantoic fusion of *Abi1* mutant embryos on the C57BL/6 background. (A) WT embryo at E8.5, dissected free from the uterus and attached to a portion of the maternal placenta (P), shows the allantois (arrow) attached to the chorion, which fuses at this stage. (B) *Abi1* homozygous mutant embryo at E8.5 is shown at identical magnification to WT embryo in A. The distal tip of the allantois (arrow) floats freely, unattached to the chorion. (C and D) H&E sections of WT (C) and *Abi1*-null embryos (D) illustrate the lack of attachment of the allantois to the chorion in *Abi1* mutants, although the allantois length is sufficient to allow contact. $\alpha 4$ integrin (E and F, arrows) and VCAM (G, arrowheads) are expressed appropriately in the chorion and allantois, respectively, of *Abi1* mutant embryos. HF, headfold; H, heart; Ch, chorion. (Scale bars: 100 μm .)

parable to that of WT littermates, but failed to initiate fusion (Fig. 1 A–D). The WT allantois undergoes de novo vasculogenesis as evidenced by the large hydropic spaces indicative of endothelial cell remodeling events occurring at the distal tip of the allantois before chorio-allantoic fusion (28) (Fig. 1C). In contrast, the allantoises of *Abi1*^{-/-} embryos exhibited impaired vasculogenesis, appearing compact and lacking spaces at the distal tip (Fig. 1D). Chorio-allantoic fusion requires reciprocal interactions between $\alpha 4$ integrin and VCAM1 expressed on the surface of the chorion and allantois, respectively (21–23). Thus, we analyzed the expression of these proteins in *Abi1*^{-/-} embryos at E8.5. We found that both $\alpha 4$ integrin and VCAM1 were expressed appropriately in these mutant embryos, indicating that loss of expression of either protein was not responsible for the observed lack of chorio-allantoic fusion (Fig. 1 E–G). These data implicate *Abi1* in chorio-allantoic fusion, and suggest that placental failure is a contributing factor to the lethality of *Abi1*-deficient mice.

Placental Abnormalities Are Associated with Chorionic Defects in *Abi1*^{-/-} Embryos. In contrast to the chorio-allantoic fusion abnormalities observed in the C57BL/6 background, *Abi1* KO mice on the CD1 strain were able to initiate chorio-allantoic fusion, but displayed dramatic defects in chorionic development by E9.5 which were often accompanied by growth retardation (Fig. 2). In these *Abi1*^{-/-} mice, the chorion appears compact and button-like, often with a prominent blood clot at the maternal interface (Fig. 2 B and C). In the absence of *Abi1* the chorion appears densely packed, whereas at this stage the WT chorion has spread circumferentially throughout the surface of the ectoplacental cone (Fig. 2 D–G). Ultrastructurally, a thickened basement membrane associated with Reichart's membrane is apparent in *Abi1*^{-/-} mice, suggestive of defects in integrin activation (Fig. 2 H–J) (29). Comparison of the phenotypes of *Abi1*^{-/-} mice on the C57BL/6 and CD1 strains suggest that defects arising in chorionic trophoblast cells may underlie the aberrant placentation observed in both of these strains.

Loss of *Abi1* Affects Cardiovascular Development. In addition to the placental and allantoic vasculogenic defects in the *Abi1*^{-/-} mice, loss of *Abi1* resulted in cardiovascular abnormalities (Fig. 3). *Abi1*^{-/-} mice on the CD1 strain displayed reduced angiogenesis in the yolk sac, edema and hemorrhage around the heart, as well as disruption of the heart layers (Fig. 3 A–G). The epicardium of *Abi1*^{-/-} mice appears disorganized (Fig. 3G), a phenotype reminiscent of the $\alpha 4$ integrin KO mice (21). In the absence of *Abi1*, the myocardium is also disorganized, and the endocardium is discontinuous (Fig. 3G). These data demonstrate a requirement for *Abi1* in cardiovascular development and suggest that cardiovascular defects in the *Abi1*^{-/-} mice may contribute to their embryonic lethality.

