

Chemotaxis: signalling modules join hands at front and tail

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Chemotaxis is the result of a refined interplay among various intracellular molecules that process spatial and temporal information. Here we present a modular scheme of the complex interactions between the front and the back of cells that allows them to navigate. First, at the front of the cell, activated Rho-type GTPases induce actin polymerization and pseudopod formation. Second, phosphatidylinositol-3,4,5-trisphosphate (PtdIns(3,4,5)P₃) is produced in a patch at the leading edge, where it binds pleckstrin-homology-domain-containing proteins, which enhance actin polymerization and translocation of the pseudopod. Third, in *Dictyostelium* amoebae, a cyclic-GMP-signalling cascade has been identified that regulates myosin filament formation in the posterior of the cell, thereby inhibiting the formation of lateral pseudopodia that could misdirect the cell.

Keywords: actin; myosin; cGMP; PtdIns(3,4,5)P₃; modules

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Introduction

Many cells sense the presence of extracellular chemical signals and guide their movement in the direction of the concentration gradient. This process, called chemotaxis, has a role in various functions such as finding food in prokaryotes, forming multicellular structures in protozoans, tracing bacterial infections by leukocytes and embryogenesis in metazoans (Baggiolini, 1998; Campbell & Butcher, 2000; Crone & Lee, 2002; Iijima *et al*, 2002). Cells that are exposed to chemoattractants can use temporal and spatial aspects of the chemoattractant gradient to orientate their movement. Prokaryotes use only the temporal component of the chemoattractant gradient, probably because, at 1–2 µm, they are too small to process spatial information. These cells exhibit a random walk, with movement steps in all directions. Cells frequently tumble and move in a new random direction. Tumbling frequency is decreased and movement in one direction is prolonged when cells experience an increase in chemoattractant concentration, that is, when moving up the gradient (Berg, 1988;

Bourret & Stock, 2002). Eukaryotic chemotactic cells are about 10–20 µm in diameter, which allows them to process both spatial and temporal information: they measure the difference in chemoattractant concentration between the ends of the cell, and then move up the gradient of chemoattractant (Zigmond, 1977). Under starvation conditions, *Dictyostelium* cells emit the chemoattractant cyclic AMP, which causes them to aggregate and eventually form fruiting bodies. *Dictyostelium* cells are able to respond to a 2% difference in cAMP concentration between the front and the back of the cell, at a mean cAMP concentration that occupies only about 1% of its 40,000 cAMP receptors (Mato *et al*, 1975). The ability to sense and respond to shallow gradients of extracellular signals is remarkably similar in simple amoebae such as *Dictyostelium* and in mammalian leukocytes.

Here, we present data suggesting that chemotaxis is regulated by the interplay of at least three signalling pathways. These signalling pathways are presumably intricately connected to the basal mechanism of pseudopod formation (Pollard & Borisy, 2003). In the absence of chemoattractants, cells extend pseudopodia in a more or less random direction, probably as self-organizing structures (Bourne & Weiner, 2002). In the presence of a chemotactic gradient, this basic mechanism of pseudopod formation is biased to produce more pseudopodia at the front of the cell, as defined by the site of the highest chemoattractant concentration. This means that, during chemotaxis, cells become functionally polarized. In *Dictyostelium* cells that have been starved for 4 h, polarity is a rather transient state, because cells quickly extend a pseudopod at the back of the cell on gradient reversal (Swanson & Taylor, 1982; Chen *et al*, 2003; Kriebel *et al*, 2003). The polarity of neutrophils is more persistent, because gradient reversal usually leads to the formation of additional pseudopodia at the front by which cells make U-turns (Zigmond, 1977). Such a rigid polarization might improve chemotaxis by enhancing the persistence of movement in the gradient, but it also makes the cells less able to respond to changes in gradient direction. Interestingly, *Dictyostelium* cells starved for 7 h are more polar than those starved for a shorter period. At the onset of *Dictyostelium* cell aggregation, many potential aggregation centres emit cAMP waves, and cells experience successive cAMP gradients from different directions. After 7 h, the aggregation centres are fixed and the chemoattractant gradients become stable.

