The Caenorhabditis elegans vab-10 spectraplakin isoforms protect the epidermis against internal and external forces

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Morphogenesis of the Caenorhabditis elegans embryo is driven by actin microfilaments in the epidermis and by sarcomeres in body wall muscles. Both tissues are mechanically coupled, most likely through specialized attachment structures called fibrous organelles (FOs) that connect muscles to the cuticle across the epidermis. Here, we report the identification of new mutations in a gene known as vab-10, which lead to severe morphogenesis defects, and show that vab-10 corresponds to the C. elegans spectraplakin locus. Our analysis of vab-10 reveals novel insights into the role of this plakin subfamily. vab-10 generates isoforms related either to plectin (termed VAB-10A) or to microtubule actin cross-linking factor plakins (termed VAB-10B). Using specific antibodies and mutations, we show that VAB-10A and VAB-10B have distinct distributions and functions in the epidermis. Loss of VAB-10A impairs the integrity of FOs, leading to epidermal detachment from the cuticle and muscles, hence demonstrating that FOs are functionally and molecularly related to hemidesmosomes. We suggest that this isoform protects against forces external to the epidermis. In contrast, lack of VAB-10B leads to increased epidermal thickness during embryonic morphogenesis when epidermal cells change shape. We suggest that this isoform protects cells against tension that builds up within the epidermis.

Introduction

Morphogenesis is the complex set of processes by which a tissue, an organ, or an organism reaches its final form. It involves numerous changes in shape and position of individual cells. These transformations are driven by molecular motors and cell adhesion complexes; they occasionally require a mechanical coupling between adjacent tissues (Schock and Perrimon, 2002).

Motors molding tissues generate forces that can have shearing effects. Thus, it is likely that they are coupled to molecules acting to preserve cell and tissue integrity. The mechanisms that protect cells against mechanical stresses have seldom been investigated in the context of morphogenesis. Their dissection has instead relied primarily on the analysis of human autoimmune or genetic diseases affecting tissue integrity, and on mouse knockout models. Biochemical and genetic studies have outlined the key function played by integrin- and dystrophin-associated adhesion complexes, plaques, and intermediate filaments (IFs;* Fuchs and Cleveland, 2002).

*Abbreviations used in this paper: ABD, actin-binding domain; BPAG, bullous pemphigoid antigen; FO, fibrous organelle; GAR, growth-arrest protein 2-related homology; IF, intermediate filament; MCAF, microtubule actin cross-linking factor; MF, microfilament; MT, microtubule; RNAi, RNA interference.

© The Rockefeller University Press, 0021-9525/2003/05/757/12 $8.00
The Journal of Cell Biology, Volume 161, Number 4, May 26, 2003 757–768
http://www.jcb.org/cgi/doi/10.1083/jcb.200302151
In the skin, for instance, the plectin and bullous pemphigoid antigen (BPAG)-ε plakins were shown to link the keratin filaments to α6β4-integrin located in hemidesmosomes (Nievers et al., 1999). Mutations in any of these proteins cause skin blistering disorders on tension or outer abrasion (Pulkkinen and Uitto, 1999; McGrath and Eady, 2001).

We are analyzing the molecular basis and the mechanical aspects of morphogenesis using a genetic strategy in Caenorhabditis elegans. C. elegans embryos elongate more than fourfold along their anterior–posterior axis in the absence of cell division. This process mainly depends on the epidermis surrounding the embryo and the underlying body wall musculature (Chin-Sang and Chisholm, 2000). The initial phase of elongation, which corresponds to a twofold increase in embryonic length, is driven by the contraction of circumferentially oriented actin microfilaments (MFs) causing changes of epidermal cell shapes (Priess and Hirsh, 1986).

Mutations affecting proteins that anchor MFs (α-actinin, β-catenin, E-cadherin) or that organize the MF bundles and regulate their contractions (rhr kinase, myosin light chain, spectrins) disrupt embryonic elongation (for review see Chin-Sang and Chisholm, 2000). The subsequent phase of elongation requires the activity of muscle cells, which assemble sarcomeres at the plasma membrane facing epidermal cells during the initial phase (Hresko et al., 1994). Genetic analysis has identified most components involved in building a functional sarcomere (Williams and Waterston, 1994; Mackinnon et al., 2002). Mutations in the corresponding genes generally prevent embryonic elongation beyond the twofold stage, resulting in a terminal phenotype termed Pat, for “paralyzed at twofold” (Williams and Waterston, 1994).

