



Prespecification and plasticity: shifting mechanisms of cell migration

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Cell migration is a universal process involving different morphologies and mechanisms in different cell types and tissue environments. Prespecified cell-type-specific patterns of cell migration can be classified into single cell migration (amoeboid, mesenchymal) and collective migration modes (cell sheets, strands, tubes, clusters). These intrinsic molecular programs are associated with a characteristic structure of the actin cytoskeleton, as well as the cell-type-specific use of integrins, matrix-degrading enzymes (matrix metalloproteinases and serine proteases), cell-cell adhesion molecules (cadherins and activated leukocyte adhesion molecule), and signaling towards the cytoskeleton (carried out by RHO GTPases). In response to the gain or loss of these key molecular determinants, significant adaptation reactions can modify the cell's shape, pattern, and migration mechanism; examples of this include the epithelialmesenchymal transition, mesenchymal-amoeboid transition and collective-amoeboid transition.

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Current Opinion in Cell Biology 2004, 16:14-23

This review comes from a themed issue on Cell structure and dynamics Edited by John A Cooper and Margaret A Titus

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DOI 10.1016/j.ceb.2003.11.001

Abbreviations

2D two-dimensional3D three-dimensional

CAT collective–amoeboid transition

EC endothelial cell
ECM extracellular matrix

EMT epithelial-mesenchymal transition
MAT mesenchymal-amoeboid transition

MMP matrix metalloproteinaseMT-MMP membrane-type MMP

ROCK Rho-associated serine/threonine kinase uPA urokinase-type plasminogen activator

uPAR uPA receptor

Introduction

Most cell types in the body, at a given time and tissue space, are capable of — and perform — migration. Initially established for fibroblasts and keratinocytes moving

across 2D substrata, the principle events leading to cell migration within 3D extracellular matrix (ECM) environments are regarded as five interdependent steps that form a continuous cycle [1,2]. Initial cell polarization is driven by localized actin polymerization to filaments that is followed by the extension of a leading pseudopod (step 1). Once the protruding pseudopod touches ECM ligands, adhesion receptors of the β1 and β3 integrin families cluster, attach to the substrate and engage cytoplasmic adaptor, signaling and cytoskeletal proteins towards the adhesion site (step 2). These circumscribed interaction zones, termed focal adhesions or focal contacts, recruit surface proteases, such as MT1-MMP (membrane-type 1 matrix metalloproteinase) and the complex of urokinase-type plasminogen activator (uPA) and its receptor (uPAR), to execute local proteolysis towards adjacent ECM proteins. Pericellular proteolysis is thought to widen extracellular scaffolds for the forward-expanding cell body (step 3). During or shortly after integrin-ligand binding, actin filaments engage with cross-linking and contractile proteins, such as myosin II, which stabilize and contract the actin strands (step 4). The shortening of membrane-anchored actin filaments results in local cell contraction, slow forward gliding of the posterior cell pole and cell translocation along the substrate (step 5). These five steps do not represent a stereotypic program but rather provide an adaptive platform that undergoes cell-type-specific modification dependent on the inherent molecular repertoire and the type of ECM environment.

This review summarizes cellular and molecular signatures of cell migration as well as their reprogramming and reactive compensation in migrating cells.

Prespecified migration mechanisms

In the process of cell differentiation and acquisition of specific tissue phenotypes, different patterns and mechanisms of cell migration emerge. These include the mechanisms controlling the migration of single cells (Figures 1a-c) or, alternatively, collective migration (Figures 1d-g). The central molecules that govern and specify such diverse migration processes comprise the following (Figure 1): matrix-binding adhesion receptors, most notably those belonging to the β1 and β3 integrin families; matrix-degrading proteases of the matrix metalloproteinase (MMP) and serine protease families (uPA/uPAR); molecules that enable cell–cell adhesion and communication (i.e. E-, N- and VE-cadherins and gap junctions); and signaling proteins that control the actin cytoskeleton, most notably the small GTPase RHO and

