Persistent random motion: Uncovering cell migration dynamics

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In this paper we study analytically the stick–slip models recently introduced to explain the stochastic migration of free cells. We show that persistent motion of cells of many different types is compatible with stochastic reorientation models which admit an analytical mesoscopic treatment. This is proved by examining and discussing experimental data compiled from different sources in the literature, and by fitting some of these results too. We are able to explain many of the ‘apparently complex’ migration patterns obtained recently from cell tracking data, like power-law dependences in the mean square displacement or non-Gaussian behavior for the kurtosis and the velocity distributions, which depart from the predictions of the classical Ornstein–Uhlenbeck process.

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1. Introduction

Studying the properties of cell movement is of fundamental interest to understand many physiological processes in living organisms (embryogenesis, wound healing, etc.) as well as their malfunctions (e.g. in tumors growth). Exhaustive research in this field is focused on the biophysical and biochemical intracellular processes driving cell signaling and motility. However, for many purposes like the analysis of tracking experiments or the implementation of migration algorithms into more general models, the use of phenomenological approaches as those based on stochastic processes is extremely helpful. Within this context, the Ornstein–Uhlenbeck (OU) process (Uhlenbeck and Ornstein, 1931) has been considered for decades the archetypal model to describe persistent random motion of cells or other particles/organisms. This process is defined by the Langevin equation

$$\frac{dv}{dt} = -\frac{1}{\tau}v + \sqrt{\frac{2D}{\tau}} \xi(t)$$

(1)

for the velocity vector $v$, where $\xi(t)$ represents a vector with white-noise components, $D$ is the diffusion coefficient characteristic of Brownian motion, and the timescale $\tau$ is often called the persistence time. Fibroblasts locomotion (Gail and Boone, 1970), the motility of lung epithelial cells (Wright et al., 2008), microvessel endothelial cells (Stokes et al., 1991) or self-motile colloids (Howse et al., 2007) are just some examples from the vast amount of this model. Even in recent studies where cells have been found experimentally to exhibit more complex migration patterns, the OU process is still the reference model to which more sophisticated approaches are compared (Dieterich et al., 2008; Li et al., 2008; Takagi et al., 2008).

The popularity of this approach is in part founded on the fact that the mean square displacement (MSD) and the velocity autocorrelation function (VACF) of cells have been often experimentally observed to fit the behavior it predicts

$$\langle r^2(t) \rangle = 2nDt(e^{-t/\tau} + t/\tau - 1)$$

(2)

$$\langle v(t)v(0) \rangle = \frac{nD}{\tau} e^{-t/\tau}$$

(3)

where (2) is known as the Füth formula (Füth, 1920) and $n$ is the space dimension. Note that Eqs. (2) and (3) describe the behavior of averaged quantities, while much less efforts have been performed in order to verify directly Eq. (1) from experimental cell trajectories, since it requires much better resolution in data. Actually, the only systematic works conducted in that way we have found in the literature led to strong discrepancies between statistics from experimental trajectories and the OU process (Takagi et al., 2008; Selmezzi et al., 2005, 2008). The results found in these references are that the acceleration perpendicular to the direction of motion exhibits zero-mean fluctuations, while the mean acceleration in the direction of motion decreases with the speed in a seemingly linear manner. The authors in Selmezzi et al. (2008) have already noted that the Füth formula can also be derived from other models of persistent motion. So that, misinterpretations of cell migration dynamics can arise from assuming that there is an univocal equivalence between (2) and (3) and the OU process.
To illustrate these ideas let us consider a reorientation model which consists of a particle traveling with constant speed $v_0$ in a 2D space and whose orientation angle $\theta$ (which determines its direction of motion) exhibits white-noise fluctuations with a characteristic strength $\sigma$. For that situation, the microscopic equations of motion are

$$\frac{d\theta}{dt} = \sigma \dot{z}(t), \quad \frac{dr}{dt} = v_0 u_r$$

where $u_r = (\cos \theta, \sin \theta)$ is a unit vector in the direction of motion and $\dot{z}(t)$ is white noise. One can now define the probability density $P(r, \theta, t)$ of finding the particle at position $r$ with orientation $\theta$ at time $t$, given the initial conditions $r_0$, $\theta_0 = (0,0)$. Applying standard techniques from stochastic calculus (Gardiner, 2004), we can obtain the Fokker–Planck equation for the probability density $P$ (see Schienbein and Gruler, 1993)

$$\frac{\partial P}{\partial t} = \frac{\sigma^2 \partial^2 P}{2} - v_0 u_r \cdot \nabla P$$

It is straightforward now to check from Eq. (5) that the MSD and the velocity auto-correlation of a particle governed by this model also satisfy Eqs. (2) and (3), with $\tau = 2/\sigma^2$, $D = 2v_0^2/\sigma^2$. So that it represents an alternative to the OU process for describing persistent motion (see Schienbein and Gruler, 1993; Othmer et al., 1988 for some seminal papers on this field). The form of the first moments of $P$ coincide for both models (1) and (4), but their microscopic differences become apparent for higher-order moments. For example, it is easy to check that the kurtosis $\langle r^2(t) \rangle / \langle r^2(0) \rangle^2$ is different in each model, though in both cases it tends to 3 for $t \to \infty$, as expected for a Gaussian variable.