The molecular mechanism underlying the mechanical coupling between muscle and epidermal cells is poorly understood. In C. elegans, the external cuticle acts as an exoskeleton onto which muscles attach in order to transform their contractions into body movements. These attachments comprise two distinct entities. Within muscle cells, dense bodies at the level of thin filaments and the M-line at the level of thick filaments anchor sarcomeres to the muscle plasma membrane. They are functionally and molecularly related to vertebrate adhesion plaques, including, for instance, an integrin dimer called PAT-2/PAT-3 (Williams and Waterston, 1994; Gettner et al., 1995; Mackinnon et al., 2002). Within epidermal cells, a series of small electron-dense plaques are found at the basal and apical membranes facing muscles and the cuticle, respectively. These plaques are reminiscent of vertebrate hemidesmosomes because they are connected to IFs (Francis and Waterston, 1985). The entire unit (plaques and IFs) is known as a fibrous organelle (FO). Besides IFs, another potential FO component, based on its distribution, corresponds to the uncharacterized protein recognized by the mAb MH5 that was generated by immunizing mice against muscle- and cuticle-associated extracts (Francis and Waterston, 1985, 1991).

Results
Identification of vab-10 alleles
Previously, we performed a genetic screen to identify loci required for embryonic morphogenesis, and reported that embryos homozygous for the chromosomal deficiency hDf17 arrest with severe morphogenic defects (Labouesse, 1997).

We could identify an embryonic lethal mutation, h1356, that recapitulates some aspects of the hDf17 mutant phenotype (Fig. 1 B), and then found that h1356 is allelic to the viable mutation vab-10(e698). This allele was originally identified for its variable head morphogenesis defects (varibably abnormal), and was later observed to cause muscle attachments to be fragile in larvae (Hodgkin, 1983; Plenfish et al., 2000). In an independent screen for lethal mutants, we identified the vab-10 allele ju281 (Fig. 1 C). Finally, after cloning vab-10, we recovered the small deletion mc44 using a molecular approach (Fig. 1 E; see Materials and methods).

vab-10 is a spectraplakin locus that generates two distinct sets of plakins
We molecularly identified vab-10 by a positional cloning strategy (see Materials and methods). vab-10 is a complex locus that generates two distinct sets of isoforms by alternative splicing of a common 5′ region to two distinct 3′ regions. The locus spans a region previously predicted to contain three genes (ZK1151.1, ZK1151.2, and ZK1151.3; Fig. 2 A). Using RT-PCR, we identified two distinct sets of isoforms that all contained ZK1151.1 exons (dark gray) spliced either to ZK1151.3 (medium gray) or to ZK1151.2 (light
The domains found in the proteins encoded by vab-10, its overall sequence similarity with Drosophila shot, and their comparable intron/exon organization qualify vab-10 as the sole C. elegans spectraplakin locus (Roper et al., 2002). The common vab-10 exons code for two potential actin-binding domains (ABD) and a plakin domain; the putative ABD is >50% identical to that of other actin-binding plakins (Andra et al., 1998; Sun et al., 2001; Lee and Kolodziej, 2002b). vab-10A–specific exons code for plectin repeats that are preceded by a short coiled-coil rod domain. Comparison with the IF-binding domain of rat plectin (Nikolic et al., 1996) identifies a small region (57% similarity) within the only cluster of five plectin repeats at the COOH terminus of VAB-10A (Fig. S2). This motif cannot be found in any Shot isoform, consistent with the absence of IFs in Drosophila. vab-10B–specific exons code for spectrin repeats, two EF-hands, and a growth-arrest protein 2–related homology (GAR) domain (Fig. 2 A); there is over 60% identity between the VAB-10B GAR domain and the GAR domain–mediating microtubule (MT) binding in MACF and Shot (Sun et al., 2001; Lee and Kolodziej, 2002b). Compared with vertebrate plakins, VAB-10A is more similar to plectin than to BPAG1-e, and VAB-10B slightly more similar to MACF than to BPAG1-a (Fig. 2 and Fig. S2).