Box 1 Viewpoint: amoeboid movement in mammalian cells

The sequencing of the Dictyostelium genome will be finished soon (see Dictyostelium Genome Analysis Consortium website at http:// genome.imb-jena.de/dictyostelium/). To date, no predicted homologues of integrins or of secreted or surface metalloproteases have been identified. Dictyostelium is therefore a likely model organism for studies on cell migration processes that do not depend on integrin and MMP functions. In a recent genetic screen at least one novel pattern recognition receptor was identified to contribute critically to adhesive interactions with substrate and migration [3*]. Termed SadA, this receptor contains several EGF-like repeats and a predicted talin binding site, thereby providing putative links between the cytoskeleton and the extracellular substrate. It is reasonable to assume that, in addition to integrins, pattern recognition receptors are retained in higher mammals to provide interactions with ECM ligands and to contribute to cell adhesion and migration in the absence of integrin function.

Amoeboid cells, such as stem cells and most leukocytes, are able to enter and move through many different organs, including skin, gut and brain. They hence retain a powerful capacity for dissemination in connective tissue and recirculation between organs and the blood system as well as the lymphatic system. Other examples of amoeboid movement in the human body are cells from lymphoma, small-cell lung carcinoma and certain mammary carcinoma [2,6]. Because of their striking capacity for efficient dissemination from very small tumor lesions, amoeboid cancers represent systemic diseases metastasizing to distant organs at their earliest stages. Taken together, amoeboid migration in mammals is a feature of nonresident cells that recirculate towards different tissues and migrate therein, yet possess cell-matrix interactions of low stringency and, hence, have limited tissue specificity.

its downstream effector, the RHO-associated serine/ threonine kinase ROCK.

Amoeboid migration

Arguably the most primitive and in some ways the most effective form of cell migration is amoeboid movement (Figure 1a; Box 1), which mimics features of the singlecell behavior of the amoeba Dictyostelium discoideum. Dictyostelium is an ellipsoid cell that has fast deformability (within seconds) and translocates via rapidly alternating cycles of morphological expansion and contraction. Although integrins are not expressed and the binding force towards the substrate is relatively low, migrating *Dictyos*telium cells utilize one or several non-integrin pattern recognition receptors (Box 1) to bind to extracellular structures [3°]. In higher eukaryotes, amoeboid movement is carried out by hematopoietic stem cells, leukocytes and certain tumor cells [4-6] (P Friedl, unpublished). These cells use a fast 'crawling' type of movement that is driven by short-lived and relatively weak interactions with the substrate. In lymphocytes and neutrophils, β1-integrinmediated adhesion is completely or partially dispensable for cell migration within connective tissue, both in vitro [7] and in vivo [8,9°]. Leukocytes are highly deformable and, because of their lack of stable focal contacts, move at high velocities (2–30 µm/min) [10]. Shape change is generated by cortical filamentous actin, which mediates cell dynamics as well as providing stiffness to the cell body, but mature focal contacts and stress fibers are lacking [7,11]. T lymphocytes and other leukocytes use proteaseindependent physical mechanisms to overcome matrix barriers, including adaptation of the cell shape to preformed matrix structures (contact guidance), extension of lateral footholds ('elbowing') [12] and squeezing through narrow spaces (constriction rings) [11]. Such shape-driven migration allows cells to glide through or circumnavigate, rather than degrade, ECM barriers [13]. The ellipsoid cell shape in amoeboid migration requires actin polymerization along the plasma membrane to stiffen and contract the cell cortex. These cortical actin dynamics are critically controlled by the small GTPase RhoA and its effector ROCK to generate cortical tension, stiffness and the maintenance of roundish cell morphology [14,15]. On the other hand, the small GTPases Rac and Cdc42 engage adaptor proteins (i.e. the WASP) that favor focalized actin assembly and generate dynamic cell protusions, such as pseudopodia, lamellipodia, and filopodia, thereby supporting cell polarization and elongation [16]. In contrast to the five-step migration paradigm, focal contacts and focalization of proteolysis are thus eliminated in amoeboid movement, whereas fast and non-focalized receptor assemblies at cell-matrix interactions are retained.