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Diffusion of intracellular messengers

During chemotaxis, pseudopodia are extended at the leading edge and retracted at the back of the cell. These processes are regulated by intracellular second messengers that are produced on the activation of chemoattractant receptors, and whose diffusion should meet two contradictory requirements. To establish and preserve a front-to-tail polarity, intracellular signalling molecules must not be allowed to diffuse very far into the cell. Conversely, the ability of the cells to detect shallow gradients requires the local integration of information from several occupied receptors, which implies that signalling molecules must diffuse over some distance. The complex mathematics of molecular diffusion can be reduced to a simple parameter, dispersion length, $L_d = \sqrt{D\tau}$, which describes the average distance that a molecule with diffusion coefficient D travels from its source before it is degraded after a life-time (the life-time is the reciprocal of the degradation rate constant k ; that is, $1/k$; Postma & Van Haastert, 2001). A molecule will not move far from its source if it diffuses very slowly and/or is degraded very quickly. Soluble second messengers such as cGMP diffuse relatively quickly, so they cannot preserve spatial information unless their diffusion is restricted or their degradation is extremely fast (a life-time of less than 10 ms). With an observed life-time of about 10 s (see Valkema & Van Haastert, 1994), the dispersion length of cGMP in *Dictyostelium* can be estimated to be about 55 μm , which is several times the length of the cell (Table 1). Although such molecules cannot preserve spatial information in a 10- μm cell, they are well suited to provide temporal and global intracellular communication. Conversely, transmembrane proteins such as receptors diffuse extremely slowly, so that ligand-occupied receptors fully retain spatial information. However, with molecules whose diffusion is so sluggish, communication between the front and the back of cells would require hours. For example, the *Dictyostelium* cAMP-receptor complex has a life-time of about 1.4 s. In this short period, the cAMP-receptor complex diffuses about 0.2 μm , which is only as far as the closest

receptor. Thus, receptors cannot even integrate spatial information locally. By contrast, membrane-localized lipids such as phosphatidylinositol-3,4,5-trisphosphate (PtdIns(3,4,5)P₃), with a dispersion length of about 2 μm diffuse sufficiently slowly to preserve spatial information, but fast enough to allow the integration of information at the front of the cell.

Heterotrimeric and small GTP-binding proteins form another interesting group of signal transducers, because they can either diffuse in the cytosol (such as many activated G protein α -subunits) or localize to the plasma membrane through lipid anchors (such as some G protein α -subunits, most G protein β -subunits and many small G proteins). In the cytosol, the diffusion of these proteins is rather fast, whereas when they are anchored to the plasma membrane, their diffusion is about 100-fold slower. In view of the average life-time of the active GTP-bound state of G proteins, it is expected that a soluble G protein has a dispersion length of about 10–20 μm , whereas a membrane-anchored G protein (or its subunits) will disperse about 1.5 μm .

Thus, soluble second messengers and small proteins are well suited to the transmission of temporal information through global signalling but their diffusion is too fast to retain spatial information. Transmembrane proteins, in contrast, are too sluggish. However, phospholipids and membrane lipid-anchored proteins have the required diffusion properties to preserve and integrate spatial information locally within a few micrometres, which corresponds to the size of a pseudopod.

