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Cell invasion through basement membranes: an anchor of understanding

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To metastasize, cancer cells must acquire the ability to breach several basement membrane barriers. Cell invasions through basement membranes also occur during normal development and immune system function, enabling organ formation and cell dispersal. The mechanisms that cells use to cross basement membranes *in vivo* remain elusive. In cancer and development, these invasions occur in complex and inaccessible environments, which are difficult to study *in vivo*. Anchor-cell invasion in *Caenorhabditis elegans* is a simple, visually and experimentally accessible model of basement membrane invasion that is beginning to reveal a network of cellular and molecular control mechanisms that regulate the fundamental cellular process of invasion through basement membranes.

Introduction

The acquisition of cell-invasive behavior, which includes the ability to adhere to and migrate through the extracellular matrix, is an essential step in cancer metastasis. There has been great interest in understanding the mechanisms that promote cell invasion in an effort to design therapies to control this aspect of cancer progression [1–3]. Cell invasions occur frequently in the course of normal animal development and during leukocyte trafficking but, in contrast to cancer, these invasions are tightly regulated [4-8]. Key genes regulating invasive cell traits in normal contexts, such as epithelialmesenchymal transitions and migratory ability, often have oncogenic roles in metastasis [5,9-12], demonstrating that a crucial element in the acquisition of invasive behavior in cancer is the inappropriate regulation of genes whose normal function is to control cell invasion.

Though necessary for successful invasion, the mechanisms that guide the movement and penetration of cells through the extracellular matrix remain one of the least understood aspects of cell-invasive behavior. The extracellular matrix is a highly diverse and dynamic protein network that provides tissue level structural support and influences the differentiation and functioning of surrounding cells [13]. Extracellular matrices take on different structures, ranging from matrix packed bone and cartilage, to more porous connective tissue, to the dense, sheet-like basement membrane. Cell invasions occur through all of these extracellular matrices in normal development and during cancer metastasis [14–17]. Given the diversity of structures, invasive cells probably have a range of mechanisms to navigate through these matrices. Even within one type of matrix, the connective tissue, invasive cells can use several distinct mechanisms to mediate migration [18–20].

A crucial bottleneck in understanding the process of cell invasion through extracellular matrices has been the challenge of studying this process in vivo [15,21]. In vitro assays, such as the Boyden chamber assay through Matrigel, have important roles in revealing potential regulators of adhesion and invasion through extracellular matrix [22-24]. In vitro assays, however, must be interpreted with caution, as they are outside the context of normal cellular interactions and the cell-associated matrix. Furthermore, unlike endogenous matrix, extracted Matrigel is not crosslinked [25], and important differences in invasion through Matrigel and in vivo connective tissue matrix have been observed recently [18,20]. Anchor cell (AC) invasion in Caenorhabditis elegans is a new model for examining cell invasion through basement membranes, in which cellular, genetic and molecular dissection of invasive behavior can be directly addressed in vivo [26]. Here, I focus on the importance of cell invasion through basement membranes in cancer and normal physiology and on new insights being gained by studying AC invasion in C. elegans.

Invasion through basement membranes in cancer and normal development

The basement membrane is a cell-associated extracellular matrix that underlies epithelia and endothelia and surrounds peripheral nerve fibers, muscles and fat cells. Basement membranes regulate tissue structure, cell function and act as barriers between tissue compartments [27]. All vertebrate basement membranes contain laminin, type IV collagen, nidogen and proteoglycans, which together form dense sheet-like structures that are 50–100 nm thick [27,28]. These major basement membrane proteins have orthologs in *C. elegans* that participate in basement membrane assembly and function [29,30], suggesting that this structure is an ancient form of matrix.