***Abi1*^{-/-} MEF Cells Display Cell Spreading Defects Downstream of $\alpha 4$ Integrin.** The phenotypes of the *Abi1*-null mice suggested that *Abi1* may play a role in $\alpha 4$ integrin-mediated morphogenetic events, and may regulate $\alpha 4$ -mediated signaling (21). To examine this possibility, we derived control (*Abi1*^{+/-}/*Abi2*^{+/-}) and *Abi1* mutant (*Abi1*^{-/-}/*Abi2*^{+/-}) MEFs, a cell type that does not endogenously express the $\alpha 4$ integrin, and engineered them to exogenously express $\alpha 4$ integrin. Mutant MEFs were derived from embryos that were null for *Abi1*, and additionally heterozygous for *Abi2*, as we found up-regulation of *Abi2* expression in MEFs that were null for *Abi1* on an *Abi2* WT background (Fig. S3B). Control MEFs were derived from a littermate that was heterozygous for both *Abi1* and *Abi2*. Control and mutant MEFs were assayed for adhesion-dependent cell spreading on VCAM1, a ligand that specifically interacts with $\alpha 4$ integrin. As expected, in the absence of $\alpha 4$ expression both control and mutant cells failed

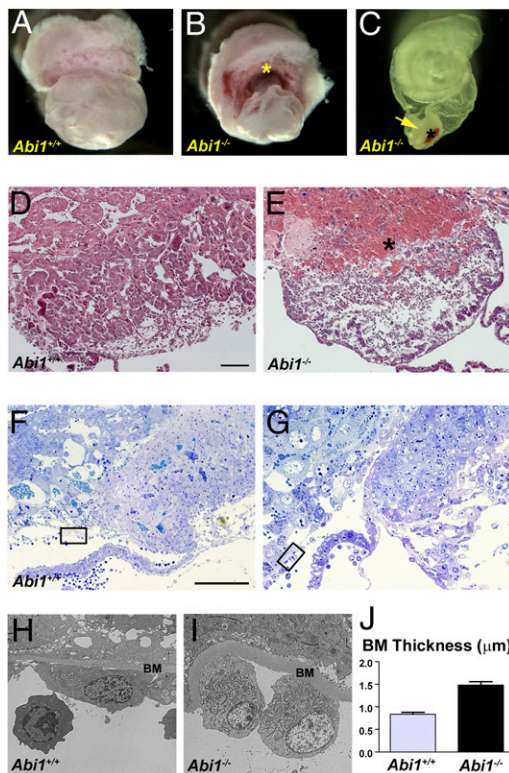


Fig. 2. Placental defects in *Abi1*-null mice on the CD1 genetic background. (A) WT embryo surrounded by yolk sac with placenta. (B) *Abi1* mutant embryo, also surrounded by yolk sac with hemorrhagic placenta (asterisk). (C) *Abi1* mutant embryo, with a portion of the yolk sac, and attached "chorionic button" (arrow) and blood clot at the maternal surface (asterisk). (D and E) H&E sections of chorion and labyrinth regions of the placentas of WT (D) and *Abi1* mutant (E) embryos. Note the blood clot adjacent to the maternal placenta in *Abi1*^{-/-} (asterisk). (F and G) Thin sections of the chorion and labyrinth regions of the placentas of WT (F) and *Abi1* mutant (G) embryos stained with toluidine blue to show cellular detail and provide orientation for ultrathin section transmission EM analysis. Boxed areas correspond to regions shown in H and I, respectively. (H and I) Transmission EM micrographs of ultrathin sections of WT (H) and *Abi1* mutant (I) embryos. Note the thickened basement membrane (BM) of Reichart membrane that surrounds the yolk sac of the embryo. (J) Quantification of basement membrane thickness in WT and *Abi1* mutant embryos ($P < 0.0001$). (Scale bars: 100 μm.)