Intracellular signalling at the front of the cell

Several excitation–inhibition models for chemotaxis have been proposed to explain the conversion of a shallow gradient of chemoattractant to activation at the front of the cell. These models all use a local activator and a global inhibitor in some way (Haugh & Lauffenburger, 1997; Parent & Devreotes, 1999; Meinhardt & Gierer, 2000; Postma & Van Haastert, 2001; Iijima *et al.*, 2002; Levchenko & Iglesias, 2002). However, the primary intracellular activator has not been fully identified. One of the proposed candidates is PtdIns(3,4,5)P₃. During chemotaxis, several proteins that contain pleckstrin homology (PH) domains translocate from the cytosol to the plasma membrane at the leading edge of both *Dictyostelium* cells and leukocytes (Parent *et al.*, 1998; Meili *et al.*, 1999; Servant *et al.*, 2000). These proteins include the cytosolic regulator of adenylyl cyclase (Crac) and protein kinase B (Akt/PKB). Several indications suggest that, *in vivo*, the PH-domain-containing proteins bind to PtdIns(3,4,5)P₃ but not to other products of the phosphatidylinositol-3-OH kinase (PI(3)K) pathway such as PtdIns(3,4)P₂. First, chemoattractants induce the accumulation of PtdIns(3,4,5)P₃ with the appropriate kinetics in both *Dictyostelium* and neutrophils (Traynor-Kaplan *et al.*, 1989; Huang *et al.*, 2003). Second, the inactivation of the gene that encodes the phosphatase and tensin homologue (PTEN) leads to higher PtdIns(3,4,5)P₃ concentrations and enhanced chemoattractant-stimulated translocation of PH-domain-containing proteins (Iijima & Devreotes, 2002; Huang *et al.*, 2003). Third, in *Dictyostelium* the concentration of PtdIns(3,4)P₂ is very low (H.M.L., D. Blero, C. Erneux and P.J.M.V.H., unpublished observations).

PtdIns(3,4,5)P₃ is synthesized by the PI(3)Ks PI3K1 and PI3K2 in *Dictyostelium* and by PI(3)K γ in neutrophils (Zhou *et al.*, 1995; Hirsch *et al.*, 2000; Huang *et al.*, 2003), and is degraded by the 3-phosphatase PTEN (Iijima & Devreotes, 2002) and by multiple

Table 1 | Dispersion length of signalling components for *Dictyostelium* chemotaxis

Compound	Diffusion coefficient, D ($\mu\text{m}^2 \text{s}^{-1}$)	Life-time, τ (s)	Dispersion length, $L_d = \sqrt{D\tau}$ (μm)
cGMP	300	10	55
G protein in cytosol	10	20	14
PtdIns(3,4,5)P ₃	1	5	2.2
G protein in membrane	0.1	20	1.4
Receptor–cAMP complex	0.027	1.4	0.20

Some measurements of cells might help in interpreting these data. The diameter of a *Dictyostelium* cell is about 10 μm and a pseudopod has a size of 2–3 μm . A *Dictyostelium* cell contains about 40,000 cAMP receptors distributed over a surface area of about 400 μm^2 , yielding an average distance between receptors of 0.1 μm . At threshold cAMP concentrations (1 nM) only about 400 receptors are occupied with cAMP, and the average distance between occupied receptors is about 1 μm . References: diffusion of cGMP, Chen *et al.* (1999); cGMP degradation *in vivo*, Valkema & Van Haastert (1994); diffusion of phosphatidylinositol-3,4,5-trisphosphate (PtdIns(3,4,5)P₃), Almeida & Vaz (1995); degradation of PtdIns(3,4,5)P₃ *in vivo*, Huang *et al.* (2003); diffusion of G protein in cytosol as diffusion of cytosolic GFP, Potma *et al.* (2001); degradation of G α GTP from interconversions of cAMP-binding sites, Van Haastert *et al.* (1986); diffusion of G proteins in membrane with lipid anchor slower than diffusion of lipid, degradation of G proteins in membrane as degradation of soluble G-protein and diffusion of receptor–cAMP complex, Ueda *et al.* (2001); life-time of receptor–cAMP complex, Van Haastert *et al.* (1986).