Genetic data validate the molecular model described Fig. 2 A, which predicts that vab-10A and vab-10B are two distinct transcription units sharing a common 5′ region. The h1356 mutation affects a splice donor site in the common region and should truncate all isoforms; e698 and ju281 are missense mutations affecting vab-10A isoforms, and mc44 is a 1-kb deletion in exons 24–25 that should eliminate the last 1,400 amino acids of VAB-10B isoforms (Fig. 2 A). Consistent with their positions, h1356 failed to complement the other vab-10 alleles, whereas ju281 and e698 did complement mc44 (Fig. 2 B). Hereafter, we refer to these alleles as vab-10(h1356), vab-10A(ju281), vab-10A(e698) and vab-10B(mc44). We suggest that vab-10(h1356) and vab-10B(mc44) are very strong or null vab-10 and vab-10B alleles, respectively, as their phenotypes did not become more severe in trans to the deficiency h2D17. In contrast, we suggest that vab-10A(ju281) and vab-10A(e698) are not vab-10A null alleles, as their phenotypes became more severe in trans to vab-10(h1356) (Fig. 2 B).

VAB-10A and VAB-10B have nonoverlapping distributions in the epidermis

The preceding molecular and genetic data predict that VAB-10A and VAB-10B isoforms should fulfill distinct functions. To address this hypothesis, we set out to examine whether their cellular and subcellular distributions are similar or different. We raised pAbs against one VAB-10A–specific and two VAB-10B–specific domains (Fig. 3). In addition, we used the mAb MH5, known to recognize an FO component (Francis and Waterston, 1991). While characterizing vab-10, we noticed that MH5 failed to stain vab-10(b1356) embryos (Fig. 4 D), suggesting that it might recognize a VAB-10 isoform. Several lines of evidence establish the specificity of these antibodies and demonstrate that MH5 specifically recognizes VAB-10A. First, rabbit polyclonal VAB-10A antibodies detected a band on Western blots that co-migrated with the band detected by MH5 in the 300–400 kD range, in agreement with previous estimates (Francis and Waterston, 1991). Second, VAB-10B antibodies detected a band that migrated above VAB-10A (Fig. 3), consistent with the predicted sizes of both plakins. Third, addition of the polypeptide used to raise the antibodies eliminated the signal recognized by VAB-10A and VAB-10B pAbs (Fig. 3), testing to their specificity. Last, we could further map the epitope recognized by MH5 within the VAB-10A–specific alternative exon 16 (Fig. S3).
In embryos, VAB-10A and VAB-10B antibodies first detected a signal at the basal and apical plasma membranes of dorsal and ventral epidermal cells, soon after the onset of differentiation (Fig. 4, A and F). As these cells became thinner during morphogenesis, staining appeared as four longitudinal rows (Fig. 4, B–E and G–L) and progressively evolved into circumferentially oriented bands located above muscle sarcomeres (Fig. 5 A). We also detected VAB-10A basally and apically in the pharynx and along mechanosensory axons as described previously (Francis and Waterston, 1991). VAB-10B was also present in the pharynx lumen, intestine lumen, nerve ring, and in body wall muscles and somatic gonad of larvae (Fig. 4, F–H, and unpublished data; a detailed account of VAB-10 distribution and role in nonepidermal tissues is hoped to be presented elsewhere). These patterns are specific because MH5 and polyclonal VAB-10A antibodies failed to detect a signal in \( vab-10 \) (h1356) and \( vab-10A \)-deficient embryos (Fig. 4, D–E, and Fig. 6 H). Likewise, VAB-10B antibodies could only detect a weak signal in the intestine and pharynx of mid-staged \( vab-10B(mc44) \) and \( vab-10(b1356) \) embryos (Fig. 4, J and L). Although we cannot exclude that VAB-10B antisera recognize a cross-reacting protein, we believe that VAB-10B is expressed in the pharynx and the intestine because surviving \( vab-10B(mc44) \) larvae had an abnormal intestine (unpublished data).

To further define the subcellular level of VAB-10A and VAB-10B in the epidermis, we determined if they colocalize with IFs and myotactin, which are associated with FOs. Using confocal microscopy, we observed that the parallel bands formed by VAB-10B are interspersed with those formed by VAB-10A (Fig. 5, D/D ; notice the alternating green and red bands). Reciprocally, we observed that the parallel bands formed by VAB-10B are interspersed with those formed by VAB-10A (Fig. 5, D/D ; notice the alternating green and red bands). Reciprocally, we observed that the parallel bands formed by VAB-10B are interspersed with those formed by VAB-10A (Fig. 5, D/D ; notice the alternating green and red bands). Reciprocally, we observed that the parallel bands formed by VAB-10B are interspersed with those formed by VAB-10A (Fig. 5, D/D ; notice the alternating green and red bands). Reciprocally, we observed that the parallel bands formed by VAB-10B are interspersed with those formed by VAB-10A (Fig. 5, D/D ; notice the alternating green and red bands). Reciprocally, we observed that the parallel bands formed by VAB-10B are interspersed with those formed by VAB-10A (Fig. 5, D/D ; notice the alternating green and red bands).
bands corresponded to the regularly spaced shallow furrows that pattern the cuticle and separate annuli ridges (Fig. 5, E–E’; for a description of annuli, see Costa et al., 1997). Immunoelectron microscopy confirmed that VAB-10A (but not VAB-10B) is restricted to FOs. As shown in Fig. 5 (F and G), VAB-10A was exclusively detected in the epidermis in areas dense in filaments, a characteristic of FOs, raising the possibility that VAB-10A might interact with IFs. Using the VAB-10B K22 antibodies, we observed gold particles in the epidermis adjacent to body wall muscles and in sarcomeres (Fig. 5 H). Although the number of particles is low, quantitative analysis indicates that staining is specific (see legend to Fig. 5). Thus, VAB-10B appears to have a disperse distribution in the epidermis and muscles.