Although it is not easy in general to elucidate the correct pattern of motion from experimental data, we think that the reorientation model has some advantages over the classical approach in Eq. (1). Firstly, it seems more natural to write the equations of motion in terms of the orientation coordinate $\theta$ (instead of using a fixed coordinate system), according to the results for the mean acceleration of cells reported in Takagi et al. (2008) and Selmecki et al. (2008). This will be specially appropriate for the case of non-isotropic particles (for example, rod-shaped), which move preferentially along one of their axes. Secondly, the simplicity of the reorientation model (4) makes generalizations of the model easier to implement, while generalizations of the OU process often involve memory kernels or other non-trivial effects. This has important implications on how experimental data is interpreted, too. When the experimental results from cell trajectories are observed to depart from the predictions of the OU process, it is often interpreted as a trace of complexity at some level. Here, we explore simple and direct generalizations of the reorientation model which can explain many of these ‘apparently complex’ patterns of motion. Experimental data, obtained from different works previously published, will be used to support these ideas (Section 4).

2. Models overview

In this Section we summarize some of the different stochastic approaches that have been recently proposed to explain the dynamics of self-propelled particles (as is the case of cells in a diluted media). This will serve to put in context the stick–slip model that will be presented and developed in Section 3.

2.1. Model in which speed and direction fluctuate independently

The assumption made in Eq. (4) that the particles travel with constant speed can be relaxed by assuming instead that speed fluctuates around a mean value $\langle v \rangle$, giving rise to particles fluctuating both in their direction of motion and their absolute speed. This model has been proposed recently to explain the motion of propelled nanoparticles (Peruani and Morelli, 2007), but application to cell migration comes straightforward. Actually, the same idea had also been explored in the context of cell motion some years before (Schienbein and Gruler, 1993), and similar models had already been used to explain migration of *Amoeba proteus* (Miyoshi et al., 2003; Masaki et al., 2007) (though the results in those works were mainly based in computer simulations).

Fig. 1A describes schematically the microscopic behavior of a particle according to this model. The dotted lines represent the cell trajectory, in which changes of direction occur and the value of the speed changes at some instants (these instants are marked by crosses). As can be seen from the situation of the crosses, random changes in direction and speed are completely independent processes. As a consequence, one can study separately the fluctuations in speed and direction. If the cells are assumed to obey a certain velocity distribution $\rho(v)$ and the speed changes occur with a characteristic (constant) rate $\beta$ then

$$\frac{\partial P(v,t)}{\partial t} = -\beta P(v,t) + \beta \rho(v)$$

(6)

gives us the evolution of $P(v,t)$, the probability density of particles traveling with speed $v$ at time $t$.

Similarly, for the orientation angle $\theta$ (defined as in the Introduction Section), provided that orientation changes are smooth, one can use a diffusion approximation

$$\frac{\partial P(\theta,t)}{\partial t} = \kappa \frac{\partial^2 P(\theta,t)}{\partial \theta^2}$$

(7)

where $\kappa$ is the diffusion parameter for the fluctuations in the direction of motion.

Eqs. (6) and (7) can be exactly solved with the appropriate initial conditions. Since changes in speed and direction are independent, one has $P(v,\theta,t) = P(v,t) \cdot P(\theta,t)$. Then, from standard statistics in 2D one can compute, for instance, the mean distance traveled by the particle in the model $\langle \text{dr}(t) \rangle = \int dt \int_0^{\infty} dv \rho(v,t) \nu$, or any moment $\langle r^2 \rangle$, $\langle v^2 \rangle$ of arbitrary order. So it is found that the MSD reads (Peruani and Morelli, 2007)

$$\langle r^2(t) \rangle = 2 \frac{\langle v^2 \rangle - \langle v \rangle^2}{(k+\beta)^2} (e^{-(k+\beta)t} + (k+\beta)t - 1) + \frac{\langle v \rangle^2}{\kappa} (e^{-\kappa t} + \kappa t - 1)$$

(8)

The first term in Eq. (8) represents Brownian motion arising from a particle with a fluctuating speed, while the second term accounts for rotational diffusion arising from the reorientation process.

For the VACF within this approach one obtains

$$\langle \text{v}(t) \text{w}(0) \rangle = \langle v \rangle^2 e^{-\kappa t} + (\langle v^2 \rangle - \langle v \rangle^2) e^{-(k+\beta)t}$$

(9)
which exhibits a double-exponential decay, compared to the single exponential one finds from the OU process.

2.2. Models with simultaneous fluctuations in speed and direction

In a recent work (Campos and Méndez, 2009) two of the authors proposed a variation of the model in Section 2.1. According to it, the time instants at which the speed of the particle changes coincide with those where the direction changes (see Fig. 1B). In the following, we will refer for simplicity to the models in Figures 1A and 1B as ‘model A’ and ‘model B’, respectively. So, model B corresponds to a motion pattern in which sojourns of constant speed are separated by (instantaneous) reorientation events. Though this is a relatively subtle modification of the previous model, the fact that now both types of fluctuations are not independent makes the present mathematical treatment quite different. In Campos and Méndez (2009) we use the framework of continuous-time random walks (CTRW) to implement this idea (see Metzler and Klafter, 2000 for a comprehensive review on such models). Within this framework one can derive an equation for the probability density \( P(r,t) \). In our specific case, this equation can only be explicitly written in the Fourier–Laplace space; so the transformation \( P(q_x,q_y,t) \rightarrow \mathcal{F}[P(q_x,q_y,t)] = \tilde{P} \) is used, with \( q_x, q_y, s \) being the transform variables of \( x, y, t \), respectively. The evolution equation takes the form (Campos and Méndez, 2009)