Mesenchymal migration

In contrast to cells that use amoeboid migration, mesenchymal cells accomplish the complete five-step migration sequence (Figure 1b). In 3D tissues, mesenchymal cells adopt a spindle-shaped, fibroblast-like morphology, as characteristic for fibroblasts, myoblasts, single endothelial cells or sarcoma cells [17,18**]. The elongated morphology is dependent on integrin-mediated adhesion dynamics and the presence of high traction forces on both cell poles [18**,19]. Blocking of integrins in spindleshaped fibroblasts, endothelial cells or tumor cells by antibody or small-molecule inhibitors causes cell retraction, acquisition of spherical shape and impaired migration rates [20,21] (N Daryab and P Friedl, unpublished). Concomitant to integrin and actin focalization at substrate-binding sites, mesenchymal cells recruit surface proteases to digest and remodel ECM [22**]. B1 and β3 integrins, MT-MMPs and other proteases then colocalize at contact regions to ECM fibers, proteolytically cleave ECM molecules near to the cell surface, and generate structural matrix defects along cell migration tracks [22°°,23]. Focal contact formation and turnover occur in the timescale of 10-120 minutes, resulting in relatively slow migration velocities (0.1–2 μm/min) in 3D models [10,19,24,25]. If other cells follow along the newly generated matrix defect, a moving cell chain evolves and is guided by matrix strands (contact guidance) (Figure 1c).

Figure 1

	Integrins	Proteases	Cadherins
(a) Amoeboid	-/+	-/+	-
(b) Mesenchymal (single cells)	+	+	-
(c) Mesenchymal (chains)	+	+	-/+
(d) Clusters/cohorts	++	++	++
(e) Multicellular sheets/strands	++	++	++
(f) Branching morphogenesis	++	++	++
(g) Vascular sprouts EC PC	++	++	++
 Integrins (β1/β3) in focal contacts Non-clustered integrins Proteases (MMPs, MT-MMPs, uPA) Cadherins (E-,N-, VE-) Basement membrane			
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Prespecified diversity in cell migration strategies. In 3D tissue environments, different cell types exhibit either individual (a-c) or collective (d-g) migration mechanisms to overcome and integrate into tissue scaffolds. (a) Amoeboid migrating cells develop a dynamic leading edge rich in small pseudopodia, a roundish or ellipsoid main cell body and a trailing small uropod. (b) Mesenchymal cells retain an adhesive, tissue-dependent phenotype and develop spindle-shaped elongation in 3D ECM (i.e. the equivalent to cell spreading on 2D substrate). For movement through 3D tissues, they form focal contacts containing focalized actin and adaptor proteins [25], whereas stress fibers are preferentially present in sessile state and contraction of anchored ECM. In the detachment zone, matrix defects and the deposition of cell surface determinants including cell fragments are seen. (c) Proteolytic migration tracks can be used by neighboring cells to arrange and migrate in a loose chain-like fashion. Such files are seen in neural crest cells upon morphogenesis [63] as well as in stromal tumors extending into connective tissue in a chain like pattern, such as melanoma (C Mayer, P Friedl, unpublished). (d) In primary cancer lesions, as detected by histopathology, detached clusters of cells are disseminated throughout the adjacent connective tissue. Leading cells provide the migratory traction and, via cell-cell junctions, pull the following group forward. (e) If an invading cancer collective maintains contact with the origin, for example via proliferation of cells in the inner mass, extended non-polarized collective In migrating mesenchymal cells, such as fibroblasts, Rac and Cdc42 generate pseudopod and lamellipod dynamics at outward edges, favoring a rapid and dynamic type of \beta1 integrin engagement towards 2D and 3D substrata [26]. Interfering with Rac and Cdc42 activity perturbs cell extension and polarized force generation, thereby severely impairing migration [26]. Rho, on the other hand, stabilizes initial integrin-substrate linkages, increases focal contact size and strength, and further thickens actin filaments through several mechanisms [27,28]. In mesenchymal or adhesive cells interacting with 2D substrata, active Rho leads to increased adhesiveness, stress fiber formation, and retardation of migration speed [27,28]; for cell migration within 3D substrata, the functions of Rho with regard to cell shape and adhesion dynamics appear to be more complex (see below). Together, the coordination and synergy between polarized cytoskeletal dynamics at the leading edge, as mediated by Rac and Cdc42, and the somewhat opposing effect of Rho-mediated adhesionstrengthening and cell contractility is thought to play a key role in adhesion-dependent cell migration and related dynamics in cell morphology [26–28].

