Biophysics/Biophysique

Chemotaxis with directional sensing during Dictyostelium aggregation

Rui Dilão a,*,b, Marcus J.B. Hauser c

a Non-Linear Dynamics Group, IST, Department of Physics, avenue Rovisco Pais, 1049-001 Lisbon, Portugal
b Institut des hautes études scientifiques, 35, route de Chartres, 91440 Bures-sur-Yvette, France
c Abteilung Biophysik, Institut für Experimentelle Physik, Otto-von-Guericke-Universität Magdeburg, Universitätsplatz 2, 39106 Magdeburg, Germany

A R T I C L E   I N F O
Article history:
Received 14 June 2013
Accepted after revision 9 October 2013
Available online 7 November 2013

Keywords:
Chemotaxis
Directional sensing
Dictyostelium
Chemotactic wave paradox

A B S T R A C T
We show that the chemotactic movements of colonies of the starving amoeba Dictyostelium discoides are driven by a force that depends on both the direction of propagation (directional sensing) of reaction-diffusion chemotactic waves and on the gradient of the concentration of the chemoattractant, solving the chemotactic wave paradox. It is shown that the directional sensing of amoebae is due to the sensitivity of the cells to the time variation of the concentration of the chemoattractant combined with its spatial gradient. It is also shown that chemotaxis exclusively driven by local concentration gradient leads to unstable local motion, preventing cells from aggregation. These findings show that the formation of mounds, which initiate multicellularity in Dictyostelium discoides, is caused by the sensitivity of the amoebae due to three factors, namely, to the direction of propagation of the chemoattractant, to its spatial gradient, and to the emergence of cAMP "emitting centres", responsible for the local accumulation of the amoebae.

© 2013 Académie des sciences. Published by Elsevier Masson SAS. All rights reserved.

1. Introduction

Chemotaxis is the phenomenon in which cells or microorganisms direct their movement as a response to the local variation of the concentration of some chemical substance. For example, in colonies of Dictyostelium discoides (Dd), localised groups of starving amoebae initiate the production of cyclic adenosine monophosphate (cAMP) that spreads in space as reaction-diffusion travelling chemical waves. In the vicinity of these spontaneously formed cAMP emission centres, the amoebae sense cAMP and direct their chemotactic movements towards the initiation centres [1]. Near these centres, amoebae rotate around a spontaneously formed hole (i.e. a zone depleted from cells), where the cell density increases locally to form aggregates (Fig. 1). At a later stage of development, aggregates are transformed into a simple multicellular organism with two main kinds of differentiated cells [2]. This is one of the simplest known mechanisms of transition from colonies of unicellular to multicellular organisms. For a detailed description of Dictyostelium aggregation and open questions on the involved biochemical mechanisms, see [2–4].

Experimental observations show that Dd cells move towards a region where cAMP chemical waves are produced, and the speed of cells is proportional to the slope or gradient of the concentration of cAMP [1,5]. On the other hand, it has been observed that cells move as long as the gradient is positive, when measured along the direction of the wave source [2,6]. “When the slope reverses, Dd cells stop moving and await for the next wave.” [2, p. 101] This fact shows that the chemotactic cell response cannot solely be determined by the local cAMP gradient [7].
Here, we investigate whether the force that drives the chemotaxis of Dd cells is solely proportional to the gradient of the chemotactic substance or whereas it also depends on the direction of propagation (directional sensing) of the time varying chemotactic signal.

If the chemotactic substance is produced at a localized source and disperses monotonically along space, the question of directional sensing is meaningless. However, if the chemoattractant propagates along space as periodically repeating wave, the hypothesis that the chemotactic motion of cells is driven by a local concentration gradient implies that the existence of a periodic variation in time of the direction of motion of the cells. This simple gradient hypothesis induces motion of the amoebae in the direction opposed to the source of the chemotactic signal, leading, asymptotically in time, to dispersive motion, preventing the amoebae from aggregation (Appendix A). This oscillatory behaviour in the aggregation pattern of Dictyostelium discoideum has never been observed and this effect is called the chemotactic wave paradox [7,8].