\[
\frac{\partial^2 \tilde{P}}{\partial q_x^2} + \frac{\partial^2 \tilde{P}}{\partial q_y^2} = \left[ s + iv(q_x \cos \theta + q_y \sin \theta) + \frac{v_0^2(q_x \cos \theta + q_y \sin \theta)^2}{s^2 + v_0^2} \right] \tilde{P} - \frac{1}{2\pi} \frac{\partial^2}{\partial t^2} \tilde{P}
\]

where the parameter \( v_0 \) measures the amplitude of the speed fluctuations. So, for \( v_0 = 0 \) it is clear that (5) will be recovered after inverting by Fourier–Laplace.

From (10), the moments of \( P(r,t) \) can also be found in a relatively easy way (see Campos and Méndez, 2009 or the Appendix for further details). The expressions for the MSD and the VACF one finds from (10) coincide with those from the model A (Eqs. (8) and (9)) after redefining the parameter \( \kappa \) as \( \sigma^2 \beta \). As it happened with the two models discussed in the Introduction, these two approaches have the same first and second-order moments, but the microscopic differences between them (sketched in Fig. 1) become evident for higher-order moments. The main advantage of this second approach based in the CTRW is that different extensions of the model can be implemented in a very natural way; this will be shown in Section 3.

2.3. Models with memory kernels

In comparison with the two previous approaches, where the microscopic details of the model (Fig. 1) are absolutely clear, many other approaches used to fit experimental data on cell migration are based on the introduction of memory functions, or similar, into well established models, e.g. the OU process. One example of this is the model presented in Selmecki et al. (2005) and lately used in Takagi et al. (2008) to fit the spontaneous movements of Dictyostelium cells. This approach is based on the stochastic differential equations

\[
\frac{dv}{dr} = -v(t) + \mu V(t) + \sigma \xi(t) \quad \text{(11)}
\]

\[
\frac{dV}{dr} = \mu V(t) - \gamma V(t) \quad \text{(12)}
\]

where \( V(t) \) represents a memory function, whose time evolution is governed by (12), while \( \mu \) and \( \gamma \) represent constant parameters. Note also that velocity-dependent friction (\( \tau^{-1} \)) and noise memory functions (\( \sigma \)) have been introduced if compared with Eq. (1). All of these functions \( V(t), \sigma v(t) \) and \( \tau^{-1}(v(t)) \) are to be adjusted from data fitting.

Though this kind of models provide excellent agreement to data, their drawback is that they require the help of mechanistic models or similar in order to justify the origin of such memory kernels so their hypothesis can be validated. The aim of our work, however, is not to criticize models based on kernels or memory effects, since the goodness of these models for fitting experimental data is clear. We just want to show that the experimental results can often be explained too by means of approaches like those in Sections 2.1 and 2.2, or direct generalizations of them. In those cases, memory effects are implicitly implemented through the time correlations introduced by the reorientation process. At this stage, we do not have any criteria to elucidate which of these approaches work better. We just think that reorientation models represent a simpler and intuitive alternative to models based on kernels. Besides, a complete analytical treatment is feasible for such models, which facilitates understanding.

3. Mesoscopic description for stick–slip models

The concept of stick–slip processes has been introduced recently in Selmecki et al. (2008). Actually, it corresponds to the ‘run and tumble’ pattern long used to describe bacterial motion (Berg, 2004). However, in order to stress the connection of our work with that in Selmecki et al. (2008) we will rather use the term stick–slip.

Stick-slip processes have been introduced as a sort of numerical models (algorithms) able to implement realistically protrusion-retraction mechanisms responsible for cell migration. Here we will show that the CTRW formalism we used in Campos and Méndez (2009) can be extended to implement these processes. By the analytical treatment of such models we expect to facilitate their understanding and check their validity for fitting cell data.

According to the discussion in the Section 2.2, we will consider a cell trajectory in 2D where the particle changes its direction of motion and absolute speed jointly at some time instants with a characteristic rate \( \beta \), while \( \sigma \) is a measure of how much the orientation can change at each of these events. Now each reorientation event is assumed to last a finite random time, controlled by a constant characteristic rate \( \beta_2 \). Regarding the distribution of speeds, we will consider here the simplest version of the model, in which the particle travels with constant speed \( v_0 \) during the motion periods. We think that the general model with an arbitrary distribution of velocities is also feasible analytically in the light of the velocity models proposed in Zaburdaev et al. (2008); that case, however, involves more complicated considerations, so we plan to deal with it in a separate forthcoming work.