to spread on VCAM1 (Fig. 4). Expression of $\alpha 4$ integrin in control cells allowed for significantly enhanced spreading on VCAM1 (Fig. 4). In contrast, *Abi1* mutant cells expressing equivalent cell surface levels of $\alpha 4$ integrin were severely impaired in their ability to spread on VCAM1 (Fig. 4). These data implicate *Abi1* in $\alpha 4$ integrin-mediated cellular behaviors, but do not address the specificity of this interaction. We tested the specificity for *Abi1* in $\alpha 4$ -mediated cell spreading by analyzing the spreading of control and mutant MEFs on a fibronectin substratum (Fig. S4). MEFs endogenously express the $\alpha 5\beta 1$ integrin heterodimer, which similar to other integrins, including $\alpha 4\beta 1$, interacts with the ECM protein fibronectin (30). *Abi1* mutant cells exhibit only a modest decrease in cell spreading on fibronectin compared with their control counterparts, indicating *Abi1* is not required for engagement downstream of the classical fibronectin receptor, $\alpha 5\beta 1$ (Fig. S4). However, when engineered to express the $\alpha 4$ integrin, *Abi1* mutant cells demonstrate a significant reduction in cell spreading on fibronectin compared with their control counterparts (Fig. S4). Together, these data support a role for *Abi1* in $\alpha 4$ integrin-mediated signaling in vitro, and suggest that aberrant $\alpha 4$ signaling may underlie the embryonic lethality of *Abi1*^{-/-} mice.

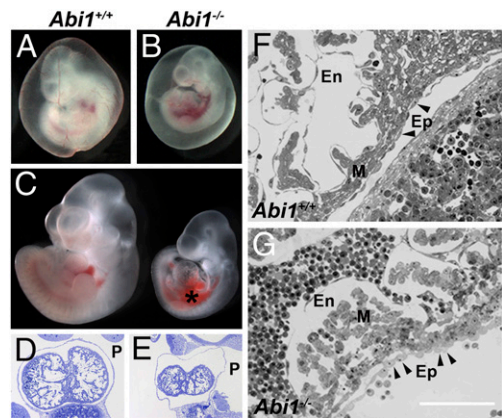


Fig. 3. Loss of *Abi1* results in cardiovascular abnormalities. (A) WT embryo with surrounding vascularized yolk sac at E10.5. (B) *Abi1*^{-/-} littermate shows defective vascularization in the yolk sac. Embryos in A and B were photographed at the same magnification. (C) WT (Left) and *Abi1*^{-/-} (Right) embryos with yolk sac removed were photographed together; the *Abi1*^{-/-} embryo displays edema and hemorrhage around the heart. (D–G) In thin sections, the *Abi1*^{-/-} heart (E) is smaller in size with a swollen pericardium and associated edema, whereas the WT heart (D) fills the pericardial cavity. At higher magnification, the epicardial (Ep), myocardial (M), and endocardial (En) layers of the heart are disorganized in *Abi1*^{-/-} embryos (G) in contrast to WT embryo (F). Of note is the ruffled appearance of the epicardial cells (arrowheads in F and G) adjacent to the myocardium in *Abi1* mutants. This phenotype is reminiscent of that of the $\alpha 4$ integrin KO mouse. (Scale bar: 100 μm.)

Abi1 Interacts with the $\alpha 4$ Tail. Paxillin was identified as a major binding partner of the $\alpha 4$ integrin, an interaction thought to regulate Rac activation and protrusive activity (2, 3). Phosphorylation of $\alpha 4$ at the leading edge of migrating cells disrupts the inhibitory adhesive interaction with paxillin, thereby allowing for directional migration (4, 31, 32). However, this model does not provide a mechanistic explanation for how phosphorylated $\alpha 4$ promotes actin-dependent lamellipodial protrusions at the leading edge. Thus, we asked whether *Abi1* might function to directly link $\alpha 4$ to positive regulation of actin dynamics at the leading edge of lamellipodial protrusions and analyzed whether these proteins could interact physically and spatially. We found that *Abi1* bound to purified $\alpha 4$ cytoplasmic tail (Fig. 5A). *Abi1* binding to the $\alpha 4$ tail was selective, as no binding was detected between *Abi1* and $\alpha 1\text{b}$, a distinct α subunit (Fig. 5A).

Abi1 is a multisubunit adaptor protein with distinct protein-protein interacting domains, including a proline-rich region and SH3 domain (Fig. 5B) (16). We found that the *Abi1* N-terminal amino acids (1–145) were required for $\alpha 4$ binding (Fig. 5C). In contrast, the C-terminal region, containing both the proline-rich region and SH3 domain, was dispensable for interaction with $\alpha 4$, as a deletion mutant of *Abi1* containing only the first 111 amino acids bound robustly to $\alpha 4$ (Fig. 5C). Moreover, by using purified GST-*Abi1* full length or GST-*Abi1*-1-111 proteins, we showed that the interaction of *Abi1* with $\alpha 4$ is direct (Fig. 5D). Thus, the *Abi1* N-terminal sequences are both necessary and sufficient for binding to $\alpha 4$.