In the experimentally oriented literature, it is implicitly assumed that Dd cells are sensitive to the direction of propagation of the cAMP wave [9–12]. On the other hand, in simulation studies, it is generally assumed that cell chemotaxis is only driven by the sensitivity of the cells to the gradient of the chemotactic substance [13–17], and directional sensing is not included in models. Early studies have shown that the amoebae motility is also sensitive to temporal gradients [18]. However, sensitivity to temporal gradients have never been considered in mathematical models of Dictyostelium aggregation. One of the results we derive here (Appendix A) is that the introduction of directional sensing leads to the sensitivity to the temporal gradient of the chemoattractant.

Since we are interested in unravelling the importance of the effects of the gradient and of the directional sensing in the Dd aggregation phenomena, we will let the cells propagate in a cAMP concentration field with a prescribed dynamics. In this paper, we omit some well-known properties of Dd aggregation, like the fact that Dd amoebae produce and relay cAMP upon sensing of an external concentration of the chemoattractant [19], and the phenomenon of streaming [2]. While the production and relay of cAMP is necessary for the establishment of a gradient field of cAMP, these effects are independent of the mechanisms responsible for chemotaxis and directional sensing.

2. Chemotaxis with directional sensing

We denote by \(X(x, y, t)\) the local concentration of some chemotactic substance. The equation of motion of an amoeba under the influence of a chemoattractant has the form:

\[
m\ddot{r} = -\lambda \dot{r} + \mathbf{f}(X(x, y, t))
\]

where, as usual, the dots represent time derivatives, \(\dot{r} = (r_x, r_y)\) are the spatial position coordinates of the amoeba, \(\lambda\) is a damping coefficient, \(m\) is the mass of the amoeba and \(\mathbf{f}\) represents a generic chemotactic force field. The term \(-\lambda \dot{r}\) describes the damped motion of the cells and has been measured experimentally [20].

As it is shown in the Appendix A, the direction dependent chemotactic force field is:

\[
\mathbf{f} = \left\{ \begin{array}{ll}
\text{grad} X & \text{if} \quad \text{sign}\left(\frac{\partial X}{\partial t}\right) > 0 \\
0 & \text{if} \quad \text{sign}\left(\frac{\partial X}{\partial t}\right) \leq 0
\end{array} \right.
\]

Introducing (2) into (1), the equations of motion of an amoeba under the influence of the chemotactic signal \(X(x, y, t)\) are:

(i) if \(\text{sign}\left(\frac{\partial X}{\partial t}\right) > 0\),

\[
\begin{align*}
m\ddot{r}_x &= -\lambda \dot{r}_x + \mu \frac{\partial X}{\partial x} \\
m\ddot{r}_y &= -\lambda \dot{r}_y + \mu \frac{\partial X}{\partial y}
\end{align*}
\]

(ii) if \(\text{sign}\left(\frac{\partial X}{\partial t}\right) \leq 0\).

\[
\begin{align*}
m\ddot{r}_x &= -\lambda \dot{r}_x \\
m\ddot{r}_y &= -\lambda \dot{r}_y
\end{align*}
\]

The equations of motion (3)–(4) have been derived under the assumption that amoebae are sensitive to the direction of propagation of a chemotactic wave, together with the condition that if the slope of the gradient reverses sign, there is no chemotactic motility.

To describe the spatial variation of the chemoattractant \(X(x, y, t)\), we consider the Ginzburg–Landau reaction-diffusion equation [21]:

\[
\begin{align*}
\frac{\partial X}{\partial t} &= \nu \dddot{X} - \beta \ddot{Y} + (X^2 + Y^2)(a \dddot{X} - b \dddot{Y}) + D_X \Delta X \\
\frac{\partial Y}{\partial t} &= \beta \dddot{X} + \nu \dddot{Y} + (X^2 + Y^2)(a \dddot{Y} + b \dddot{X}) + D_Y \Delta Y
\end{align*}
\]