Following the lines of CTRW models (see Campos and Méndez, 2009; Metzler and Klafter, 2000 and the references therein), we define \( P(r,t) \) as the probability density for the particle to be located at position \( r \) with orientation \( \theta \) at time \( t \). The label \( i = 1 \) stands for those particles waiting (while reorienting) at that position, while \( i = 2 \) denotes the particles in motion passing through that point. The total density of cells is then computed as \( P = P_1 + P_2 \). Also, we need to introduce \( J_i(r,t) \) as the probability density of particles with orientation \( \theta \) at time \( t \) stop moving at position \( r \) (\( i = 1 \)), and those which start moving at position \( r \) after a waiting period (\( i = 2 \)). The evolution of the model is then given by the system of mesoscopic equations

\[
J_i(r,t) = \int d\omega \int_0^{2\pi} d\theta \int_0^t dt' J_i(r-\omega_{r,t},\theta_{r,t},t-t') \mathcal{L}(\theta_{r,t}) \psi_i(\theta_{r,t},t) + \frac{\delta \omega_{r,t}}{2\pi} \quad \text{(13)}
\]
\[ J_2(\mathbf{r}, \theta, t) = \int_0^t dt' J_1(\mathbf{r}, \theta, t-t') \phi(t') \]  
\[ P_1(\mathbf{r}, \theta, t) = \int d\mathbf{r} \int_0^t dt' J_1(\mathbf{r}, \theta, t-t') \phi^*(t') \]  
\[ P_2(\mathbf{r}, \theta, t) = \int d\mathbf{r} \int_0^t dt' J_2(\mathbf{r}-\mathbf{r}', \theta, t-t') \psi(\mathbf{r}', \theta, t) \]

where we have assumed for simplicity that initially all the particles start waiting from position \( \mathbf{r} = 0 \) at \( t = 0 \) (see the last term in Eq. (13)). The first term on the right in (13) represents the contribution from those particles that started moving at time \( t-t' \) with orientation \( \theta \) and then stop after covering a distance \( \mathbf{r} \) in a travel time \( t' \) (according to the probability distribution \( \psi(\mathbf{r}, \theta, t) \)) and take a new orientation value \( \theta' \) according to the distribution \( T(\theta' - \theta) \). In Eq. (14) we apply the waiting times distribution \( \phi(t) \) for particles which stopped moving at time \( t-t' \) and start moving again at time \( t \). The distribution \( \phi(t) \) in (15) stands for the probability that the particle has not started moving yet after a waiting time \( t' \). Finally, in (16) \( \psi(\mathbf{r}, \theta, t) \) denotes the probability that a particle has not stopped after a time \( t \) since it started moving, during which it has covered a distance \( \mathbf{r} \) in the direction \( \theta \).

Since we have assumed in Section 3 that moving and waiting periods are governed by constant rates \( \beta \) and \( \beta_2 \), respectively, we take the corresponding probability distribution functions to be exponential (that is, a Markov assumption). So that, for the constant speed case we have \( \psi(\mathbf{r}, \theta, t) = \beta e^{-\beta t} \delta(\mathbf{r}-v_0 \mathbf{u}_0) \) and \( \phi(t) = \beta e^{-\beta t} \), where \( \mathbf{u}_0 \) represents, as mentioned above, a unit vector in the direction of motion. Correspondingly, the other distributions will read \( \psi^*(t') = e^{\beta t'} \delta(\mathbf{r}-v_0 \mathbf{u}_0) \) and \( \phi^*(t) = e^{\beta t} \). For further details on this kind of models, the reader is addressed to Campos and Méndez (2009), Zumofen and Klafter (1993) and Méndez et al. (2004).

For the case of a smooth reorientation process (smooth changes in \( \theta \), the closed equations for \( P_1 \) and \( P_2 \) can be easily found and solved in the Fourier–Laplace space (all the details are given in the Appendix). The moments of these distributions can thus be found easily; for instance, we obtain for the ensemble-averaged MSD

\[ \langle r^2(t) \rangle = \int_0^{2\pi} d\theta \int r^2 P(\mathbf{r}, \theta, t) dr = 2v_0^2 \left( p_0(t) + p_1(t)e^{-\lambda_1 t} + \sum_{k=2}^5 p_k e^{-\lambda_k t} \right) \]

where the functions \( p_0(t), p_1(t) \) are first-order polynomials in \( t \) and the \( p_k(t) \) (for \( k = 2 \ldots 5 \)) are independent of \( t \) (the exact expressions for all of them are provided in the Appendix). Likewise, the characteristic exponents \( \lambda_k \) read

\[ \lambda_1 = \beta + \beta_2 \]  
\[ \lambda_2 = \frac{1}{2}(\beta + \beta_2 + \sqrt{\beta^2 + \beta_2^2 - 4\beta \beta_2 \sigma^2}) \]  
\[ \lambda_3 = \frac{1}{2}(\beta + \beta_2 - \sqrt{\beta^2 + \beta_2^2 - 4\beta \beta_2 \sigma^2}) \]  
\[ \lambda_4 = \frac{1}{2}(\beta + \beta_2 + \sqrt{\beta^2 + \beta_2^2 - 4\beta \beta_2 \sigma^2}) \]  
\[ \lambda_5 = \frac{1}{2}(\beta + \beta_2 - \sqrt{\beta^2 + \beta_2^2 - 4\beta \beta_2 \sigma^2}) \]

From Eq. (17) and the explicit expression for \( p_0(t) \) (see (32)) one finds out how the asymptotic diffusion coefficient depends on the characteristic rates in the model

\[ D = \lim_{t \to \infty} \frac{\langle r^2(t) \rangle}{4t} = \frac{v_0^2}{4\beta \sigma^2} \left( \frac{\beta}{\beta_0} \right)^2 \]

On the other side, the VACF can be evaluated from (17) (since the motion process is isotropic) by applying the Green–Kubo relation (Kubo et al., 1991)

\[ \frac{\langle (x(t) - x(t_0))^2 \rangle}{dt_0} = 2 \int_{t_0}^{t} \langle v_x(t) v_x(t_0) \rangle dt \]

where the first term on the right is the friction kernel and \( m \) is the mass of the particle. For further details on the VACF, the reader is addressed to Klafter and Blumen (1986), Méndez et al. (2000) and Klafter and Shinbrot (1996).