Epithelial-derived tumors, also known as carcinomas, represent ~90% of all cancers in the USA and Europe [31]. The metastatic spread of these cancers is the most common cause of death in cancer patients [21,32]. When epithelial-derived tumors metastasize, the first barrier that invading cells breach is the basement membrane

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underlying the epithelium. Freed cells traverse the stromal connective tissue, a gelatinous extracellular matrix rich in proteoglycans that surrounds glands and blood vessels, and then cross the basement membranes of blood or lymphatic vessels to disperse in these vessels, a process known as intravasation [15]. During extravasation, the invasive cells cross back through the vascular basement membrane into new tissues where they can form micrometastases [21].

Invasions through basement membranes also occur during normal development and immune system functioning, in which tightly regulated cells undergo basement membrane crossings to help construct tissues and disperse cells. Examples include primate trophoblast cells that breach basement membrane underlying the endometrium to establish a functional placenta [4], neural crest cells and myoblasts that cross basement membranes during their migrations [33,34], capillary sprouts that penetrate through the surrounding vessel basement membrane to form new vessels [6,35] and leukocytes crossing through perivascular basement membrane to sites of injury and infection [8]. Despite the significant barrier function of basement membranes, invasive cells have adopted successful strategies to target and breach this dense and highly cross-linked matrix repeatedly.

AC invasion in *C. elegans*: a stereotyped cell invasion event

The simplicity, complete transparency and amenability to genetic and genomic approaches have made C. elegans an important model system for addressing fundamental biological processes [36]. During larval development in C. elegans, the uterine and vulval tissues attach together to form a uterine-vulval connection that is an essential structure in the adult for mating and moving embryos from the uterus to the outside environment. The uterine and vulval cells in C. elegans initially develop separately, isolated from one another by distinct gonadal and ventral epidermal basement membranes [37]. The AC, a single specialized gonadal cell, crosses these basement membranes during the mid L3 larval stage with a basolateral process that then moves between the central vulval cells (the 1° vulval lineage cells) to initiate attachment and properly align the future uterine-vulval connection (Figure 1) [26,38]. AC invasion is invariant from animal to animal: the AC breaks through both basement membranes in ~ 30 min and always invades at the mid L3 larval stage between the central 1° vulval lineage cells [26].

The cellular mechanisms that control the targeting and timing of AC invasion

During the late L2 to early L3 larval stage, the AC induces three epidermal precursor cells along the ventral midline to generate the vulval cells of the hermaphrodite through the epidermal growth factor (EGF) signaling pathway [39]. The epidermal precursor cell P6.p, which is closest to the AC, is induced to a 1° fate and gives rise to the eight specialized vulval cells positioned at the center of the vulva by the end of the L3 stage. The 1° vulval cells have a crucial role in regulating the timing and targeting of AC invasion by means of a diffusible cue(s) [26]. When 1° vulval cells are removed, the AC fails to break down and invade through the basement membranes in most animals. Experimental manipulations moving the 1° vulval cells from their normal central position results in ACs that invade toward these inappropriately placed cells in response to their chemotactic invasion cue(s) (Figure 2a–c).

The ability of the AC to respond to the 1° vulval cue also controls the timing of its invasion. Taking advantage of mutations in *C. elegans* that alter developmental timing events in specific tissues (heterochronic mutants), it has been possible to examine AC invasion in animals in which the 1° vulval cells form precociously during the late L2 larval stage [40,41]. Rather than invading precociously, the AC invades at or near the normal time of invasion at the mid L3 larval stage [26]. Thus, although the signal to invade is present, apparently the AC is not ready to invade at the L2 stage.

A third mechanism regulates the timing of AC invasion. Whereas most ACs fail to invade in vulvaless animals, $\sim 20\%$ of ACs nevertheless break through the basement membrane with small protrusions, which enter the underlying hypodermis during the mid L3 stage, the normal time of invasion [26]. This AC behavior indicates that an unidentified extrinsic signal from nonvulval cells or an AC cell-autonomous program helps drive AC invasion weakly during the mid L3 stage. Together, the 1° vulval cue, the vulval-independent mechanism and the capacity of the AC to invade ensure that AC invasion is executed with spatial and temporal precision.