VAB-10A is necessary for epidermis attachment to the ECM and VAB-10B to link the apical and basal epidermal plasma membranes

The distinct distributions of VAB-10A and VAB-10B suggest that these isoforms carry out different functions. Consistent with this notion, vab-10A and vab-10B mutants exhibit different terminal phenotypes. Specifically, embryos with no (or strongly reduced) VAB-10A activity arrested before the two-fold stage of elongation showed no or very little muscle contractions, but had an apparently normal intestine and pharynx (Fig. 1, C and D). vab-10B(mc44) mutants also failed to elongate beyond the 2.5-fold stage; however, most could hatch (Fig. 1 E) to generate larvae that generally died during the L1 stage with a very irregular and lumpy body morphology. The phenotype of vab-10B(h1356) embryos was more variable, but not more severe (Fig. 1 F).

The paralysis of vab-10A mutants indicates that muscles are not functional, and the lumpiness of vab-10B mutants primarily unveils epidermal defects. Because VAB-10A is as-
associated with FOs, which are envisioned as key structures for muscle–epidermis–cuticle attachment, loss of VAB-10A function would be expected to cause the epidermis to detach from the ECM, whereas loss of VAB-10B function should have different consequences. To address this issue, we analyzed vab-10A and vab-10B mutants by light and electron microscopy.

To examine the morphology of muscles and the epidermis, we visualized these tissues with specific antibodies. We used an mAb against a muscle protein with the same distribution as paramyosin (Schnabel, 1995), and VAB-10A antibodies (in vab-10B–deficient embryos) or VAB-10B antibodies (in vab-10A–deficient embryos); we also used VAB-10A antibodies to stain vab-10A(ju281) embryos, as ju281 does not affect VAB-10A synthesis. In wild-type animals, muscles are closely apposed to the epidermis at early stages of elongation (Fig. 6, A–C). Similarly, we found that in VAB-10A–deficient embryos, the positions and shapes of muscle and epidermal cells were normal until the 1.5-fold stage of elongation (Fig. 6, G–J). However, in vab-10B(RNAi) (Fig. 6, D–F) and vab-10B(mc44) (unpublished data), at a stage when muscles are not yet functional, muscle cells were located at an unusually large distance from the edge of the embryo on its dorsal side; in these embryos, VAB-10A was detected at two positions, along muscles and along the edge of the embryo. Because VAB-10A is basal and apical in younger wild-type embryos (Fig. 4 A), we interpret this image to mean that muscles still adhere to the epidermis, but that the thickness of epidermal cells dramatically increased.
As muscles become functional (normally at the 1.7/1.8-fold stages), they remain closely apposed to the epidermis in wild-type embryos (Fig. 6 M). However, in vab-10(b1356), vab-10A(RNAi), and the most severely affected vab-10A(ju281) embryos, they gradually collapsed to a central position (Fig. 6 K) while remaining attached to each other and organized into four enlarged quadrants. We believe that muscle activity is directly responsible for this phenotype because muscles were still found attached in the head and tail regions when muscle activity was blocked by simultaneously inducing RNA interference (RNAi) against vab-10A and against myo-3 (Fig. 6 L), the main myosin heavy chain gene (Williams and Waterston, 1994). In addition, VAB-10A and VAB-10B became mutually dependent for their maintenance in the epidermis. VAB-10A was absent in the bodies of most vab-10A(ju281) (Fig. 6 N) and vab-10B(mc44) (Fig. 6 O) embryos where muscles had detached. Similarly, VAB-10B was locally undetectable in some vab-10A(ju281) embryos mainly in areas where muscles had detached (Fig. 6, P–R, arrows), and was essentially absent at the end of embryogenesis (Fig. 6 K).