The N-Terminal Region of *Abi1* Is Required for Rescue of $\alpha 4$ -Mediated Cell Spreading on VCAM1. Biochemical analysis demonstrated that the N-terminal region of *Abi1* was required for binding to the cytoplasmic tail of the $\alpha 4$ integrin; thus, we wished to assess the role of this region of *Abi1* in $\alpha 4$ -mediated cell spreading. For these experiments, $\alpha 4$ -integrin expressing control MEFs (*Abi1*^{+/-}/*Abi2*^{+/-}) reconstituted with GFP vector or $\alpha 4$ -expressing mutant MEFs (*Abi1*^{-/-}/*Abi2*^{+/-}) reconstituted with GFP vector, WT *Abi1*-GFP, or the $\Delta 1$ –145 *Abi1*-GFP were plated on VCAM1 (Fig. 6). We found

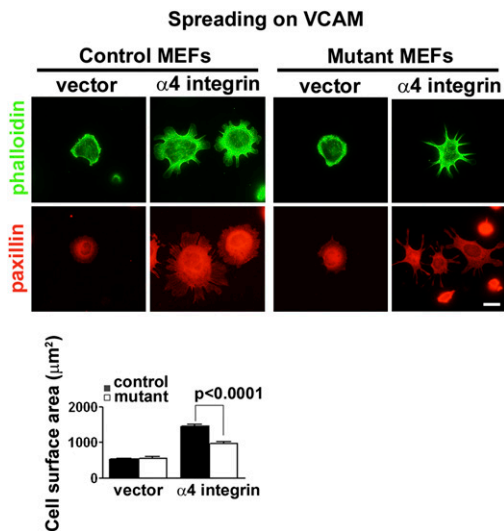


Fig. 4. *Abi1* mutant MEFs expressing $\alpha 4$ integrin show diminished cell spreading on VCAM1. Control (*Abi1*^{+/+}/*Abi2*^{+/+}) or mutant (*Abi1*^{-/-}/*Abi2*^{+/+}) MEFs were transduced with vector alone or vector containing WT full-length $\alpha 4$ integrin. Cells were sorted by FACS to obtain cell populations expressing equivalent amounts of $\alpha 4$, and these cells were plated immediately on coverslips coated with VCAM1. After 2 h, cells were fixed and processed for indirect immunofluorescence for detection of paxillin (red). Actin staining with phalloidin is shown (green). Quantification of cell spreading was analyzed by measuring cell surface area of phalloidin stained cells using Zeiss software (*Bottom*). *Abi1* mutant cells expressing $\alpha 4$ integrin exhibit significantly reduced spreading compared with their control counterparts on VCAM1 substratum. (Scale bar: 10 μm .)

that defective spreading of the $\alpha 4$ -expressing *Abi1*-deficient (mutant) cells was rescued by expression of exogenous WT GFP-tagged *Abi1*, but not the N-terminal deletion mutant, $\Delta 1$ -145, which lacks

the ability to bind to the $\alpha 4$ integrin (Fig. 6). These data demonstrate that the N terminus of *Abi1* couples $\alpha 4$ -integrin receptor engagement to actin cytoskeleton-mediated cell spreading.

Endogenous *Abi1* and Phosphorylated $\alpha 4$ Colocalize at the Leading Edge. To test whether endogenous $\alpha 4$ and *Abi1* could interact, we used A7r5 cells, a rat smooth muscle cell line that expresses $\alpha 4$ as well as *Abi1*, and responds to scratch wound stimuli by accumulation of S988-phosphorylated $\alpha 4$ at the leading edge (33). Indeed, endogenous *Abi1* coimmunoprecipitated with endogenous $\alpha 4$ integrin, and these immunoprecipitates contain S988-phosphorylated $\alpha 4$ (Fig. 7*A*). In cells plated on VCAM1, *Abi1* and S988-phosphorylated $\alpha 4$ colocalized at the leading edge as detected by confocal microscopy (Fig. 7*B*). Thus, *Abi1* binding to the $\alpha 4$ integrin at the leading edge of migrating cells or in polarized epithelium provides a direct link to the regulation of actin dynamics during tissue morphogenesis.