5-phosphatases (Loovers *et al*, 2002). Recent studies have shown that PI3K1 and PI3K2 translocate from the cytosol to the front of *Dictyostelium* cells, whereas PTEN becomes localized to the back (Funamoto *et al*, 2002; Iijima & Devreotes, 2002). In *pten*-null cells, the absence of 3-phosphatase activity leads to an increase in the membrane area with bound PH_{Crac} -GFP (the PH domain of Crac fused to green fluorescent protein) and also results in an expansion of the region from which F-actin-filled pseudopodia are extended (Iijima & Devreotes, 2002). Thus, where $\text{PtdIns}(3,4,5)\text{P}_3$ is formed in the cell, its diffusion kinetics and its apparent ability to induce F-actin formation make $\text{PtdIns}(3,4,5)\text{P}_3$ an ideal second messenger for chemotaxis. However, the chemotactic defect of *Dictyostelium pi3k1-lpi3k2-* or mammalian *PI(3)K γ* -null cells is only partial and results in a decrease in speed rather than in poor directionality of cell movement (Fig 1). These modest defects are also observed in wild-type cells that have been treated with moderate concentrations of the PI(3)K inhibitors LY294002 and wortmannin (Funamoto *et al*, 2002; Iijima *et al*, 2002; Wang *et al*, 2002; Huang *et al*, 2003). Furthermore, it has been shown in *Dictyostelium* that the amino-terminal segment of PI(3)K expressed in *pi3k*-null cells translocates to the leading edge, suggesting that signalling molecules upstream of PI(3)K are already highly localized (Funamoto *et al*, 2002).

Another candidate for the initial intracellular activator are small G proteins. In neutrophils, it has been shown that chemoattractant-induced stimulation of Rho GTPase-family members has an essential role in cellular polarization (by means of Cdc42) and F-actin formation (by means of Rac1) (Wang *et al*, 2002; Srinivasan *et al*, 2003; Xu *et al*, 2003). Small GTP-binding proteins are positively and negatively regulated by guanine exchange factors (GEFs) and GTPase-activating proteins (GAPs), respectively (Etienne-Manneville & Hall, 2002). It is conceivable that the interplay between GEFs and GAPs causes a local accumulation of active small GTP-binding proteins at the future front of the cell, which induces actin polymerization and pseudopod formation. A similar signal transduction cascade with a putative small GTP-binding protein has been proposed for *Dictyostelium* (Chung *et al*, 2001a).

If the PI(3)K/PTEN system is not responsible for the primary processing of the chemoattractant gradient, what is the function of $\text{PtdIns}(3,4,5)\text{P}_3$ synthesis and the translocation of PH-domain-containing proteins to the leading edge? Strikingly, *Dictyostelium* cells stimulated with uniform cAMP concentrations exhibit a biphasic PH_{Crac} -GFP response: there is first a uniform translocation of PH_{Crac} -GFP to the cell boundary, which is followed by a second persistent translocation of PH_{Crac} -GFP to smaller patches (Postma *et al*, 2003). These patches of PH_{Crac} -GFP are also observed in latrunculin-treated cells (Fig 2). Latrunculin A inhibits actin filaments and leads to symmetrical spherical cells without pseudopodia (Parent *et al*, 1998), which indicates that patches of PH_{Crac} -GFP are formed in the absence of actin filaments, pseudopodia or other visible spatial cues. The patches that form under conditions of uniform cAMP distribution have essentially the same properties with respect to size, life-time and intensity as the single patch of PH_{Crac} -GFP that is formed at the leading edge of cells in a cAMP gradient (Fig 2). PH_{Crac} -GFP patches are self-organizing; that is, PH_{Crac} -GFP patches are triggered by cAMP but the cAMP concentration has no effect on the size, intensity or life-time of the patches. About 10 s after a patch is formed, a pseudopod is extended from the area of the cell with a PH_{Crac} -GFP patch.