The results described (along with Fig. 6) suggest that VAB-10A is essential to anchor muscle cells to the epidermis, and that VAB-10B is required to maintain the distance between the apical and basal plasma membranes of the epidermis. EM analysis allowed us to refine these conclusions. In wild-type embryos, sarcomeres are small relative to larval and adult stages, and the dense bodies resemble electron-dense plaques at the plasma membrane instead of the elongated plaques spanning the sarcomeres of larvae and adults (Fig. 7 A). However, in the overlying epidermis,
FOs had the same appearance as in adults, looking like regularly spaced electron-dense dots found between the developing cuticle or the basal lamina and the epidermis (Fig. 7 A). In vab-10(h1356) (Fig. 7 E) and vab-10A(RNAi) (Fig. 7 F), sarcomeres appeared severely disorganized and were not closely apposed to the epidermis. Some greyish material, which probably corresponds to ECM material, accumulated between muscles and the epidermis. Within epidermal cells, we observed fewer FOs than normal in vab-10A(ju281), vab-10A(RNAi), and vab-10(h1356) embryos with gaps between the cuticle and the epidermis (Fig. 7, B, C, and E). One reason why the cuticle–epidermis gap was not wider, except occasionally (Fig. 1 C, arrow), could be that once muscles have pulled away, there is no more tension exerted to widen it. Because we failed to observe similar defects in control embryos, we conclude that VAB-10A is essential to assemble or maintain FOs. In contrast, muscles remained closely apposed to epidermal cells in vab-10B(RNAi) embryos, but the thickness of epidermal cells was increased even in areas where the number of FOs was normal (Fig. 7 D).

**VAB-10A is essential to organize FO components, VAB-10B to strengthen actin filaments**

The potential cytoskeleton-binding domains of VAB-10 proteins and the association of VAB-10A with FOs provide some clues as to their functions in epidermal cells. Thus, we examined if these plakins were required for the organization of the cytoskeleton and FO components. Immunostaining with mAbs recognizing IFs and myotactin reveals a regular pattern of circumferentially oriented bands that becomes distinct in wild-type pretzel-staged embryos (Fig. 8, A and C; see also Fig. 5, B and C). However, IFs and myotactin almost never form parallel bands in vab-10A(RNAi) embryos, except occasionally in the head or the tail, and were generally absent dorsally (Fig. 8, B and D). Ventrally, IFs often collapsed to form thick bundles (Fig. 8 B, arrowheads), which first became apparent at mid-embryogenesis, whereas
myotactin was punctate (Fig. 8 D). These defects, although less severe, were also observed in vab-10A(ju281)– and VAB-10B–deficient embryos (unpublished data).

Using phalloidin staining, we found that actin MFs could form in vab-10A(RNAi) embryos and generally had a normal length, but were often not circumferentially oriented (Fig. 8, E and F). In vab-10B(RNAi) embryos, phalloidin staining was reproducibly fainter, to the point that actin MFs could hardly be detected in some areas (Fig. 8 G). Both types of defects could be seen in vab-10(h1356) homozygous mutants, in that actin MFs were frequently disorganized and shorter or absent (Fig. 8 H). These data suggest that VAB-10 proteins possess cytoskeleton organizing properties characteristic of plakins.

Discussion

Our data demonstrate that vab-10 corresponds to the C. elegans spectraplakin locus, a plakin subfamily defined by Drosophila shot and vertebrate BPAG1 and MACF1 loci (Roper et al., 2002). By alternative splicing of a common 5' region to different sets of 3' exons, vab-10 generates two distinct classes of isoforms with essentially nonoverlapping subcellular distributions and different functions in the epidermis. Isoforms related to plectin and BPAG1-e (termed VAB-10A) are essential to maintain the epidermis–ECM attachment. Isoforms related to MACF and BPAG1-a (termed VAB-10B) maintain a connection between the apical and basal epidermal plasma membranes during morphogenesis. A model for VAB-10 distribution and function is presented in Fig. 9 and discussed further in the next paragraph.