where \(\dddot{X} = X - X`, \dddot{Y} = Y - Y`, \Delta = \frac{\partial^2}{\partial x^2} + \frac{\partial^2}{\partial y^2}\) is the two-dimensional Laplace operator, \(D_X\) and \(D_Y\) are the diffusion coefficients, and \(\nu, \beta, a, b, X`, \gamma\) and \(Y`\) are parameters. The functions \(X\) and \(Y\) represent chemical variables. In bounded two-dimensional domains with zero-flux
boundary conditions, if $v > 0$ and $a < 0$, the reaction-diffusion equation (5) has a homogeneous and unstable spatially extended steady state with constant values, $X(x, y, t) = X'$ and $Y(x, y, t) = Y'$. A local perturbation of this unstable spatial steady state leads to spatial oscillations of $X$ and $Y$. For these parameter values, the ordinary differential equation part of equation $(5)$ ($DX = DY = 0$) has a limit cycle in the $(X, Y)$ phase space with radius $r = \sqrt{-v/a}$.

All the equations are written in dimensional form and, in the following, we use the CGS system of units.

For $v > 0$ and $a < 0$ and particular choices of the initial conditions, the numerical solutions of the Ginzburg–Landau reaction-diffusion equation (5) in bounded domains and with zero-flux boundary conditions produce wave-like target (i.e. concentric) and spiral propagating patterns [21], similar to the ones observed in the early phase of aggregation of Dd colonies. Therefore, in order to investigate numerically the aggregation properties of the model equations (3)–(4), we have integrated numerically equation (5) in a circular region inside a square of side length $Ndx = 600 \times dx$. We have considered zero-flux boundary conditions on the circular region, and the parameter values considered were $v = 1 \text{s}^{-1}$, $\beta = 0.5 \text{s}^{-1}$, $a = -1 \text{cm}^2/(\text{s} \text{ mol}^2)$, $b = 1 \text{cm}^2/(\text{s} \text{ mol}^2)$, $D_X = D_Y = 0.000001 \text{ cm}^2/\text{s}$, $X' = Y' = 1.5 \text{ mol/cm}^2$, $dt = 0.005 \text{s}$ and $dx = \sqrt{6D_X dt} = 0.000173 \text{ cm}$ [22].

In Fig. 2, we show the time evolution of 1000 amoebae, calculated from (3)–(4), coupled with the chemotactic signal $X(x, y, t)$, propagating as a reaction-diffusion spiral wave emanating from the central region of the two-dimensional domain. The chemotactic signal $X(x, y, t)$ has been calculated with equation (5). The spiral wave has been induced by perturbing four contiguous lattice sites with $X$ and $Y$ taking values on the limit cycle of the diffusion-free Ginzburg–Landau equation (5) [21,22]. The values of the variables $X$ and $Y$ for these four sites have a phase advance of $\pi/2$ in phase space. Initially, the amoebae move towards the central region of the circular domain and, after a transient time, all the cells rotate near the tip of the spiral, forming a ring as shown in Fig. 2d. The radius of the ring is related to the diffusion coefficients of $X$ and $Y$, i.e. the radius increases with increasing diffusion coefficients.

Comparing the numerical simulation of Fig. 2d with the aggregation patterns of Dd cells in Fig. 1, we conclude that the rotating ring pattern of *Dictyostelium discoideum* colonies is similar to the rotating ring pattern obtained in these simulations. This suggests that the hole observed in Fig. 2d is due to the rotation of the tip of the reaction-diffusion spiral wave associated with the dynamics of the chemoattractant $X$. Chemotactic models without the directional sensing condition, introduced in (3)–(4), would lead to a non-aggregative behaviour as in the case of the simulation in Fig. 3 of Appendix A.

For the chosen parameter values, the circular region of the simulations of Fig. 2 has the diameter, $d = 600 \times dx = 0.1039 \text{ cm}$, and the wave length of the cAMP spiral wave, which is approximately $0.1039/6 = 0.017 \text{ cm}$ (compared with [2], pp. 130, Fig. 8.15). Assuming that an amoeba, initially at the periphery of the circular domain, reaches the annular region after $t = 80 \text{s}$, as simulations have shown, the order of magnitude of the amoeba speed is $v = d/(2 \times 80) = 0.00064 \text{ cm/s} = 6.4 \mu\text{m/s}$. The wave length of the cAMP wave and the speed of the amoebae