Collective migration modes

In collective migration, cells maintain their cell-cell iunctions and move as multicellular connected strands or chords into tissues; examples include invading epithelial strands or tubes, vascular sprouts and tumor clusters (Figure 1d–g). Specialized forms of collective migration are the horizontal migration of epithelial cell sheets across 2D substrates, as observed in the gut mucosa epithelium upon self-renewal or in keratinocytes migrating across provisional wound matrix, as well as in slow border cells migrating through the cell-rich scaffold of the developing ovary in Drosophila [29]. Collective migration is seen experimentally in 2D in vitro wound scratch assays [28] and in 3D ECM cultures [30]. The leading edge of a moving cell group, both in 2D and 3D migration models, is formed by one or several cells that utilize actinmediated ruffles and integrin-dependent traction to execute steps 1-4 of the migration cycle [31**,32]. In cell cultures, migrating cancer cell collectives develop preferential integrin and protease (MT1-MMP, MMP-2) engagement in a subset of cells at the leading edge [33]. The junctions within invading collectives are stabilized by cadherins, members of the immunoglobulin superfamily (e.g. NCAM or activated leukocyte adhesion molecule) [34-36] and gap-junctional cell-cell communication [37,38]. The rear of the leading cell(s) maintain

the adhesive interaction with other cells, so the retraction of the trailing edge has an important modification: as it glides along the ECM structure, neighboring cells are dragged forward along the established migration track by means of cell-cell adhesion [30,34,39]. While the leading cells generate actin- and integrin-mediated traction, a linear cortical actin network extends along cell-cell junctions into deeper regions of the collective, suggesting that cortical actin plays a role in sustaining collective integrity [31°,32,40]. In de-differentiated tumor cell groups, rather amorphous cell strands and masses that lack an inner lumen extend within the tissue (Figures 1d,e). By contrast, non-neoplastic developing glandular ducts (Figure 1f) and blood vessels (Figure 1g) contain polarized cells that form an inner lumen and newly produce a surrounding basement membrane [41,42]. Sprouting blood vessels, unlike other groups of cells undergoing collective migration, use cadherins to recruit pericytes as a second cell type; these pericytes then participate in the *de novo* synthesis of an encircling basement membrane [43,44].

A special and more complex example of collective migration is the 'mass' movement in morphogenesis, as seen during the convergent extension of the vertebrate embryo [45] or the closure of the dorsal surface in the *Drosophila* embryo [46]. In both cases, movement is carried out by complex multicellular sheets that contain cells linked to each other by cell-cell junctions and other forms of cellcell communication; the cells move along the underlying or surrounding tissue substrate to form epithelia or organs [45,46]. Besides the well-studied mechanisms of 'active' collective migration, morphogenic movements additionally comprise biophysically barely understood 'supracellular' forces that go beyond single cell dynamics to change the shape and position of cells and entire tissues, for example by folding cell and matrix sheets or extending and contracting entire body regions [46].

Plasticity in cell migration

While it is generally assumed that differentiated cells retain their migration mode once they have acquired it, recent data suggest that the gain or loss of generically prespecified components in the migration cycle can cause an adaptive switch in migration pattern and mechanism. This phenomenon is here termed plasticity or transition.

Physiological transitions

The process by which circulating precursor cells acquire a differentiated phenotype in the peripheral organs follows

(Figure 1 Legend Continued) strands or sheets extend into the stroma. (f) Non-neoplastic collective migration is seen in budding epithelial ducts and glands, termed branching morphogenesis. While cells of the leading edge engage in cell-ECM interactions and proteolytic ECM remodeling, the emerging matrix defect is collectively filled by following cells. The collective then generates a basement membrane along the outside tissue interface while an inner lumen is formed. (g) Collective sprouts of endothelial cells form new blood vessels by moving and maintaining cell-cell junctions. Guided by one 'pathfinder' cell, the chain matures into a growing strand containing a lumen. While the strand moves forward, pericytes are recruited and engaged by cadherins. Endothelial cells and pericytes then produce a basement membrane. EC, endothelial cell; PC, pericyte. Arrows indicate the direction of migration. For the functions of integrins, proteases and cadherins, please refer to the main text.