Discussion

Analysis of *Abi1* mutant mice uncovered an unexpected mechanistic link between *Abi1*-regulated cytoskeletal dynamics and $\alpha 4$ integrin-dependent morphogenetic events. Mice knockout for $\alpha 4$ or *Abi1* exhibit defective chorio-allantoic fusion and cardiovascular defects. Chorio-allantoic fusion is regulated by the binding of $\alpha 4$ integrin on the surface of the chorion to VCAM1 on the surface of the allantois (21–23). The defects in placental development observed in the absence of *Abi1* suggest that loss of *Abi1* in the chorionic mesothelium impairs $\alpha 4$ -mediated signaling, ultimately leading to chorionic abnormalities and placental failure in the C57BL/6 or CD1 genetic background. *Abi1*-null mice, similar to $\alpha 4$ integrin and VCAM1 KO mice, display defects in cardiovascular development with impaired vessel maturation in the yolk sac, defective adhesion of epicardial cells to the adjacent myocardium, and general edema around the heart. Interestingly, mice knocked out for the *Abi*-binding protein WAVE2 also exhibit cardiovascular abnormalities and die

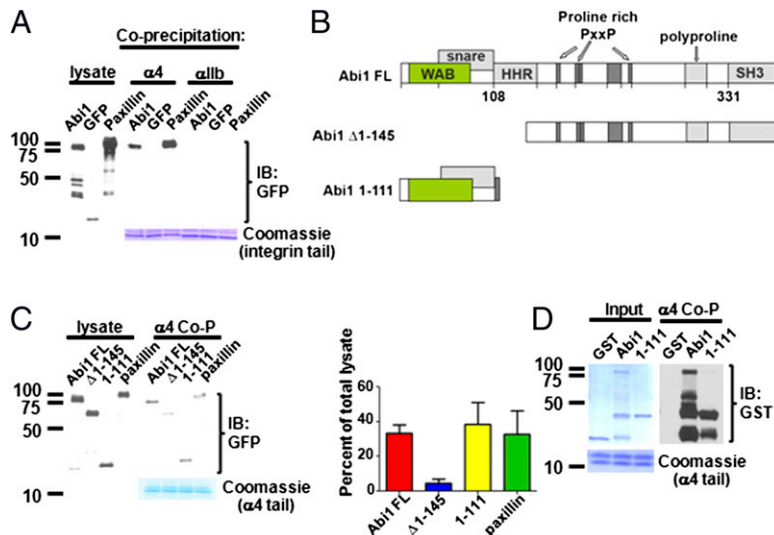


Fig. 5. *Abi1* protein binds to $\alpha 4$ integrin. (*A*) Bacterially expressed $\alpha 4$ or $\alpha 1b$ integrin cytoplasmic tails were incubated with lysates from 293T cells expressing GFP-*Abi1*, GFP-paxillin, or GFP alone. Detection of integrin tail-bound proteins by Western blotting for GFP showed that *Abi1* bound to $\alpha 4$ but not to $\alpha 1b$ (*Top*). Coomassie staining showed equal loading of integrin cytoplasmic tails in the pull-down assays (*Bottom*). (*B*) Structural diagram of *Abi1* full-length (FL) and indicated fragments. *Abi1* contains proline-rich and SH3 domains in its C terminus, and the WAVE binding (WAB), SNARE, and homeodomain homology region (HHR) in its N terminus. (*C*) Identification of $\alpha 4$ -binding site on *Abi1* was carried out as described in *A*. Deletion of the N-terminal region of *Abi1* decreases binding to the $\alpha 4$ cytoplasmic tail, whereas the N-terminal *Abi1* (1–111) fragment is sufficient for binding to the $\alpha 4$ tail. (*D*) Direct binding of purified $\alpha 4$ tail to purified GST-*Abi1* full-length and GST-*Abi1*-1-111 N-terminal fragment, but not to GST alone. Coomassie staining of GST fusion proteins (*Upper*) and $\alpha 4$ integrin cytoplasmic tails is shown (*Bottom*). Protein molecular weight markers are indicated (*A*, *C*, and *D*, *Left*). Co-P, coprecipitation.