Similar mechanisms regulate invasion in other cells. In a mouse model for intravasation, cells from metastatic MnTLn3 rat tumors polarize, migrate towards and cross through the basement membranes of blood vessels that generate a chemotactic signal, possibly EGF from tumorassociated macrophages [42–44]. Many metastatic cancers express the chemokine receptor CXCR4, which could help target invasion during extravasation to particular tissues expressing the CXCR4 ligand CXCL12 [45–47]. The chemokine receptor CCR1 is also expressed on trophoblast cells as they differentiate to an invasive phenotype [48]. Furthermore, similar to the AC, the time that trophoblast cells can invade is tightly regulated [49].

The AC is polarized

During invasion, only the basolateral portion of the AC extends an invasive process through the basement membrane and between the 1° vulval cells. The apical end of the AC always remains attached to neighboring uterine cells. The AC is thus polarized; the basolateral region of the AC has migratory mesenchymal characteristics and the apical portion has cell-cell adhesion epithelial qualities. Molecular markers further confirm this polarity. The Fat-like cadherin CDH-3 localizes specifically to the invasive basolateral membrane of the AC during invasion [50], whereas AJM-1, an apical adherens protein expressed in epithelial cells, forms junctions in the apical region of the AC (Sherwood, D.R., unpublished). Several cells in development and cancer maintain cell-cell junctions yet have active cellular Review



Figure 1. AC invasion through basement membrane initiates uterine–vulval attachment. Panels (**a**–**d**) viewed with Nomarski optics (left), overlaid with the GFP fluorescence signal from a *cdh-3*:GFP transcriptional reporter that drives GFP expression in the AC (center), and in a schematic showing the basement membrane (BM) in red (right). Anterior is to the left, ventral is down and the scale bar is 10 µm. (a) At the early L3 stage, the ectodermal precursor cell P6.p (bracket) has been specified to a 1° vulval cell fate. The gonadal AC (arrow) is directly over the P6.p cell and does not cross the gonadal and ventral epidermal basement membranes (arrowhead) that separate the gonad and vulval cells. With Nomarski optics increased contrast is observed where there are differences in thickness and refractive index of the specimen. This is the case when the basement membrane is intact between the uterine and vulval cells: a dark high contrast line represents the intact basement membrane. (b) During the mid L3 stage, the 1° fated P6.p cell has divided (bracket) and the basolateral portion of the AC (arrowhead) has broken through and initiated movement across the basement membranes and begun to directly contact the 1° vulval cells. (c) At the mid-to-late L3 stage, the basolateral region of the AC has completed crossing through the basement membranes, is firmly attached to the 1° fated vulval cells, which have now formed four cells (bracket), and is invading between the central 1° vulval cells (arrowhead). (d) By the early L4 stage, the AC has completed invasion and moved between the 1° vulval cells to the apex of the invaginating vulva (arrowhead). The 1° vulval cells have divided again (eight cells total) and shifted out of the plane offocus. Their position on one side of the AC is indicated by the dotted lines in the schematic image. (e) Immunolocalization of the basement membrane are tightly juxtaposed, thus they are difficult to distinguish. The hole in the basement membranes (arrow), corresponds with the invasiv

protrusions. For example, many carcinomas, including epithelial prostate cancer, lobular breast cancer and melanoma, invade *en masse* rather than as single cells: the invading cells are attached by cell-cell contacts at one end, while extending cellular protrusions from the other [51]. The mechanisms that generate and maintain asymmetry of dynamic cytoskeletal activity at the protrusive membrane while promoting adhesions at the apical membrane are only beginning to be understood [5,51,52]. Thus, it will be important to understand the mechanisms that establish AC polarity and how this polarization impacts AC invasion.