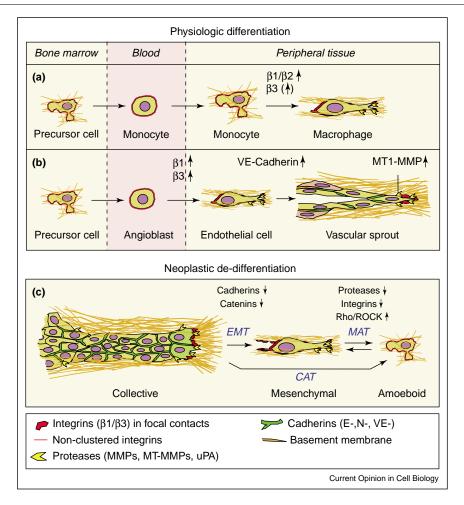
a program of genetic prespecification that involves position change as well as alterations in the cells' molecular repertoire.

Macrophage development

An amoeboid stem cell (hemangioblast) in the bone marrow differentiates into a monocyte which, after exit from the bone marrow, blood passage and emigration into the peripheral tissue, becomes a resident macrophage (Figure 2a). Migrating monocytes express low levels of β1 or β3 integrins on their surface and use amoeboid

shape change for their migration (see movies 4 and 5 at http://www.bloodjournal.org/cgi/content/full/2002-12-3791/DC1). Macrophages are stromal-type elongated cells that show strongly upregulated \(\beta \)1 and \(\beta \)3 integrins [47,48], focalized integrin-ligand interactions and an actin cytoskeleton that is rearranged from diffuse to strongly focalized [49]. The macrophage adopts a spindle-shaped yet mobile stromal phenotype and, upon granulomatous tissue reaction and cell fusion with other macrophages, can create multinucleated giant cells [49]. Although a complete dynamic mapping of these transition stages

Figure 2



Transition of cell migration in physiologic differentiation and neoplastic de-differentiation. Migratory plasticity in non-neoplastic differentiation occurs via the gain of adhesion molecules and proteases and of cell-cell interactions. (a) The development of a tissue macrophage requires the amoeboid stem cells in the bone marrow to differentiate sequentially into an amoeboid myeloic precursor cell and then a monocyte, which emigrates from the bone marrow. After circulation in the blood stream, the monocyte is recruited into the tissue, upregulates β1, β2 and, to some extent, β3 integrins and develops into a slowly moving macrophage of spindle-shaped morphology. (b) For formation of new vessel sprouts, amoeboid angioblasts are recruited from the bone marrow into the tissue to undergo differentiation into single endothelial cells (ECs) of mesenchymal phenotype. ECs interact with neighbor ECs via VE-cadherin and other adhesion receptors forming a multicellular sprout. (c) The reverse processes take place in neoplastic de-differentiation. After the loss of cell-cell junctions, for example through reduced expression or cleavage of cadherins, individual mesenchymal cells detach from the collective and migrate via integrin and protease usage (epithelial-mesenchymal transition, EMT). If integrin or protease functions are weakened in mesenchymal cells, as well as by an increase in ROCK activity, the transition towards amoeboid movement (mesenchymal-amoeboid transition, MAT) occurs. Collective-amoeboid transition (CAT) results from the dissociation of multicellular collectives into amoeboid single cells, as seen after inhibition of $\beta 1$ integrins in melanoma explants.

is presently lacking, macrophage development represents the transition from amoeboid to mesenchymal characteristics.