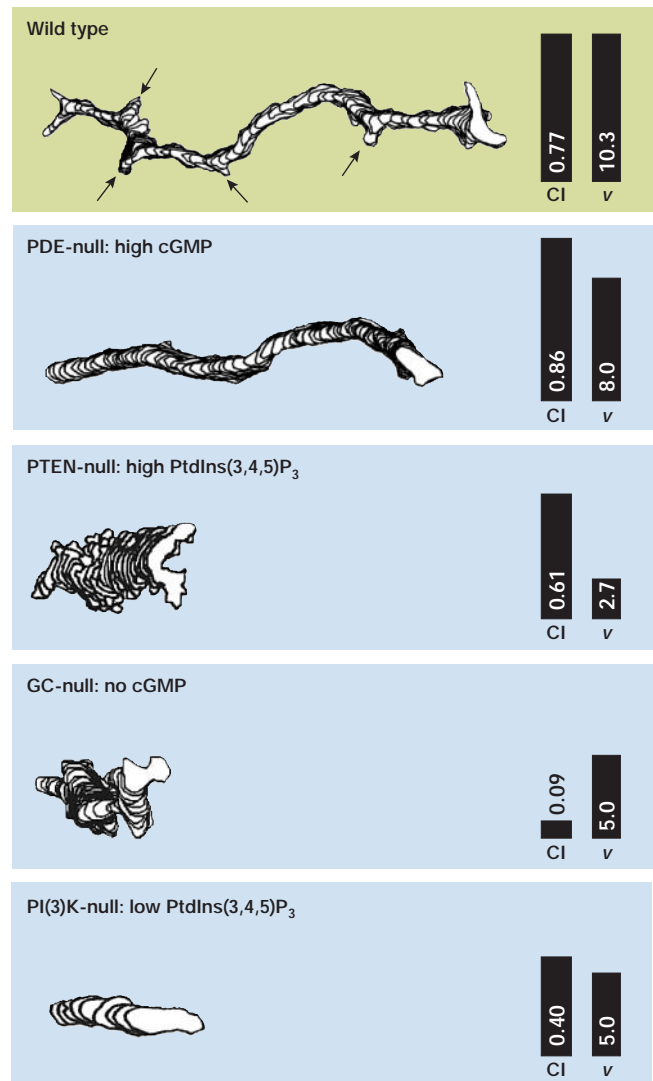


Fig 1 | Chemotaxis of *Dictyostelium* mutants. The tracks of *Dictyostelium* mutants moving for 10 min in a cAMP gradient were obtained by computer-assisted analysis. The arrows in the upper panel indicate the formation of lateral pseudopodia, which are absent from phosphodiesterase (PDE)-null cells. The numbers in each panel refer to the chemotaxis index (CI), which is the efficiency of orientating in the gradient, and the speed of movement (v , in $\mu\text{m s}^{-1}$). Data are from Bosgraaf *et al* (2002) and Iijima & Devreotes (2002), and are corrected for the differences between the wild types in these three reports. GC, guanylyl cyclase; PI(3)K, phosphatidylinositol-3-OH kinase; PTEN, phosphatase and tensin homologue.

Furthermore, pseudopodia with PH_{Crac} -GFP patches show movement steps that are about double in size those of pseudopodia without PH_{Crac} -GFP patches. In uniform cAMP, patches of PH_{Crac} -GFP are induced by very low cAMP concentrations, the response does not show perfect adaptation, and PH_{Crac} -GFP patch formation is readily reversible on the removal and readdition of cAMP (Chen *et al*, 2003; Postma *et al*, 2003; M.P., J. Roelofs, J. Goedhart, H.M.L., A.J.W.G. Visser and P.J.M.V.H., unpublished observations). We propose two functions for the $\text{PtdIns}(3,4,5)\text{P}_3$