vab-10 and other spectraplakin loci

vab-10 shares with other spectraplakin loci the potential to generate a complex set of isoforms, including isoforms with COOH-terminal plectin repeats (VAB-10A) and isoforms with COOH-terminal spectrin repeats (VAB-10B). VAB-10 isoforms possess a potential ABD and VAB-10B a potential microtubule-binding domain, which are well conserved; in contrast, the potential IF-binding domain of VAB-10A isoforms is less well conserved. Future experiments should allow us to test whether these domains are indeed functional. However, our data do not support the existence of a giant vab-10 transcript similar to BPAG1-b with both central plectin and COOH-terminal spectrin repeats, or imply that such an isoform must be rare and have a minor nonessential function. First, we failed to identify a giant transcript by RT-PCR experiments. Second, VAB-10A and VAB-10B isoforms have different tissue and subcellular distributions, except in the epidermis of early embryos. Finally, vab-10A mutations complement vab-10B mutations. Our identification of multiple vab-10 alleles, in an organism that has IFs, provides a more complete genetic analysis of spectraplakin function. In vertebrates, the multiplicity of plakins raises issues of genetic redundancy and complicates the analysis of their protective function during morphogenesis. In Drosophila, which has no IFs, previous analyses did not specifically address the function of plectin repeat–containing isoforms (Roper et al., 2002).

FOs are molecularly and functionally homologous to hemidesmosomes

Genetic analysis of vab-10 mutants strongly suggests that VAB-10A is an essential component of FOs, which have been proposed to mediate muscle–cuticle attachment (Francis and Waterston, 1985). Our data provide the first unambiguous demonstration that FOs are indeed essential to anchor muscles to the cuticle. First, we show by immunoelectron microscopy that VAB-10A is concentrated in FOs. Next, we demonstrate that in the absence of VAB-10A, the number of FOs is reduced, and the epidermis detaches from

Figure 8. vab-10 is required to organize other FO components and actin MFs. Late-stage embryos were stained with the IF-specific mAb MH4 (A and C), the myotactin-specific mAb MH46 (C and D), or with rhodamine-conjugated phalloidin (E–H). (A and B) Wild-type threefold stage embryo; actin MFs are evenly spaced along what could correspond to lateral membranes (C, arrowheads). (E) Wild-type threefold stage embryo; actin MFs are evenly spaced and oriented along the circumference (arrows). (F) vab-10A(RNAi) embryo; many actin MFs are irregularly oriented. (G) vab-10B(RNAi) embryo; actin MFs were reproducibly thin and difficult to visualize. (H) vab-10(h1356) embryo; actin MFs were short or absent and randomly oriented. Bar, 10 μm.
VAB-10B maintains epidermal integrity

Our results demonstrate that VAB-10B and VAB-10A roles differ. Loss of VAB-10B function leads to increased epidermal thickness at a time when muscle cells are not yet functional, a phenotype that is not observed in VAB-10A–deficient embryos of the same age. This change was visible predominantly on the dorsal side of the embryo, where the epidermis is strongly bent and formed by several cells that fuse at approximately the same time to form a large syncytium called hyp7. Thus, one appealing possibility could be that VAB-10B acts to protect cells against mechanical forces generated within cells that change their shapes. VAB-10B could do so in several ways. For example, Drosophila Shot is concentrated at the end of MT bundles in neurons and tendon cells, and is required to anchor these bundles (Gregory and Brown, 1998; Prokop et al., 1998; Lee and Kolodziej, 2002b). By analogy, VAB-10B, which has well conserved actin- and MT-binding domains, could attach to cortical actin and anchor MTs, which are generally oriented along the apico–basal axis in epithelial cells (Fig. 9). An alternative possibility, which takes into account the large size of VAB-10B (4950 aa), the thinness of the epidermis in muscle contact areas (200 nm in early embryos and 50 nm in later embryos), and plakin potential to form dimers through their rod domains (Leung et al., 2002), is that VAB-10B molecules located on the basal and apical membranes dimerize to maintain epidermal thickness (Fig. 9).

Localized change of epidermal thickness might explain why muscles also detach from the epidermis in VAB-10B–deficient embryos. IFs might be unable to maintain a connection between FO plaques when the epidermal thickness increases. In addition, as muscle cells are known to help maintain FOs in the adjacent epidermis (Hresko et al., 1999), perhaps localized weakening of FO integrity triggers a chain reaction, whereby muscle partially detaches, causing further FO instability and increased muscle detachment.

Organization of the cytoskeleton

The mechanism that controls the formation of alternating VAB-10A and VAB-10B circumferential bands is intriguing. The fact that both VAB-10B and actin filament bundles are localized at furrows separating annuli, and the observation that the actin cytoskeleton is perturbed in VAB-10B–deficient embryos, suggest that VAB-10B could play a key role in the initial events that organize VAB-10 and actin distri-
bution. Because VAB-10A and VAB-10B are mutually dependent on each other for their stabilization, the establishment of the alternating pattern of VAB-10B with VAB-10A could be governed by rules of self-organization as often observed for the cytoskeleton (for a discussion on self-organization, see Misteli, 2001).