Endothelial cell differentiation

Endothelial precursor cells (hemangioblasts) undergo a transition from an amoeboid phenotype through a transient mesenchymal stage to a terminal collective pattern (Figure 2b). Hemangioblasts leave the bone marrow as amoeboid cells characterized by a roundish appearance on adhesive substrates such as fibronectin, low surface levels of β1 and αv integrins, and an exclusively cortical actin cytoskeleton lacking stress fibers [50,51°] (J Varner, personal communication). At the extravasation site, the interaction of circulating angioblasts with blood vessels requires, like the transendothelial migration of leukocytes, the function of integrin α4β1 (Varner J, personal communication). Once in contact with the tissue ECM, a differentiation program towards a mesenchymal adhesive phenotype occurs, consisting of increased attachment and spreading, acquisition of stress fibers, and the strong upregulation of β 1, $\alpha v\beta$ 3 and $\alpha v\beta$ 5 integrins, VCAM, and endothelial differentiation markers [51°] (J Varner, personal communication). As the final differentiation step, the gain of cell-cell junctions and the formation of multicellular sprouts and tubes results in a collective phenotype of initially primitive — later matured vessels [52,53]. The tip of the sprout is formed by a pathfinder cell using β1 and αvβ3 integrins and MT1-MMP for collagen remodeling [54]. In all these steps, cell movement is essential for appropriate endothelial cell positioning and assembly into multicellular vessels.

Other examples of plasticity of cell shape and migration mode might include myotube formation (mesenchymal) from myogenic precursor cells (amoeboid) and fibroblast differentiation (mesenchymal) from recirculating stem cells (amoeboid) in regenerating wounds.

What happens now, if characteristics of cell differentiation are lost, as seen in neoplastic cell dedifferentiation?

Reverse transitions

Tumor cell migration models have provided useful insights into the transition stages and underlying mechanisms that allow the conversion from differentiated and collective migration towards more simple single-cell movements, here referred to as 'reverse transition'.

Epithelial-mesenchymal transition

During progressive de-differentiation in epithelial cancer, the conversion from multicellular growth and invasion to mesenchymal single cell migration is termed the epithelial-mesenchymal transition (EMT) (Figure 2c) (see [55] and references therein). The primary step is the loss of cell-cell junctions via several mechanisms. These include reduced cadherin expression, loss-of-function

mutations in cadherin and catenin signaling pathways, and deregulated function of proteases leading to degradation of cadherins and other cell-cell adhesion molecules [56]. Because integrin and protease functions and promigratory signaling remain intact, detaching cells adopt a mesenchymal type of single-cell migration [55,57]. As represented by the transition of highly differentiated epithelial cancer towards a sarcomatous stromal phenotype, EMT is considered as a key event in tumor progression [55].

Mesenchymal-amoeboid transition

If mechanical or signaling pathways that stabilize cell-ECM interactions are weakened, mesenchymal movement can convert towards amoeboid migration (Figures 2c and 3). Known mechanisms leading to mesenchymalamoeboid transition (MAT) are the abrogation of pericellular proteolysis using protease inhibitors, the strengthening of RHO/ROCK signal pathways, and the weakening of integrin-ECM interactions by antagonists (Figure 2c).