The mechanics of basement membrane crossing

The *in vivo* interactions between invading cells and the basement membrane have been difficult to visualize and understand. For example, in studying the metastatic spread of epithelial-derived tumors, it has been hard to



Figure 2. AC invasion in wild-type animals and *fos-1a* mutants. All animals viewed at the mid-to-late L3 stage and shown with Nomarski (left), fluorescence (left center), overlaid (right center), and schematic (right) images. **(a)** Basement membrane staining (red, small arrow) and *cdh-3*::CFP expression in the AC (green, arrow) show an AC in a wild-type animal that has breached and crossed through a gap in the underlying basement membranes (arrowhead), and is now attached to the central 1° vulval cells (bracket). **(b)** A wild-type animal in which the AC (expressing *cdh-3*::CFP in green) has invaded and contacted the underlying 1° vulval cells that express the marker *egl-17*::YFP (purple, bracket). **(c)** A wild-type animal in which the 1° vulval cells (expressing *egl-17*::YFP in purple) have been displaced from their normal site directly under the AC. In response to a chemotactic cue from the 1° vulval cells, the AC directs an invasive process through the basement membrane (arrowhead) and toward the vulval cells (small arrow). **(d)** and **(e)** *Fos-1* is expressed in the AC. In a *fos-1a* loss-of-function mutant, the AC fails to remove the underlying basement membrane and contact the 1° vulval cells, although they express *egl-17*::YFP normally (bracket, e). **(f)** Moving the 1° vulval cells in a *fos-1a* mutant reveals that the 1° vulval cells generate a chemotactic cue and the AC can extend a basolateral process (small arrow) towards this signal; however, this process flattens at the site of the basement membrane, indicating an inability to breach this barrier. Reproduced, with permission. from [50].

determine whether tumor associated macrophages, the activities of the tumor cells themselves, or both actively create breaches in basement membrane to enable tumor cell dispersal [53]. Similarly, the mechanisms used by leukocytes to cross through perivascular basement membranes are unknown [8]. Visual observations combined with forward genetic and molecular approaches are beginning to reveal how the AC interacts with basement membrane to break through this barrier during its invasion.

The fos-1 gene is the C. elegans ortholog of the vertebrate fos gene family. Fos proteins form heterodimers with Jun or other bZIP proteins to create the AP-1 transcription factor complex [54]. An isoform of the fos-1 gene, fos-1a, is a crucial regulator of basement membrane removal during AC invasion. In fos-1a mutants, ACs never invade at the mid L3 stage (Figure 2d,e) and only occasionally invade in a delayed and partial manner [50]. In fos-1a mutants, the AC continues to respond to the chemotactic cue generated by the 1° vulval cells and extends cellular processes toward 1° vulval cells. These processes, however, cannot cross through the basement

FOS-1A is expressed at high levels in the AC during invasion and restoration of FOS-1A expression in the AC in fos-1a mutants rescues invasion [50]. The other isoform of fos-1, fos-1b, does not rescue invasion in fos-1a mutants, indicating that these two isoforms of fos-1 have distinct activities [50]. In vertebrates there are four fos genes (cfos, fosB, fra-1 and fra-2) that, similar to fos-1a and fos-1b, have different functional properties. c-fos and fra-1 are overexpressed in several metastatic cancers [55–57], including breast, prostate, pancreatic and thyroid, and their ectopic expression in cell lines stimulates cellinvasive behavior through reconstituted extracellular matrix [58,59]. These observations suggest that fos is a conserved component of a regulatory pathway underlying cell-invasive behavior in development and cancer and that c-fos and fra-1 are required in metastatic cancer cells to promote their movement across basement membranes.

membrane and instead flatten at this site (Figure 2f).