Elongation of the C. elegans embryo depends on the epidermal cytoskeleton and on a communication between muscles and epidermal cells. We have shown that VAB-10 isoforms are essential for embryonic elongation. Future analysis of vab-10 function should be invaluable to understand how spectraklins can modulate cytoskeletal dynamics and signaling between tissues during morphogenesis.

Materials and methods

Worm culture and strains

C. elegans strains were grown on NGM plates and fed with OP50 Esche- richia coli strain at 20°C unless otherwise stated (Brenner, 1974). The alle- lele vab-10(e698) (Hodgkin, 1983) and the chromosomal deficiency hDf17 were obtained from the Caenorhabditis Genetics Center (University of Minnesota, Minneapolis, MN).

vab-10 mutations and cloning

The ethyl methane sulfonate–induced lethal mutation h1356 belongs to a collection of eight lethal mutations isolated in the Rose laboratory (University of British Columbia, Vancouver, Canada). The mutation ju281 was iso- lated after ethyl methane sulfonate mutagenesis in screens for larval lethal mutations performed in the Chisholm laboratory (University of California, Santa Cruz, Santa Cruz, CA). We recovered the allele vab-10(zk1151.3)-mc44 by PCR screening after trimethylpsoralen/UV mutagenesis using two pairs of nested primers located 2.3 kb apart in vab-10 exons 23–24 (vab-10B–spec- cific). One pool carrying a 1-kb deletion (mc44), among 96 pools of 25 F1 mutagenized animals, was sib-selected; heterozygous animals segregating mc44 were outcrossed six times to wild-type animals and balanced. The molecular lesions affecting vab-10 alleles were identified by sequencing each predicted vab-10 exon after PCR amplification from h1356- and ju281–arrested embryos or e698 adults. Potential mutations were con- firmed on an independent embryo/animal. The sequence of mc44 was deter- mined from mc44–arrested larvae. Complementation tests between vab-10 alleles were performed by crossing vab-10(x)/+ males with vab-10(y)/+ hermaphrodites. Mating partners were transferred to fresh plates twice daily, and the number of unhatched eggs, arrested larvae, normal males, and hermaphrodites was counted; retarded larvae (putative trans-heterozy- gotes) were placed on separate plates and were examined separately.

To molecularly identify vab-10, we used RNAi (Fire et al., 1998) against predicted genes mapping under hDf17 (www.wormbase.org). RNAi against ZK1151.1, ZK1151.2, and ZK1151.3, the three genes predicted within cosmid ZK1151 (GenBank/EMBL/DDJB accession no. ZK1151.3), re- sulted in phenotypes similar or almost identical to those of strong vab-10 alleles, suggesting that they define a single locus, RT-PCR experiments were performed using the one-step RT-PCR kit (QIAGEN) starting from to- tal RNA (see Fig. S1). The GenBank/EMBL/DDJB accession nos. for vab-10 sequences are AJ505815 and AJ505816.

RNAi

Double-stranded RNA (dsRNA) used for RNAi was prepared and injected as described previously (McMahon et al., 2001). RNAi against vab-10 common exons (positions 25635–26646 of ZK1151) generated a phenotype indistin- guishable from that of h1356 mutant embryos. RNAi against vab-10A–spec- cific exons was obtained with either of two regions from the large vab-10A–specific exon (positions 14630 to 15756, or 19490 and 20417 of ZK1151); both dsRNAs produced the phenotypes shown in Fig. 1 E. RNAi against vab-10B–specific exons was obtained by co-injecting two dsRNAs from different areas of vab-10B (positions 5807 of Y47H9B and 518 of ZK1151, plus 2018 and 3136 of ZK1151). This resulted in mc44-like phenotypes within 24–30 h after injection, then phenotypes became weaker; injection of each dsRNA separately resulted in weaker and less penetrant phenotypes.