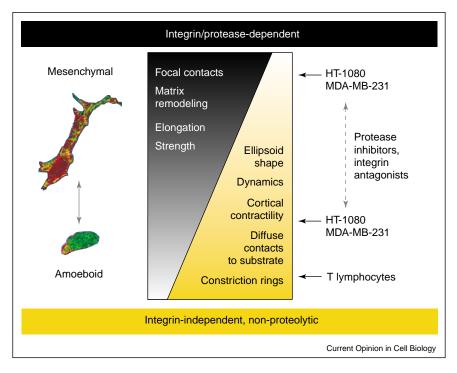
Loss of protease function

Mesenchymally migrating tumor cells such as HT1080 fibrosarcoma and MDA-MB-231 mammary carcinoma cells cease their proteolytic migration after the addition of protease inhibitors that target MMPs, ADAMs, cathepsins and serine/threonine proteases [22**]. Instead of widening the pericellular space and cleaving ECM fibers, the cells then switch to amoeboid behavior involving vigorous shape change and the ability to squeeze through narrow regions, thereby rescuing their migration independently of pericellular proteolysis [22.]. Consistent with an amoeboid phenotype, both β1 integrin distribution and filamenous actin adopt a diffuse cortical pattern, reminiscent of migrating lymphocytes [13]. MAT after inhibition of surface proteases was confirmed in vivo for tumor cells injected into the mouse dermis [22**]. A similar phenotypic change was obtained in fibroblasts populating excisional wounds in rats treated with the broad-spectrum MMP inhibitor GM6001 (Agren MS, personal communication). Despite MMP inhibition, these fibroblasts retain the capacity to infiltrate the fibrin-rich wound matrix; however they now exhibit roundish amoeboid morphology coupled to a strongly reduced capacity to remodel the wound and generate scar contraction (Agren MS, personal communication). How protease inhibitors interfere with the regulation of integrin and cytoskeletal dynamics and thereby reprogram mesenchymal cells towards amoeboid movement remains unknown.

Activation of ROCK

Similar transition from mesenchymal to amoeboid movement occurs in HT1080 cells that penetrate thick 3D matrigel layers after constitutive activation of the Rhoeffector kinase ROCK [58**]. ROCK acts by increasing

Figure 3



Shifting adhesion and matrix degradation: the Ying and Yang of single cell migration. Migrating mesenchymal HT1080 fibrosarcoma cells and MDB-MB-231 breast cancer cells are spindle-shaped cells that use integrins and proteases for adhesive and proteolytic interactions to ECM substrate, respectively [22**]. On the other hand, T lymphocytes maintain migration by integrin-independent mechanisms and further do not generate pericellular proteolysis [10.11]. Between both extremes, cells may also fulfill the criteria for mixed phenotypes. Abrogating integrin and protease function can result in the reversible reprogramming of the migration type and the conversion from mesenchymal to amoeboid, as seen in HT-1080 and MDA-MB-231 cells [22**].

myosin-II-mediated actin filament stabilization and contraction [28,59]. Active RhoA is required for diffuse cortical actin polymerization and cell retraction in dividing cells [60] and overexpression of constitutively active ROCK causes cortical contraction and cell rounding in originally mesenchymal cell lines, such as 3T3 fibroblasts and HT1080 cells [58**,61]. Active ROCK not only causes HT1080 cells to loose their mesenchymal characteristics and convert to a roundish, contracted shape: driven by small filopodia and blebbing-type cell protrusions that contain cortical actin, the cells convert to a protease-independent migration type, reminiscent of amoeboid movement [58**]. Because ROCK activation generates cortical actin and cell rounding in some cells [15,58°,62], yet stabilizes stress fibers and causes cell spreading, flattening, and migratory cell arrest in other cell types [28], it appears that additional endogenous or environmental cofactors determine if cell-substrate adhesions are stabilized or weakened by Rho/ROCK-mediated signals.

Downregulating integrin functions

Reducing attachment forces without interfering with cell contractility prompts cell rounding and transition towards amoeboid movement. In 3D collagen substrate, such approaches include the following: selection for low endo-

genous β1 integrin expression by FACS sorting; titrating β1-integrin-mediated adhesion downwards by blocking antibody; interfering with the cytoplasmic β1 integrin domain by a dominant-negative peptidomimetic; and abrogating integrin expression by knockout strategies. All these procedures uniformly generate a roundish yet dynamic cell shape, an even surface distribution of (residual) \$1 integrins, and a non-focalized cortical layer of filamentous actin, supporting amoeboid migration in 3D collagen lattices (N Daryab, C Brakebusch and P Friedl, unpublished).

Together, these findings indicate that cells can use a spectrum of migration modes ranging from adhesive to nonadhesive and from proteolytic to protease-independent (Figure 3) [13].

Collective-amoeboid transition

In analogy to EMT, the transition from collective invasion to amoeboid single-cell crawling is obtained if cellcell and cell-ECM interactions are simultaneously weakened (Figure 2c). In multicellular clusters emigrating from melanoma explants, the inhibition of β1 integrins by adhesion-perturbing antibody abolishes collective movement yet induces the detachment of individual amoeboid cells from the collective [31.1]. In the continuous presence of antibody, detached cells utilized amoeboid shape change and constriction rings and developed a nonclustered, diffuse distribution of surface β1 integrins, similar to that observed in migrating lymphocytes [31^{••}]. Although the molecular mechanism for dispersing cell collectives towards amoeboid single cells through the \beta1-integrin pathway requires further investigation, these observations suggest that collective migration can directly, or indirectly via a mesenchymal migration step, give raise to amoeboid single cell dissemination.