Three AC-specific transcriptional targets of *fos-1a* have been identified that are diverse in structure and function: ZMP-1, a GPI-linked matrix metalloproteinase (MMP) with matrix degrading activity [60,61], CDH-3, a Fat-like protocadherin [62] that functions in vertebrates to regulate the dynamics of cellular protrusions [63] and hemicentin, a recently identified conserved extracellular member of the immunoglobulin superfamily [64] that promotes cell associations with basement membranes. These proteins localize at the site of invasion. CDH-3 is found on the invasive basolateral membrane and ZMP-1 localizes to apparent lipid raft microdomains concentrated near the invasive membrane. Hemicentin is deposited in the basement membrane under the AC presaging the future site of invasion where it might mediate AC adherence to the basement membrane to target it for removal [50]. Surprisingly, animals lacking zmp-1, cdh-3 and hemicentin function have only a slight delay in AC invasion, indicating that there are additional and perhaps many fos-1a targets that contribute to basement membrane removal. Whereas in vitro studies have largely focused on the role of MMPs and other proteases in breaching the basement membrane [22], these in vivo observations indicate that basement membrane removal is probably a complex and highly regulated process involving MMPs, as well as several other proteins with diverse functions.

How do the 1° vulval cue and FOS-1A activity coordinate invasion?

The 1° vulval cue and the activity of *fos-1a* are required to execute AC invasion in the mid L3 larval stage. An attractive model for how AC invasive activity could be coordinated would be for the 1° vulval cue to stimulate *fos-1a* expression or activity. This, however, is not the case. In vulvaless animals FOS-1A is expressed and localized to the AC nucleus normally. Furthermore, the expression of the *fos-1a* targets, *zmp-1* and *cdh-3* are not altered and hemicentin is deposited normally in the basement membrane under the AC [50]. Thus, in vulvaless animals, *fos-1a* is active, and it functions to initiate invasion, as

evidenced by the deposition of hemicentin in the basement membrane. However, the 1° vulval signal is required to complete invasion, as invasion fails to occur in most vulvaless animals [26]. The 1° vulval signal might provide the AC with directional information to target invasive processes more effectively or could contribute a key component necessary for the AC to break through the basement membrane.

Future prospects

Analysis of the mechanisms that control AC invasion through the basement membrane has only recently been initiated (Figure 3). Forward genetic and genomic approaches [65-67] combined with the visual cell biological assays described here should identify the molecular pathways that stimulate and target invasion, the mechanisms that establish cell polarity, and the genes that promote basement membrane removal. It is likely that these studies will continue to identify genes with known oncogenic functions (e.g. fos-1), as well as those not previously implicated in regulating invasion or metastasis (e.g. hemicentin). Reverse genetic approaches will also be useful. AC and vulval-specific *cis*-regulatory elements [68] can knock down or target overexpression of genes in C. elegans that are implicated in regulating invasion in metastatic cancers. The specific functions and interactions of genes and genetic networks in the invasion process can then be determined to gain a systems level understanding of how distinct steps in AC invasion are controlled and coordinated. The knowledge gained from these studies might ultimately contribute to the development of novel cancer therapeutics. An understanding of cell invasion through basement membranes is a daunting challenge, but the AC is beginning to break down barriers that have impeded our understanding of this fundamental biological process.



Figure 3. The cellular and molecular networks regulating AC invasion through basement membrane. During the late L2 to early L3 larval stage, the AC induces the 1° fate in the underlying P6.p ectodermal precursor cells through the EGF signaling pathway. FOS-1A is also expressed at high levels and active in the nucleus of the AC at this time (blue circle), where it controls the expression of (1) ZMP-1 (red spots), a GPI-linked MMP that localizes to apparent lipid raft microdomains located near the invasive basolateral membrane; (2) CDH-3 (green line), a Fat-like protocadherin that is found on the invasive basolateral membrane of the AC; and (3) hemicentin (orange, rough line), an extracellular immunoglobulin superfamily member that is secreted directly under the AC before invasion. During the mid L3 stage, the 1° fated vulval cell divides and generates a chemotactic cue that initiates invasion. ZMP-1, CDH-3 and hemicentin appear to be members of a large number of transcriptional targets of FOS-1A that promote removal of the AC and found in large aggregates surrounding the site of invasion, and the basolateral region of the AC begins inserting itself between the central 1° vulval cells. The apical region remains attached to neighboring cells within the gonad and retains an epithelial character.

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