4. Experimental data for cell migration

In this section we will try to show how reorientation models, as the stick–slip case introduced in the previous Section, can explain many of the departures from the OU process that have been found recently in different cell migration experiments. To complete our discussion we will also compare with the results obtained from the models A, B, C and D. We will discuss different parameters reported in experimental works and will try to fit the experimental data to the stick–slip process presented here. For such fitting procedure we will mainly use the data collected in Takagi et al. (2008) for vegetative and starved (after 5.5 h) Dicyostelium discoideum cells (since this is one of the most exhaustive works we have found in the literature), plus some data from Dieterich et al. (2008) and Selmeczi et al. (2008).

(a) Mean-square displacement: This is probably the most recurrent parameter in studies on cell migration. The OU process predicts, according to (2), a transition from ballistic to diffusive behavior which occurs at timescale \( \tau \). However, departures from that formula have increasingly appeared during the last years. Most of these departures consist of a power-law scaling of the type \( \langle r^2(t) \rangle \sim t^\alpha \) (Dieterich et al., 2008; Takagi et al., 2008; Upadhyaya et al., 2001; Thurner et al., 2003; Potdar et al., 2009). This easily connects with the theory of anomalous diffusion, which is very popular among theoreticians at present (Metzler and Klafter, 2000), so concepts like ‘Levy flight’ or ‘long-range memory’ are now relatively common within this field. According to theoretical works, anomalous diffusion has different origins: (i) long-range time correlations exist in the particles dynamics, which can (be a consequence of different internal or external mechanisms, or (ii) movement constraints are imposed by heterogeneous (fractal-like) media (Ben-Avraham and Havlin, 2005) or by crowding effects (Banks and Fradin, 2005). In most works where power-law
dependences of the MSD have been found, the influence of external mechanisms or constraints has been minimized. Thus, the only explanation seems to be that internal mechanisms in the cell should be responsible for the emergence of memory effects; this has been the interpretation suggested by most of the authors. However, this idea is quite speculative, since experimental evidence for these long-range memory mechanisms is scarce, in the best case. According to that, we can ask: ‘Is there an alternative to explain why these power-law dependences are observed so often without invoking such concepts from anomalous diffusion?’.

In our opinion a positive answer can be provided by the reorientation models studied here (alternative answers have also been explored in the literature: for example, it has been recently shown Hapca et al. (2009) that heterogeneous diffusivities in a set of individual particles can lead statistically to the emergence of anomalous diffusion at the ensemble level).

For the models depicted in Fig. 1 the form of the MSD is given by (8). According to that, the MSD will be mainly determined by the relative values of the characteristic rates \( \beta \) and \( \kappa \). For \( \beta \ll \kappa \), the formula is completely governed by \( \kappa \) and so a single ballisic-to-diffusive transition will be found, as in the OU process. For \( \beta \gg \kappa \) two transitions at very different timescales (one for each term in (8)) occur, so we observe a sequence ballistic–diffusive–ballistic–diffusive (Peruani and Morelli, 2007). Interestingly, for intermediated values of \( \beta \) the MSD often fits quite well a power-law dependence \( \langle r^2(t) \rangle \sim t^\alpha \) with an exponent \( \alpha \) which lies between 1 and 2. Exactly the same behavior holds for the expression (17) we have derived for the stick–slip model.

These results for the MSD have been discussed in detail in Campos and Méndez (2009), where they have been used to interpret data from experiments with nanoparticles. Our hypothesis is that the same could apply to cell migration data, so some of the experimental works where anomalous diffusion has been reported could actually correspond to a behavior like that in (8).

There are some facts that seem to support this idea:

(i) The values of the anomalous exponents \( \alpha \) found experimentally (in the absence of crowding or geometrical constraints) are always between 1 and 2.

(ii) The power-law tails in the MSD often extend over just one or two decades of time values (in the best case) due to experimental limitations. So, it is not clear that the asymptotic regime has been reached yet.

By ‘asymptotic regime’ we mean here that the observational times have reached the condition \( t \gg \kappa^{-1} \), where diffusive behavior should be recovered. This condition is equivalent to the ‘criterion for distinguishing true superdiffusion from correlated random-walks’ given in Viswanathan et al. (2005). The authors in that reference already detected the formal problem reported here and discuss its implications on the analysis of cell motility data as well as for foraging in animals. So, this controversy between anomalous or ‘similar to anomalous’ is not a new problem.

Our hypothesis is tested by fitting the experimental data from Takagi et al. (2008) to the result (17) and to the models A and B (Eq. (8)). Fig. 2 (left) shows that both models (solid and dotted lines, respectively) can fit reasonably well the experimental results, though apparently the data seems to exhibit a power-law dependence (as claimed originally by the authors in Takagi et al., 2008). For the sake of completeness, we also show (Fig. 3) the data fitting corresponding to the MSD results in Dieterich et al. (2008). We stress, however, that from this fitting procedure it is very difficult to determine which models fits the data best; anyway, this is not the aim of our paper. To perform such a detailed study it would be necessary to have access to the original raw data, and even in that case it is unlikely that concluding results could be found.