Conclusions

Taking these findings together, the conversion from multicellular to mesenchymal and/or amoeboid migration may represent stereotypic processes that can be achieved by diverse mechanisms. Although amoeboid movement is detected in cells of low differentiation state, such as stem cells and leukemia and cancer cells, it is also a feature of certain non-neoplastic differentiated cells, such as T lym-

Box 2 Viewpoint: essential and optional components in cell migration

For practical reasons, it will be important to distinguish between cellular and molecular mechanisms that are essential for cell movement and those that fulfill optional supportive or modifying functions in determining migratory shapes and patterns and are dispensable for migration after functional ablation.

Essential functions

For movement of virtually all cell types on 2D as well as in 3D ECM, essential functions include the acquisition of cell polarity, pseudopod protrusion, cortical stiffness, cytoskeletal contractility and dynamics, as well as physical (but not necessarily adhesive) contact with ECM environments. The capacity to change shape, the key feature of cell movement, appears to require actin polymerization and depolymerization and the function of ROCK in controlling myosin II activity. Consequently, inhibition of actin polymerization by cytocalasin D or latrunculin and/or interference with myosin-IImediated sliding of actin filaments, for example by inhibiting ROCK [28,58**,64,65], causes profound impairment in amoeboid and other migration modes.

Optional factors

Optional factors determine a particular migration phenotype, but do not limit the capacity to migrate if cellular and/or molecular compensation mechanisms rescue motility. Some of these optional factors have previously gained major attention on the basis of their impact in 2D migration models, although in more complex 3D tissues or in vivo models migration may be sustained by compensation strategies. Candidate pathways include:

- Integrin-mediated attachment and traction because loss of attachment might be compensated by less stringent cell-matrix interactions and shape change
- Proteolytic degradation of the ECM its loss can be compensated by physical mechanisms, such as shape change, cell squeezing and seeking for trails of least resistance
- Cell-cell adhesion because cell-cell junctions favor collective cell dynamics and migration while the loss of cell-cell contacts will prompt single cell migration
- Activating chemotactic factors and cytokines virtually all cell types retain autochthonous cytoskeletal oscillation, polarity and random motility in the absence of migration-promoting factors

phocytes, monocytes and neutrophils [11]. In higher mammals (Box 1) amoeboid migration is hence likely to embody a physically optimized migration mode that allows easy cell traffic towards and between structurally different tissue compartments. Although cell differentiation and tissue segregation lead to prespecified migration patterns that are characteristic for each cell type, a basic capacity to revert to ultimately amoeboid migration appears to be retained in some, if not many, mammalian cell types.

Molecular targeting of specific pathways may thus yield more complex adaptation reactions than previously appreciated. Pharmacological interference with adhesive, proteolytic and signaling pathways that maintain a differentiated tissue phenotype but are not essential for cell movement may favor cell reversion to a less differentiated behavior and dissemination mode (Box 2). On the basis of these concepts, the efficiency of drugs designed to interfere with, or redirect, different sorts of migration processes will depend on the target cell's capacity to compensate its migration ability by cellular and molecular adaptation. Consequently, an understanding of the mechanisms that sustain amoeboid rescue mechanisms in higher eukaryotes as well as in *Dictyostelium* will be required for a better targeting of migration processes.

Acknowledgements

I am very grateful to Katarina Wolf and Anthea Messent for helpful suggestions on the manuscript. I further acknowledge Eva-B. Bröcker for continuous support and insightful discussions. The original work of the author's laboratory was supported by grants from the DFG (FR1155/2-3 and FR1155/5-1), Ministry for Science and Education, the Wilhelm-Sander-Foundation and the Interdisciplinary Center for Clinical Research (IZKF), Würzburg.

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