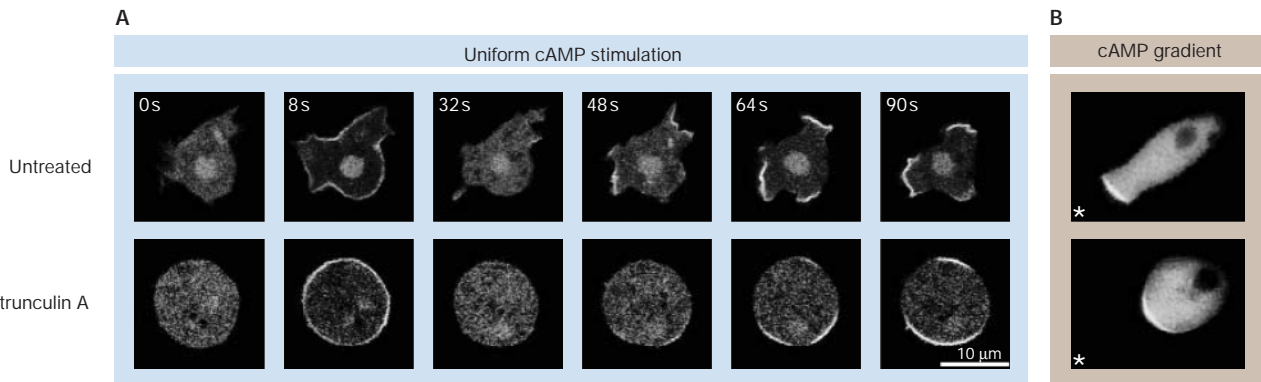


Fig 2 | Localization of phosphatidylinositol-3,4,5-trisphosphate by using PH_{Crac}-GFP. **(A)** *Dictyostelium* cells were stimulated in a perfusion chamber with a uniform cAMP concentration of 1 µM. In untreated cells, PH_{Crac}-GFP transiently translocates to the entire membrane in about 8 s, which is followed by persistent translocation of PH_{Crac}-GFP to small patches. About 10 s after a PH_{Crac}-GFP patch appears, a pseudopod is formed at the position of a PH_{Crac}-GFP patch (Postma *et al*, 2003). In the lower panels, cells were pretreated for 5 min with 1 µM latrunculin A, which induces the depolymerization of actin, the inhibition of pseudopod formation and the rounding up of the cell. A uniform cAMP concentration still induces a localized response with patches of PH_{Crac}-GFP in these spherical cells. **(B)** Cells were stimulated locally with cAMP by using a micropipette. The position of the pipette is indicated by an asterisk (data from Parent *et al*, 1998; Parent & Devreotes, 1999). The multiple patches induced by uniform cAMP have the same size and life-time as the single patch induced by the cAMP gradient, and they closely correlate with pseudopod formation. Crac, cytosolic regulator of adenyllyl cyclase; GFP, green fluorescent protein; PH, pleckstrin homology.

synthesis that occurs at these PH_{Crac}-GFP patches at the leading edge. First, the absence of adaptation implies that cells constantly monitor changes in the direction of the chemoattractant gradient that triggers new patches of PtdIns(3,4,5)P₃. Second, the induction of PtdIns(3,4,5)P₃ patches by very low cAMP concentrations and the increased movement steps of pseudopodia with patches suggest that the PI(3)K pathway increases the efficiency of the cells to detect and process shallow gradients.

Intracellular signalling at the back of the cell

The suppression of pseudopodia at the back of the cell is important for efficient chemotaxis, and this seems to be mediated by myosin filaments in the cortex at the back of the cell, both in *Dictyostelium* and in leukocytes (Stites *et al*, 1998; Eddy *et al*, 2000; Xu *et al*, 2003). Previous work in *Dictyostelium* suggests that intracellular cGMP is a good candidate for the messenger that induces myosin filament formation and the suppression of lateral pseudopodia (Liu & Newell, 1993). Recently, the genes that encode guanylyl cyclases, cGMP-phosphodiesterases and cGMP-target proteins were identified, and these have been used to resolve the function of cGMP signalling in *Dictyostelium* chemotaxis (Bosgraaf *et al*, 2002; Goldberg *et al*, 2002; Roelofs *et al*, 2002). Cells that lack two phosphodiesterases contain very high concentrations of cGMP, which are accompanied by an increase in the number of myosin filaments in the cortex and the suppression of lateral pseudopodia (Fig 1). Cells without guanylyl cyclases or without the cGMP targets show very poor chemotaxis (Bosgraaf *et al*, 2002). Although cGMP clearly mediates myosin filament formation, cGMP might have other functions that are unknown at present, because the deletion of guanylyl cyclases has a more drastic effect on chemotaxis than the deletion of myosin (Wessels *et al*, 1988; Bosgraaf *et al*, 2002). In addition, cGMP might not be the only mediator of myosin filament formation. A small residual myosin response remains detectable in guanylyl-cyclase-null cells (Bosgraaf *et al*, 2002), which might be mediated by the activation of PAKa, a member of the p21-activated kinase family, by PI(3)K (Chung &