---

**Fig. 2.** (Color online) Time evolution of the position of 1000 amoebae in a two-dimensional circular domain, calculated with equations (3)–(4), for (a) $t = 0 \text{s}$, (b) $t = 40 \text{s}$, (c) $t = 60 \text{s}$ and (d) $t = 80 \text{s}$. Green spots represent amoeba positions. Red and blue correspond to regions of high and low local concentrations of the chemoattractant $X$, respectively. The chemotactic signal is a reaction-diffusion spiral wave originated at the central point (emission centre) of the two-dimensional domain and has been calculated with equations (5). The parameters of the dynamics of the amoebae are $m = 0.0001 \text{ g}$, $\lambda = 0.0001 \text{ g/s}$ and $\mu = 0.00001 \text{ g cm}^2/\text{s}^2$. The parameters associated with the dynamics of the chemotactic signal are described in the text.
are within the orders of magnitude of the observed values [1]. In this simple calibration, the simulation time relates with the real time of the motion of amoebae by a scaling constant also affecting $m$, $\lambda$, and the chemoattractant concentration.

We have also tested the amoeba motion in the presence of a concentric chemotactic waves or target waves, generated by the Ginzburg–Landau equation (5). In this case, the amoebae move towards the centre of the circular domain, accumulating near the wave initiation centre (Fig. 4). The target wave pattern was generated from a spatially uniform initial condition on the unstable steady state $\langle X = X', Y = Y' \rangle$ with a small perturbation at the centre of the circular region. After reaching the region of the initial perturbation, all the amoebae stop moving.

To understand the role of the cAMP emission centres in Dd aggregation, in Fig. 5, we show the distribution of 1000 amoebas at time $t = 80$ s, calculated from a random initial distribution of the chemotactic signal $X(x, y, t)$ and of the amoebae. From this simulation, we conclude that if the chemoattractant is not produced from emission centres, there are no aggregation movements towards localised spatial regions. In order to have aggregation towards a finite number of localised spots, the chemotactic signal must be initiated in the emission centres. Other simulations have shown that if the aggregation centres are within a short distance, the interference between the reaction-diffusion waves destroy the process of aggregation. This simulation shows that the spontaneously formed cAMP emission centres, are necessary for the aggregation and formation of the slime mould and the subsequent multicellularity process.

3. Discussion

From the model just described, we conclude that the typical circular ring pattern found in the aggregation of Dd colonies is explained by the sensitivity of the amoebae to the direction of propagation and to the gradient of the chemoattractant, together with the spiral shape of the
propagating chemoattractant signal. In other models without directional sensing [23], spiral chemoattractant waves do not produce this rotating motion of the Dd cells. This dynamic effect is at the origin of the multicellularity and of the formation of the mould [2], prior to the emergence of the Dd slug.

The sensitivity of amoebae to the gradient of concentration is due to the large number of cAMP receptors (of the order of 50,000) distributed along the plasmic membrane of amoebae [2]. The density of occupation of the cAMP receptors along the amoeba’s external membrane is a sensor for the gradient of chemoattractant, being at the origin of the motility of the cells against the chemotactic gradient. Dd amoebae can detect a 1% difference in concentration of the chemoattractant between the front and the back of the cells [5,24]. This spatial information together with the local sensitivity of the cells to the time variation of the concentration of chemoattractant, determines the directional sensing of amoebae. If the local concentration of chemoattractant decreases, the binding rate of cAMP to its receptors also decreases, justifying the sensitivity of the cells to the time variation of the concentration of chemoattractant, in agreement with observations [18].

These biological mechanisms explain why amoebae are sensitive to the temporal and spatial derivatives of the chemotactic field, explaining the stop moving condition, as observed in the Dd aggregation [6,12]. On the other hand, our simulations results are consistent with the polarization effects observed in the amoebae during aggregation [11]. We have concentrated this investigation on the roles of the cAMP gradient and directional sensing, solving the well-known chemoattractant wave paradox [8]. Consequently, in the present approach, we did not consider other well-known effects observed in the aggregation of Dd colonies as streaming [2,13], and cAMP production by amoebae [19].

Disclosure of interest

The authors declare that they have no conflicts of interest concerning this article.