Generation of VAB-10 antibodies

To produce antibodies, GST fusions with RT-PCR fragments encoding VAB-10A residues G2837–Q3436, VAB-10B residues D2697–R3862, or VAB-10B residues E3863–K9455 were purified using DEAE-Sepharose fol- lowed by glutathione-Sepharose 4B columns, and were used to immunize two male New Zealand rabbits. Sera were purified by incubation for 3 h with fusion proteins that had been transferred onto nitrocellulose filters, followed by washing with 0.15 M NaCl for 30 min and PBS for 5 min. Spec- ific antibodies were eluted by 0.1 M glycine-HCl (pH 2.5) for 10 min, and 1 M Tris-HCl (pH 8.0) was added to neutralize the solution. These antibod- ies were used to detect VAB-10B isoforms (1/1,000 dilution) on Western blots of total worm extracts separated by electrophoresis on 5% acryl- amide gels.

Fluorescence microscopy

Embryos were fixed and stained by indirect immunofluorescence or by rhodamine-conjugated phallolidin as described elsewhere (Costa et al., 1997). Primary antibodies were diluted 1:1,000 for VAB-10A– and VAB-10B–immunopurified pAbs, or 1:50 for MH15, MH4, MH46, MH13, MH24, NE8/46C, and 5.6.1.1 mAbs (Miller et al., 1983; Francis and Waterston, 1991; Schnabel, 1995). MH-series antibodies were a gift from Chelly Hresco and Bob Waterston (Washington University, St. Louis, MO), NE8/ 4C6 from the Medical Research Council, and 5.6.1.1 from David Miller (Vanderbilt University, Nashville, TN). Primary antibodies were detected using Cy3- or FITC-conjugated secondary antibodies. Stacks of images ev- ery 0.5 μm (0.2 μm for images in Fig. 5 and Fig. 8, E–H) were captured us- ing a confocal microscope (model Sp1; Leica); generally, 12–15 confocal sections were projected using the Tcstk software (McMahon et al., 2001) and then processed using Adobe Photoshop.

EM

Mutant embryos laid between 6 and 9 h before processing were prepared for EM as described elsewhere (McMahon et al., 2001). Only nonruptured embryos that had reached the 1.5–1.7-fold stage were fixed. Thin sections were examined with a microscope (model EM208; Philips). We examined wild-type or heterozygous mutant/+ background, three embryos and five larvae; h1356, four embryos; ju281, four embryos; vab-10A/RNAi) three embryos; and vab-10B/RNAi, three embryos and six larvae. In each case, at least 10–15 sections at different levels were examined using a 10,000– 15,000 magnification.

Immunoelectron microscopy

Wild-type adults were prepared for post-embedded immunolabeling as previously described for studies in other invertebrates (McDonald, 1999). In brief, young adults were immobilized by high pressure freeze, fixed in anhydrous methanol with 4% PFA and 1.5% uranyl acetate at −90°C, and infiltrated at −45°C with lowicryl HM20 (PolyScience, Inc.) that was then polymerized by UV exposure. Thin sections (~80 nm) were incubated with either VAB-10A 1:1,000, VAB-10B 1:2,000, or no primary anti- body, and were labeled with gold colloidal suspensions of a 15-nm diame- ter coupled to GAR-IgG (BBInternational) according to manufacturer’s rec- ommendations (Aurion). Specificity of staining was judged based on comparing the number of gold beads localized in muscle, epidermis, cuti- cle, and resin in primary antibody–stained and control grids.

Online supplemental material

Fig. S1 presents some of the RT-PCR experiments that lead us to propose the structure of vab-10 transcripts shown in Fig. 2. Fig. S2 pre- serves, in further details, alignments between the cytoskeleton-binding do- mains of VAB-10 and other plakins. Fig. S3 describes how we mapped the epitope recognized by the mAb MH5 to the vab-10A exon 16. Online supplemental material available at http://www.jcb.org/cgi/content/full/ jcb.200302151/DC1.

We are indebted to Chelly Hresco for the gifts of MH4, MH5, MH27, and MH46 antibodies. We also thank David Miller and the Medical Research Council for additional antibodies; the Caenorhabditis Genetics Center for strains; Philippe Rostaing, Erik Jorgensen, andannonce for expert technical assistance with immunoelectron microscopy; the EM facility of IFR 83 for access to the high pressure machine; Jean-Luc Vonesch and Marcel Boeglin for assistance with confocal microscopy; and Elisabeth Georges-Labouesse for critical reading of the manuscript. J.M. Bosher was supported by fellowships from the Fondation pour la Recherche Médicale and the AFM, B.-S. Hahn by the Ministère de la Recherche, and R.M. Weiner by the INSERM “postes verts” program. This work was supported by institutional funds from CNRS, INSERM, Hôpitaux Uni- versitaires de Strasbourg, and grants to M. Labouesse from the Association pour le Recherche Sur la Cancer and the EEC-TMR program. Light camera...
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