(b) Velocity autocorrelation function: We have already shown the results for the VACF one obtains in the stick–slip model (Eq. (21)) and for reorientation models A and B (Eq. (9)). Such multi-exponential functions are able to explain all the results of cell migration in 2D we have found in the literature. For example, this double-exponential behavior has been explicitly observed in experiments for HaCaT cells, dermal keratinocytes or different developmental stages of Dictyostelium discoideum (Takagi et al., 2008; Selmeczi et al., 2005, 2008; Maeda et al., 2008). The single exponential decay found in other works will be recovered when fluctuations in the speed are much larger than fluctuations in the direction (\( \beta \ll \kappa \)). Also, in those cases when the time resolution of observations is not good enough to reproduce accurately the region \( t < (\beta + \kappa)^{-1} \), it will be impossible to observe experimentally the second exponential term and a single exponential will be found. The only exception is the case reported in Dieterich et al. (2008), where the migration of epithelial Madin–Carby canine kidney cells has been investigated. There, a power-law scaling for the VACF has been observed. However, this behavior is found barely over one decade of time values (see Dieterich et al., 2008, Fig. 3), so it is plausible that other curves could be able to fit well the data, too.

The comparison between Eq. (21) (stick–slip model), Eq. (9) (obtained from models A or B) and the data for Dictyostelium cells obtained from the experimental setting in Takagi et al. (2008) and Selmeczi et al. (2008) shows an excellent agreement too (Fig. 2, right and Fig. 4, respectively). This reinforces the main idea of the

![Fig. 2.](image-url)
The present paper, which is to show that stochastic reorientation models provide an alternative description to cell migration patterns. Note that in Fig. 2 we have fitted simultaneously the MSD and the VACF just with the four parameters of our model (see Fig. Caption); the agreement found can be interpreted as a verification of the Green–Kubo formula (20). More free parameters were used in the original (Takagi et al., 2008) for the same fitting, though it must be noted that the authors there also fitted the behavior of the cell acceleration as a function of the velocity (this cannot be done in our case since we are not using a Langevin description).

(c) Kurtosis: Another parameter of special interest is the kurtosis $\langle \dot{r}^4(t)/\langle \dot{r}^2(t) \rangle^2 \rangle$. It is known that this parameter must tend to 3 for a diffusive regime characterized by a Gaussian distribution. This is not true, however, for anomalous diffusion, since in that case the classical central limit theorem no longer applies. Unfortunately, the values of the kurtosis as a function of time are only available for two of the references discussed throughout this paper (Dieterich et al., 2008; Takagi et al., 2008). In the former (Dieterich et al., 2008) the asymptotic behavior of the kurtosis was found to tend asymptotically to the value 2.3. This has been interpreted as a departure from Brownian models, and so by the existence of a long-range memory effect. In the latter (Takagi et al., 2008), the tendency of the kurtosis towards 3 as $t \to \infty$ is found (note that in Takagi et al., 2008 the kurtosis is defined as $\langle \dot{r}^4(t)/\langle \dot{r}^2(t) \rangle^2 \rangle - 3$ so there is a shift in the results there). However, this asymptotic value is reached from above which is a departure from the OU process, where the kurtosis is a monotonically increasing function (Uhlenbeck and Ornstein, 1931).

In fact, we show now that both results can be explained (qualitatively, at least) from our reorientation models. Using the same calculation methods as reported in Campos and Méndez (2009) and Peruani and Morelli (2007), the fourth moment $\langle \dot{r}^4(t) \rangle$ can be calculated for the stick–slip model and the two other models considered here. However, the corresponding expressions one obtains are huge so we cannot reproduce them here; we just show the corresponding plots for some arbitrary parameters (Fig. 5). Note that the two models in Fig. 1 yield now different results, since (as stated above) the microscopic differences between them become apparent for higher-order moments.

Fig. 5 confirms that the kurtosis exhibits a non-monotonic behavior which is in a qualitative agreement with the results in Dieterich et al. (2008) and Takagi et al. (2008). Also, the stick–slip model presented here is able to explain asymptotic deviations from the value 3 expected for Gaussian variables (Fig. 5, right). In summary, this shows that there is no need to account for memory effects or long tailed distributions to explain such results. We must admit that a quantitative agreement with experimental data, using the parameter values obtained from the fitting in Fig. 2, has been tried and the results are quite poor, specially for short times (Fig. 6). Note, however, that the models proposed in Dieterich et al. (2008) and Takagi et al. (2008) were not able to fit the short and intermediate-time regime for the kurtosis, neither. This is in fact a usual problem, which may be due to the fact that discretization in data acquisition (both in time and space) strongly affect the kurtosis values, specially for small times. This is another aspect of extreme importance which requires further investigation (Potdar et al., 2009).