Acknowledgments

We would like to thank Christiane Hilgardt for helpful discussions. We would like to thanks the comments of one of the anonymous reviewers of this paper. Travel support by the European Science Foundation research network program FUNCODYN (Functional Dynamics in Complex Chemical and Biological Systems) is gratefully acknowledged. The first author would like to thank IRES, where the final version of this paper has been prepared.

Appendix A

We consider a chemotactic substance distributed along a one-dimensional domain. We assume that the concentration of this substance evolve in time and space as a one-dimensional travelling wave, symmetrically along the positive and the negative x-directions. This centrally symmetric wave is generated at the point \( x = 0 \) and has radial propagation speed \( c > 0 \). To simplify, we assume that the concentration of the chemotactic substance propagates according to the wave type law:

\[
\phi(x, t) = \begin{cases} 
A(1 - \cos(ct - |x|)), & \text{if } |x| \leq ct \\
0, & \text{if } |x| > ct
\end{cases}
\]  

(6)

where \( A \) is the concentration amplitude, and \( t \geq 0 \).

At time \( t = 0 \), we consider an immobile amoeba at position \( x_0 > 0 \). Assuming that the response of the amoeba to the chemotactic signal is solely proportional to its gradient, as it is generally assumed in the literature [13,15,16], the one-dimensional equation of motion of the amoeba is:

\[
mx' = -\lambda x + \frac{\partial \phi}{\partial x}
\]

(7)

where \( \lambda \) is a damping constant and \( x(t) \) is the position of the amoeba at time \( t \). Equation (7) describes the chemotactic response of an amoeba without directional sensing.

To implement directional sensing in the model equation (7), we now introduce vectorial notation. The effect of the chemoattractant \( \phi(x, t) \) on the amoeba is assumed to be described by the force:

\[
\vec{F} = \left( \frac{\partial \phi}{\partial x} \right) \vec{n} \vec{e}_x = \left( \frac{\partial \phi}{\partial x} \right) n_x \vec{n} \vec{e}_x
\]

(8)

where \( \vec{n} = n_x \vec{e}_x \) is the direction of propagation of the chemical wave. In this one-dimensional case, \( n_x = \pm 1 \) and \( \vec{e}_x \) is the usual direction versor. As the wave (6) propagates radially from the origin of coordinates \( x = 0 \), if \( x > 0 \), then, \( n_x = +1 \). If \( x < 0 \), then \( n_x = -1 \). To introduce the directional sensitivity of the amoeba, we assume that the amoeba moves towards the origin \( x = 0 \) when it senses a positive slope of the gradient, measured in the negative x-direction (\( \frac{\partial \phi}{\partial x} < 0 \)). The amoeba stops moving when the slope of the gradient reverses sign. Therefore, the response of the amoeba to the chemical signal with directional sensing is described by the effective force:

\[
\vec{F} = F e_x = \begin{cases} 
\left( \frac{\partial \phi}{\partial x} \right) n_x \vec{n} \vec{e}_x = \left( \frac{\partial \phi}{\partial x} \right) n_x \vec{n} \vec{e}_x, & \text{if } n_x < 0 \\
0 \vec{e}_x, & \text{if } n_x \geq 0.
\end{cases}
\]

In order to determine a closed form for the direction versor \( n_x \), we consider the scalar wave field, \( \phi(x, t) = \cos(ct - x) \), where \( c \) is a positive constant, \( |x| \leq ct \) and \( x, t \in \mathbb{R} \).

As \( c > 0 \), the scalar field \( \phi(x, t) \) propagates in the positive x-direction and, for each fixed \( t = t_1 \), we can have two cases: (i) if, \( \frac{\partial \phi}{\partial x} \big|_{x=t_1} < 0 \), then, \( \frac{\partial \phi}{\partial x} \big|_{x=t_1} > 0 \); (ii) if, \( \frac{\partial \phi}{\partial x} \big|_{x=t_1} > 0 \), then,
\[ \frac{\partial \phi}{\partial t} |_{t=t_1} < 0. \] Therefore, in both cases, \( \text{sign} \left( \frac{\partial \phi}{\partial x} \right) < 0, \) where \( \text{sign}(x) = 1 \) for \( x > 0, \) and \( \text{sign}(x) = -1 \) for \( x < 0. \)