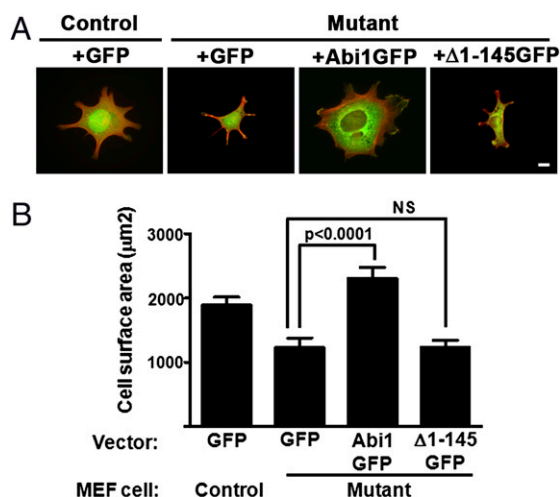


Fig. 6. The N-terminal region of Abi1 is required for rescue of $\alpha 4$ -mediated cell spreading on VCAM1. The $\alpha 4$ -expressing control ($Abi1^{+/+}/Abi2^{+/+}$) MEFs were reconstituted with GFP vector and $\alpha 4$ -expressing mutant ($Abi1^{-/-}/Abi2^{+/+}$) MEFs were reconstituted with GFP vector, GFP-tagged WT Abi1 or the GFP-tagged $\Delta 1$ -145 Abi1 deletion mutant. Cells were sorted by FACS to obtain cell populations expressing equivalent amounts of $\alpha 4$ as well as GFP, and these cells were plated immediately on coverslips coated with VCAM1. (A) Representative images of each cell type are shown. Merged images reveal GFP or the GFP fusion proteins (green) with phalloidin-stained actin (red). (B) Quantification of cell spreading was analyzed by measuring cell surface area of phalloidin stained cells using Zeiss software. Abi1 mutant cells expressing $\alpha 4$ integrin and GFP displayed significantly reduced spreading compared with their control counterparts. This defect was rescued by adding back GFP-tagged WT Abi1 (compare GFP with GFP-Abi1; $P < 0.0001$, two-tailed Student t test), whereas GFP-Abi1 $\Delta 1$ -145 was unable to rescue [GFP vs. GFP- $\Delta 1$ -145; not significant (NS)]. (Scale bar: 10 μm .)

midgestationally (24). WAVE2-deficient mice exhibit impaired vascular remodeling in response to growth factor stimulation (24). Further, mouse mutants of Nap1, a direct binding partner of Abi1 in the WAVE complex, also exhibit defective chorio-allantoic fusion, and other developmental phenotypes associated with defects in cell migration and adhesion (34). Interestingly, heterozygous insertional mutants of Nap1, which display neural tube closure defects, also have dramatic reductions in vessel diameter (35). Thus, the Abi1-WAVE2-Nap1 complex may regulate a subset of signals downstream of integrin receptors during cardiovascular development.

In contrast to other integrin subunits, $\alpha 4$ integrin signaling occurs primarily in the cell periphery, rather than in focal adhesions or focal complexes (4, 33, 36). Indeed, expression of $\alpha 4$ in CHO cells promotes protrusion of broad lamellipodia in response to scratch wounding (36). Furthermore, phosphorylated $\alpha 4$ integrin preferentially localizes to the leading edge of migrating cells and this localization is linked to positive regulation of lamellipodial extensions and enhanced cell migration (37, 38). Unexpectedly, we found that Abi1 interacts with the purified $\alpha 4$ cytoplasmic tail and colocalizes with the endogenous phosphorylated $\alpha 4$ integrin at the leading edge of spreading cells. To date, the model for the regulation of $\alpha 4$ -mediated directional migration hinges on the interaction of $\alpha 4$ with paxillin, which negatively regulates Rac activity, and the termination of this association at the leading edge by phosphorylation of the $\alpha 4$ cytoplasmic tail. Paxillin and $\alpha 4$ remain associated at the sides and rear of the cell, where paxillin recruits the ADP ribosylation factor GTPase activating protein (Arf-GAP), leading to the inhibition of Rac (32). Directional cell migration was thereby proposed to occur by relieving an inhibitory interaction at the leading edge of the cell; however, the signals