(d) Speed and velocity distributions: In the reorientation models considered here the speed probability distribution is a model input, so it cannot be used to fit the experimental data; however, it is convenient to include some comments about it. Usually, a standard assumption is to consider a Gaussian distribution for $v$. However, several works have reported experimental evidence of non-Gaussian (exponential or power-law-like) tails for these distributions. To be more specific, one of the components of the velocity (for example, $v_x$) is observed to exhibit this unexpected behavior (Takagi et al., 2008; Selmecci et al., 2005; Upadhyaya et al., 2001; Thurner et al., 2003; Czirok et al., 1998). This is clearly a departure from the OU process, where all the components of the velocity are Gaussian variables. Instead, note this does not happen for the reorientation models explored here. If one wants to find for the reorientation model the probability distribution $Q(v_x)$ of the velocity component $v_x = \nu \cos \theta$, this can be done easily from probability theory. The formula (Rohatgi, 1976)

$$Q(z) = \int_{-\infty}^{\infty} f(x) \frac{1}{|x|} dx$$

Fig. 3. MSD comparison between the results from the stick–slip model presented here (solid lines), the models A and B (dotted lines) and the experimental data extracted from Dieterich et al. (2008) for Madin–Darby canine kidney cell strains of the wild type and NHE-deficient (symbols, see legend). The fitted parameters are, for the stick–slip model, $\beta = 1.08 s^{-1}, \beta_2 = 0.19 s^{-1}, \sigma = 0.36, v_0 = 1.19 \mu m/s$ (wild-type) and $\beta = 1.02 s^{-1}, \beta_2 = 0.27 s^{-1}, \sigma = 0.30, v_0 = 1.42 \mu m/s$ (NHE-deficient). For the other case we obtain $\beta = 0.70 s^{-1}, \sigma = 0.017 s^{-1}, \langle v \rangle = 0.83 \mu m/s$, $\langle v^2 \rangle = 2.92 \mu m^2/s^2$ (wild-type) and $\beta = 0.14 s^{-1}, \sigma = 0.011 s^{-1}, \langle v \rangle = 0.49 \mu m/s$, $\langle v^2 \rangle = 1.01 \mu m^2/s^2$ (NHE-deficient).

Fig. 4. VACF comparison between the results from the stick–slip model presented here (solid lines), the models A and B (dotted lines) and the experimental data for AX4 strains of Dictyostelium cells obtained in Selmecci et al. (2008) (asterisks). The fitted parameters are, for the stick–slip model, $\beta = 1.205 s^{-1}, \beta_2 = 1.08 s^{-1}, \sigma = 0.41, v_0 = 5.43 \mu m/s$. For the models A and B we obtain $\beta = 2.07 s^{-1}, \sigma = 0.10 s^{-1}, \langle v \rangle = 4.95 \mu m/s$, $\langle v^2 \rangle = 63.2 \mu m^2/s^2$. 

$$\langle \dot{r}^4(t)/\langle \dot{r}^2(t) \rangle^2 \rangle$$
allows one to find the probability distribution of the product $Z = XY$ of two random variables $X$, $Y$, with $f(x,y)$ being the joint distribution of these two variables. For any reorientation model without a preferential direction of motion, it can be assumed that the distribution of the $\theta$ values is uniform, which leads to a probability distribution $\sim (1 - \cos^2 \theta)^{-1/2}$ for the variable $\cos \theta$.

Using this result and assuming a Gaussian distribution for the speed, with a characteristic exponent $g$, Eq. (22) can be applied. It leads to the analytical solution

$$Q(v_x) \sim e^{-\gamma v_x^2/2} K_0 \left( \frac{\gamma v_x^2}{2\pi} \right)$$  \quad (23)$$

where $K_0(\cdot)$ denotes the modified Bessel function of zeroth order. Clearly, the expression (23) exhibits a non-Gaussian behavior for small and intermediate times (in general it resembles very much an exponential decay). At long times, however, it will tend to a Gaussian distribution, as the central limit theorem requires. So, we see that, provided our window of experimental times cannot reach the asymptotic regime $t \gg K^{-1}$ an exponential decay of the velocities will be found, as observed in Selmecki et al. (2005) and Czirok et al. (1998). This cannot explain, however, why power-law tails have been observed in other experimental works. In those works, it seems that cell interactions were important (this happens for example in the case of Hydra cells studied in Upadhyaya et al., 2001), while our approach here is restricted to free cells. In other cases, velocities are computed from measuring the cell displacements at scales smaller than the cell size (Takagi et al., 2008). For those observations cell deformation is expected to play a major role in the short-time dynamics, so biomechanical
5. Conclusions

A reorientation model, equivalent to the stick–slip process introduced recently in Selmecki et al. (2008) has been presented and analytically explored from a mesoscopic CTRW formalism in order to show its interest for cell migration. In addition, we have provided an extensive review of those experimental works on cell migration where departures from the OU process have been detected. By analyzing one by one the main statistical parameters measured in those works we have been able to show that our stochastic reorientation models are compatible with most of the results observed. Again, we do not mean that motion of cells must correspond necessarily to such microscopic patterns. Basically the contribution of our work lies in showing that many departures from the OU process, which had been previously attributed to memory kernels or other complex effects, also admit a simpler interpretation. Since biological systems have multiple and complex internal states, we think there is a great value in identifying non-uniqueness in models that can describe experimental data. Models based on memory kernels, as discussed in Section 2, rely on the hypothesis that the interactions between the cells and the surrounding media leads to the appearance of some specific memory effects, which requires justification from biophysical (mechanistic) models. The approach presented here avoids that drawback. Besides, the results obtained in Selmecki et al. (2008) and Takagi et al. (2008), where the Langevin equation (1) has been tested directly, seem to support our general idea that reorientation mechanisms models represent a more appropriate way to describe cell motion. While our approach can provide a quite general framework for the persistent motion of free cells at low densities, it is clear that more complex situations involving cell interactions or external stimuli are out of the scope of the present work. More sophisticated approaches will be required for those situations, and it is not clear yet if the CTRW formalism used here will be useful still in that case. Note, for example, that even extremely simple interactions between particles can lead to quite complicated collective and cooperative phenomena, a topic which is at present under investigation (see Ginelli et al., 2010 and the references therein).