Assume now that the scalar field propagates in the negative \( x \)-direction, that is, \( c < 0. \) (i) If \( \frac{\partial \phi}{\partial t} |_{t=t_1} < 0, \) then, \( \frac{\partial \phi}{\partial t} |_{t=t_1} < 0. \) (ii) If \( \frac{\partial \phi}{\partial t} |_{t=t_1} > 0, \) then, \( \frac{\partial \phi}{\partial t} |_{t=t_1} > 0. \) Therefore, in both cases, \( \text{sign} \left( \frac{\partial \phi}{\partial t} \right) > 0. \)

So, the direction of propagation of the one-dimensional scalar field \( \phi(x, t) \) is:

\[ n_x = - \text{sign} \left( \frac{\partial \phi}{\partial t} \right). \] (10)

Introducing (10) into (9), the effective force on the amoeba is:

\[ \vec{F} = \vec{F}_x = \begin{cases} \left( \frac{\partial \phi}{\partial x} \right) \sigma_x, & if \ - \text{sign} \left( \frac{\partial \phi}{\partial t} \right) < 0 \\ 0 \sigma_x, & if \ - \text{sign} \left( \frac{\partial \phi}{\partial t} \right) \geq 0. \end{cases} \] (11)

Hence, the equation describing the motion of the amoeba is:

\[ m \ddot{x} = - \lambda \dot{x} + \vec{F}. \] (12)

where \( \vec{F} \) is given by (11). Equation (12), together with (11), describes the chemotactic force on the amoeba, taking into account directional sensing.

In Fig. 3, we show the time evolution of the position of a cell with initial condition \( x_0 = 2 \) and \( \dot{x} = 0, \) calculated with equations (7) and (11)–(12), respectively, and chemotactic field (6). Initially, the cell is immobile. For \( t \geq 0 \), \( \sigma = 2/c, \) the movement of the amoeba is equally described by the two model equations and the amoeba starts to move in the direction of the point \( x = 0. \) For the case of simple chemotaxis, the motion is described by equation (7). For larger values of \( t, \) the he slope of the chemotactrant concentration changes sign and the amoeba reverses the direction of motion. In this case, the motion of the amoeba is oscillatory and numerical simulations show that the cell slowly deviates from the origin of coordinates. Asymptotically in time, the position of the amoeba goes away from the source of the chemotactrant. A simple linear analysis shows that equation (7) is linearly unstable around the phase space fixed point \( (x = 0, \dot{x} = 0). \)

In contrast, using model equation (12), with the force (11), amoebae direct their motion towards the origin of coordinates, where the source of the chemotactrant is located.

The simulations in Fig. 3 show that a model based on equation (11)–(12) describes the observed motility in the aggregation of Dictyostelium discoideum colonies. On the other hand, the model based on equation (7) shows that wave driven chemotaxis without directional sensing leads to divergent motion.

For wave propagation in two-dimensional domains, the directional sensing condition in (11) and the direction versor (10) are simply generalised. As the gradient is normal to the level curves of a surface, in two space dimensions, the normalized direction of propagation is then:

\[ \vec{N} = - \frac{\left| \frac{\partial \phi}{\partial x} \right|}{\text{grad} \phi} \text{signal} \left( \frac{\partial \phi}{\partial x} \right) \vec{e}_x \]

\[ - \frac{\left| \frac{\partial \phi}{\partial y} \right|}{\text{grad} \phi} \text{signal} \left( \frac{\partial \phi}{\partial y} \right) \vec{e}_y \]

\[ = - \frac{1}{\text{grad} \phi} \left( \frac{\partial \phi}{\partial x} \right) \text{signal} \left( \frac{\partial \phi}{\partial x} \right) \vec{e}_x \]

\[ - \frac{1}{\text{grad} \phi} \left( \frac{\partial \phi}{\partial y} \right) \text{signal} \left( \frac{\partial \phi}{\partial y} \right) \vec{e}_y \]

and \( \| \vec{N} \| = 1, \) provided \( \| \text{grad} \phi \| = 0. \)

References