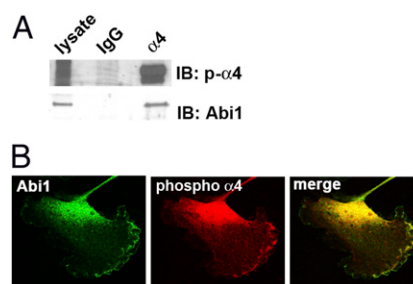


Fig. 7. Endogenous Abi1 interacts with endogenous $\alpha 4$ integrin. (A) Lysates from A7r5 cells were subjected to immunoprecipitation with the HP2/1 antibody specific for $\alpha 4$ integrin. Coprecipitation of endogenous Abi1 with $\alpha 4$ integrin containing 5988-phosphorylated $\alpha 4$ integrin was detected by using a phospho-specific antibody. Specificity was shown by incubation of lysates with isotype IgG control. (B) Indirect immunofluorescence of endogenous Abi1 and 5988-phosphorylated $\alpha 4$ integrin proteins revealed colocalization at the leading edge of cells spread on VCAM1 by using confocal microscopy.

that link $\alpha 4$ to enhanced actin polymerization and protrusive activity at the leading edge have remained poorly understood. We have identified Abi1 as a key mediator of $\alpha 4$ -mediated signaling at the leading edge. Abi1 interacts and colocalizes in discrete puncta at the leading edge of lamellipodia with phosphorylated $\alpha 4$. Moreover, Abi1 interacts with the purified cytoplasmic tail of $\alpha 4$ in vitro and KO of *Abi1* dramatically diminishes cell spreading of fibroblasts engineered to express $\alpha 4$ on both VCAM1 and fibronectin.

The $\alpha 4\beta 1$ integrin heterodimer plays an important role in diseases of the cardiovascular and immune systems (39). Antagonists of $\alpha 4$ have shown promise in the treatment of autoimmune disorders such as multiple sclerosis and Crohn's disease, but the associated toxicities have led to the design of alternative approaches other than those directed at blocking integrin-ligand binding (40, 41). Blocking $\alpha 4$ signaling may have important implications with respect to ablation of tumor vasculature, as it has been recently demonstrated that homing of progenitor cells to tumor areas undergoing active angiogenesis uses the $\alpha 4$ integrin/VCAM1 counter receptor pair (42, 43). Moreover, exciting new findings have shown that suppression of $\alpha 4\beta 1$ function in the endothelium inhibits tumor lymphangiogenesis and metastasis in mouse models (44). Current approaches to antagonize $\alpha 4$ function have focused on development of drugs that disrupt the interaction of $\alpha 4$ with paxillin (41). Our data suggest that development of inhibitors of the Abi1- $\alpha 4$ interaction may be an alternative therapeutic strategy for the treatment of metastatic tumors, inflammatory immune disorders and other $\alpha 4$ -dependent pathologic processes.

Materials and Methods

Generation of *Abi1*^{-/-} Mice. The mouse *Abi1* locus was targeted by homologous recombination using standard methods and confirmed by Southern blotting and PCR analysis as described in Fig. S1. Additional experimental details are provided in SI Materials and Methods. Included in this information are reagents and protocols with respect to coprecipitation and in vitro binding assays, cell culture, cell spreading assays and immunofluorescence microscopy as well as statistical analysis performed.

ACKNOWLEDGMENTS. The authors thank Cheryl Bock of the Duke Transgenic and Knockout Mouse Facility for generation of Abi1 KO mice and Matthew Grove for preparation of the targeting vector and identification of positive embryonic stem cell clones. We also thank Sam Johnson of the Duke Light Microscopy Facility for excellent advice, Rob Wechsler-Reya for use of the stereoscope, Phillip Christopher of Duke Electron Microscopy Services, Donna Webb for insights on evaluation of focal adhesion dynamics, and James Cross for insights into the placental phenotypes. This study was supported by National Institutes of Health Grants CA070940 (to A.M.P.), HL084102 (to A.M.P.), and AR27214 (to M.H.G.).

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