Firtel, 1999; Chung *et al*, 2001b). In neutrophils, myosin filament formation seems to be mediated by a Rho-stimulated kinase (Xu *et al*, 2003) rather than by a cGMP-stimulated kinase. Although these pathways are different in neutrophils and in *Dictyostelium*, the end result is essentially identical: myosin filaments are formed in the back and at the sides of the cell that mediate the contraction of the back and inhibit the formation of lateral pseudopodia.

In *Dictyostelium*, chemoattractant-induced cGMP formation is tightly regulated by adaptation, such that only an increment in the chemoattractant concentration leads to a higher concentration of cGMP (Van Haastert & Van der Heijden, 1983). These temporal properties of cGMP formation could be important for chemotaxis. As long as cells move up the gradient, the cGMP concentration is higher, leading to the inhibition of pseudopodia at the back of the cell. When the chemoattractant concentration no longer changes with time, or when the gradient reverses, the cGMP concentration declines, and soon the back of the cell regains the ability to form pseudopodia. This temporal component of chemotactic signalling in *Dictyostelium* is similar to the temporal mechanism of chemotaxis in prokaryotes (Berg, 1988).

As mentioned above, it is unlikely that cGMP can form spatial gradients inside cells because a cGMP molecule can diffuse on average five cell lengths before it is degraded. What prevents the formation of myosin filaments at the front, if cGMP cannot form a gradient in the cell? Recent studies show that myosin heavy-chain kinase A (MHCK-A) translocates to the leading edge (Liang *et al*, 2002; Steimle *et al*, 2002). It is known that phosphorylated myosin is bent and does not form filaments (de la Roche & Cote, 2001), suggesting that myosin filaments at the leading edge fall apart. It seems that temporal and spatial mechanisms collaborate: cGMP induces myosin filaments globally as long as the chemoattractant concentration increases with time, whereas filaments dissociate at the front of the cell as a result of the local activity of MHCK-A. In this way, myosin filaments become restricted to the back of the cell, where they retract the uropod and suppress lateral pseudopodia.