Anyway, even for the non-interacting particles considered here many questions remain open. For example: if stochastic reorientation can really explain most of animal migration patterns, do then these patterns correspond to any functional advantage as an optimal search strategy (in the line of the results reported in Friedrich, 2008)? Is it really possible to describe the short-time dynamics of cell migration (which is rather dominated by cell deformations) just by stochastic approaches based in point particles? How does the time-discretization introduced in data analysis affect all these results? These questions are just to illustrate that understanding cell migration patterns still represents a really challenging task.

Introducing all the probability distribution functions given in the text into (13)–(16), and transforming from the real coordinates \((\mathbf{r}, t)\) to Fourier–Laplace coordinates \((\mathbf{q}, \omega, t)\), this system of equations takes the much more manageable form

\[
\mathcal{F}_1(\mathbf{q}, \omega, t) = \int_{0}^{2\pi} d\theta \frac{\beta(\theta-\theta)}{s + \beta + iq\omega \cos(\theta) + q\omega \sin(\theta)} \mathcal{F}_1(\mathbf{q}, \omega, t) + \frac{1}{2\pi}
\]

\[
\mathcal{F}_2(\mathbf{q}, \omega, t) = \frac{\beta_2}{s + \beta_2} \mathcal{F}_1(\mathbf{q}, \omega, t)
\]

\[
\hat{P}_1(\mathbf{q}, \omega, t) = \frac{1}{s + \beta_2} \mathcal{F}_1(\mathbf{q}, \omega, t)
\]

\[
\hat{P}_2(\mathbf{q}, \omega, t) = \frac{1}{s + \beta + iq\omega \cos(\theta) + q\omega \sin(\theta)} \mathcal{F}_2(\mathbf{q}, \omega, t)
\]

where we use the hat (‘) to denote the Fourier–Laplace transform. In the following, we will obviate for simplicity the explicit dependence of \(\mathcal{F}\) and \(\hat{P}\) in their variables. Now, if we assume that orientation changes are smooth we can expand (24) for \(\theta \to \theta_0\), which yields up to second order

\[
\mathcal{F}_1 = \left(1 + \sigma^2 \frac{\partial^2}{\partial \theta^2}\right) \frac{\beta_1}{s + \beta_1 + iq\omega \cos(\theta) + q\omega \sin(\theta)} \mathcal{F}_1 + \frac{1}{2\pi}
\]

\[
\mathcal{F}_2 = \frac{\beta_2}{s + \beta_2} \mathcal{F}_1
\]

These equations can be formally inverted to the real coordinates \((\mathbf{r}, t)\) but a solution for \(P_1(\mathbf{r}, t)\) and \(P_2(\mathbf{r}, t)\) cannot be explicitly found. However, the MSD is easier to compute in the Fourier–Laplace coordinates from \(<r^2(\mathbf{q})> = \frac{1}{s + \beta + iq\omega \cos(\theta) + q\omega \sin(\theta)} \hat{P}_1 + \frac{1}{2\pi}
\]

\[
\hat{P}_1 = \frac{1}{s + \beta_2 + iq\omega \cos(\theta) + q\omega \sin(\theta)} \hat{P}_2
\]

This expression, inverted from the Laplace coordinate \(s\) back to \(t\), takes finally the form given in Eq. (17). For the sake of completeness, we provide the exact expressions of the \(p_k\)'s one obtains so:

\[
p_0(t) = \frac{2\beta_1(\beta + \beta_2)^3 + 3\beta^2(2\beta + \beta_2)^2 + 3\beta_1 \beta_2 + 3\beta_2 \beta^2 \sigma^2 (\beta + \beta_2) t}{\beta^3 \sigma^4 (\beta + \beta_2)^3}
\]

\[
p_1(t) = \frac{-2\beta_2(2\beta + \beta_2)^2 + \beta_1 \beta_2 \sigma^2 (2\beta + \beta_2) + 3\beta_1 \beta_2 t}{\beta^3 \sigma^4 (\beta + \beta_2)^3}
\]

\[
p_2 = \frac{2\beta_2 \sigma^2 (2\beta + \beta_2)^2 + 2\beta_2 \beta_3 (\beta - \beta_2)}{\beta^4 \sigma^4 (2\beta + \beta_2)^3}
\]

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Appendix

The system of Eqs. (13)–(16) can be explicitly solved for some particular cases in order to reach an expression for the MSD.
\[ p_3 = \frac{2\beta_2(1 + \sigma^2) - 2\lambda_2(\beta - \beta_2)}{\beta_2^2\sigma^4(2\lambda_2 - \beta_2)} \]
\[ p_4 = \frac{2\lambda_4(\beta_2 - \beta_4)}{\beta^2\sigma^4(2\lambda_4 - \beta_2)} \]
\[ p_5 = \frac{2\lambda_5(\beta_2 - \beta_4)}{\beta^2\sigma^4(2\lambda_5 - \beta_2)} \]

(32)

References


References


References