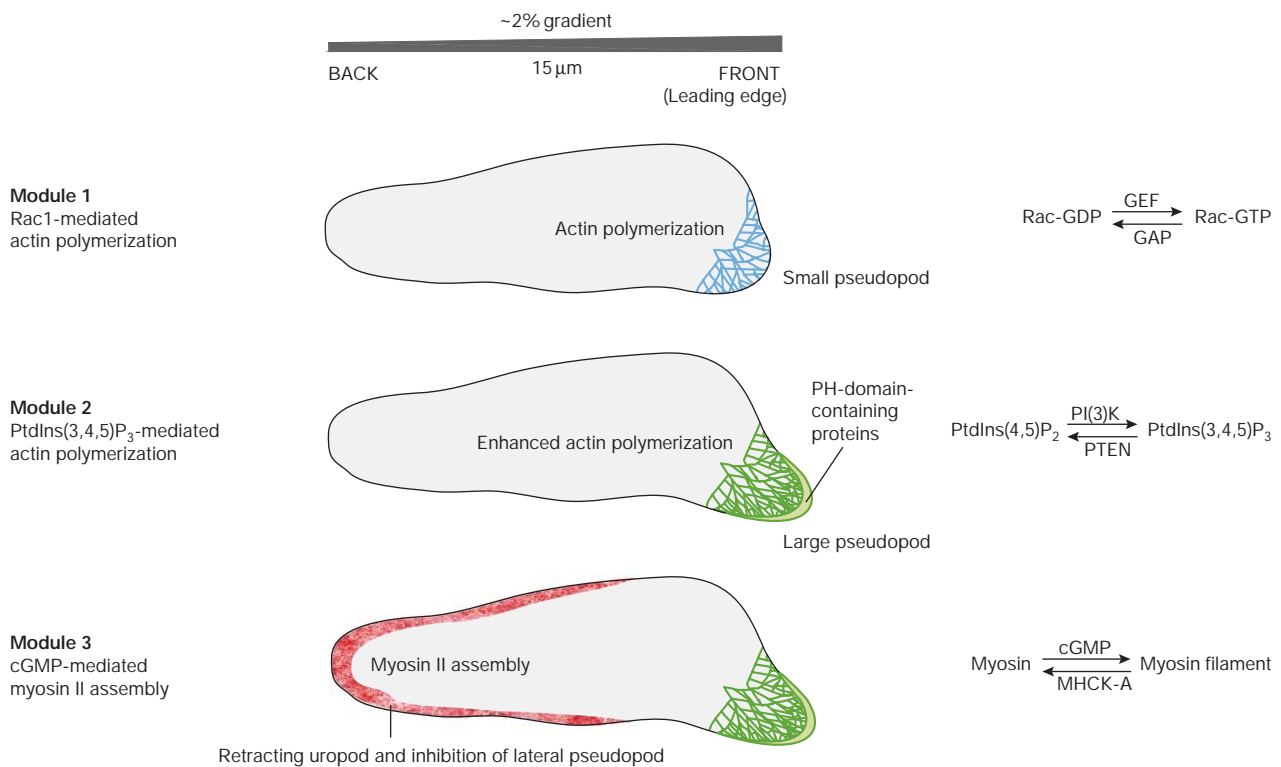


Fig 3 | Modules in chemotaxis. The first module provides the formation of active small G proteins of the Rho/Rac group, leading to actin polymerization and a small pseudopod at the front of the cell. It is proposed that the differential spatial activation of stimulatory guanine exchange factors (GEFs) and inhibitory GTPase-activating proteins (GAPs) leads to the accumulation of GTP-bound G protein at the higher chemoattractant concentrations. The polarity induced by the first module is used by the second module to trigger a patch of phosphatidylinositol-3,4,5-trisphosphate (PtdIns(3,4,5)P₃) at the leading edge, which enhances actin polymerization and extension of the pseudopod. The third module inhibits the formation of pseudopodia in the back and at the sides of the cell, and induces retraction of the uropod. An increase in cGMP concentration leads to myosin filament formation throughout the cortex, while myosin becomes phosphorylated and is depolymerized in the front of the cell by a specific kinase (myosin heavy-chain kinase A (MHCK-A)) that translocates to the leading edge. Other pathways (PAKa in *Dictyostelium* and Rho in neutrophils) might also contribute to myosin filament formation.

In conclusion, our present knowledge, which has been gleaned from different systems such as mammalian neutrophils and protozoan amoebae, suggests that gradient sensing during chemotaxis does not depend on a single molecular mechanism that informs cells where to go, but instead consists of several modules of interconnected signalling networks (Fig 3). Efficient chemotaxis is the result of a refined interplay between these modules to transmit and integrate spatial information (such as PtdIns(3,4,5)P₃) and temporal information (for example cGMP). The spatial signals are formed in a highly nonlinear way and spread relatively slowly so that high local concentrations appear in patches. Temporal signals spread swiftly and serve as global inhibitors; their formation might be subject to adaptation so that they are only generated when the stimulus concentration increases with time. These properties of chemotaxis are surprisingly similar in *Dictyostelium* and leukocytes, cells that diverged during evolution about a billion years ago. Through future research it will be exciting to uncover additional modules of regulatory interactions that add to the wonderful complexity of cellular navigation systems.

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