FROM CELL MEMBRANES TO ULTRACOLD GASES: CLASSICAL AND QUANTUM DIFFUSION IN INHOMOGENEOUS MEDIA

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Sciences

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Outlook

- A. Classical
 - Anomalous diffusion in biological systems
 - Kinetic Ising Models
- B. Quantum
 - Quantum Kinetic Ising Models
 - Quantum Brownian Motion

Ordinary Random Walk

Time-averaged mean squared displacement along a single trajectory:

T-MSD
$$(t_{lag} = m\Delta t) = \frac{1}{N-m} \sum_{i=1}^{N-m} (x (t_i + m\Delta t) - x (t_i))^2$$

Ensemble-averaged mean squared displacement over J trajectories:

$$\text{E-MSD}(t_{lag} = m\Delta t) = \frac{1}{J} \sum_{j=1}^{J} \left(x_j \left(t_i + m\Delta t \right) - x_j \left(t_i \right) \right)^2$$

An ordinary RW is

- diffusive: MSD~ t^{β} , with $\beta=1$
- ergodic: T-MSD and E-MSD coincide at long times
- stationary: MSD independent of the total observation time (at large times)





Diffusion in structured media



A random walker may be strongly affected by:

- repeated interactions
- transient binding
- clustering
- crowding
- multi-scale, or scale-free, disorder

Live cell membrane



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- multi-scale, or scale-free, disorder

Trans-membrane receptors

Single particle tracking of pathogen-recognizing receptors on live cellular membranes

(M. García-Parajo's group @ ICFO)





60 frames/s 20nm position accuracy

Manzo, Torreno-Pina, PM, Lapeyre, Lewenstein & García-Parajo, Phys. Rev. X (2015)



Different scaling of T-MSD and E-MSD → weak ergodicity breaking!

Time&Ensemble averaged MSD: TE-MSD ~ $D_{TE}(T) \cdot t_{lag}$

The TE-MSD depends on the total observation time T - non-stationary!



Continuous-Time Random Walk

- CTRW: a fat-tailed distribution of waiting times $\sim t^{-1-\beta}$ with $\beta \le 1$ (so that $< t_{wait} > = \infty$) induces non-stationary (thus non-ergodic) subdiffusion with E-MSD $\sim t^{\beta}$ and TE-MSD $\sim D_{TE}(T)^* t_{lag}$, with $D_{TE} \sim T^{\beta-1}$
- Widely used model for transport in disordered media (initially developed for amorphous solids)

Montroll & Weiss, 1965; Montroll & Scher, 1973

10¹

10⁰

escape time (s)

10⁻¹

 10^{-2}

• However, are trapping events present in the ICFO experiment?

only 5% of the trajectories contain events compatible with transient trapping

and excluding these trajectories yields a very similar E-MSD exponent $\boldsymbol{\beta}$

Δx, Δy (μm) 500 nm -0 50 nm -0. 500 nm 2 0 1 3 Δ t (s) Rтн cdf 0.4 500 nm 0

0.2

 For CTRW, the long-time dynamics is dominated by anomalous trapping events, so that the escape-time distribution at long times becomes independent of the trapping radius R_{TH}

500 nm

Strongly varying diffusivity

- Maps of receptor motion on the cell membrane highlight the presence of patches with strongly varying diffusivity
- Many possible reasons: crossing regions of low/high viscosity, transient binding, clustering, ...
- Employ a likelihood-based Bayesian algorithm to detect *time-dependent changes of diffusivity within a single trajectory*:



Theoretical model

- Ordinary Brownian motion with a diffusivity that varies randomly, but it's constant on time intervals with random duration (or patches with random sizes)
- Assume a distribution of diffusion coefficients $P_D(D) \sim D^{\sigma-1}e^{-D/b}$ fast decay a and a conditional distribution of transit times $P_{\tau}(\tau|D) \sim D^{\gamma}e^{-\tau D^{\gamma}/k}$ mean transit each area be
- Three possible regimes:

(0) $\gamma < \sigma$: the long-time dynamics is compatible with regular Brownian motion ($\beta = 1$) (I) $\sigma < \gamma < \sigma + 1$: the average transit-time diverges \rightarrow non-ergodic sub-diffusion ($\beta = \sigma/\gamma$) (II) $\gamma > \sigma + 1$: both $<\tau >$ and $< (r_{area})^2 >$ diverge \rightarrow non-ergodic sub-diffusion ($\beta = 1 - 1/\gamma$)

PM, Manzo, Torreno-Pina, García-Parajo, Lewenstein & Lapeyre, Phys. Rev. Lett. (2014)

es randomly.



mean transit time $\sim D^{-\gamma}$ each area has radius r $\sim(\tau D)^{-1/2}$

Comparison with experiments



Instead, mutated receptors with impaired function (Δ_{rep}) may be best simulated with $\gamma < \sigma$, i.e., they perform usual ergodic and diffusive Brownian motion

Transport properties strongly linked to molecular function

Manzo, Torreno-Pina, PM, Lapeyre, Lewenstein & García-Parajo, Phys. Rev. X (2015)

Include receptor/membrane interactions?

[see Poster P.03 "Random walks in Random Environments" by M.-A. García-March]

Model the membrane as an Ising lattice: $H = -J \sum \sigma_i \sigma_{i+1}$

Link diffusivity to local membrane state (e.g. faster diffusion on 1 background)

Allow walker to locally modify the membrane state

$$= -J \sum_{i} \sigma_{i} \sigma_{i+1} \qquad \sigma_{i} = \uparrow, \downarrow$$

Kinetic Ising Model for the membrane

• Kinetic Ising Model:
$$\dot{P}(\sigma, t) = \sum_{\sigma'} [w(\sigma' \to \sigma)P(\sigma', t) - w(\sigma \to \sigma')P(\sigma, t)]$$

w: transition rate

• Interacting membrane: $w(\sigma_i \to -\sigma_i) = \frac{\alpha}{2} \left[1 - \frac{\gamma}{2} \sigma_i (\sigma_{i-1} + \sigma_{i+1}) \right]$

- $\alpha/2$: single spin-flip rate $\gamma>0$: IFM $\gamma<0$: AFM
- Detailed balance at equilibrium: $P_{eq}(\sigma_1, \dots, \sigma_i, \dots, \sigma_n)w(\sigma_i \to -\sigma_i) = P_{eq}(\sigma_1, \dots, -\sigma_i, \dots, \sigma_n)w(-\sigma_i \to \sigma_i)$
- $P_{eq}(\sigma_1, \dots, \sigma_i, \dots, \sigma_n) \propto \exp[\beta J \sigma_i(\sigma_{i-1} + \sigma_{i+1})]$ $q_i(t) = \langle \sigma_i(t) \rangle$ $r_{i,k}(t) = \langle \sigma_i(t) \sigma_k(t) \rangle$

•
$$P(\sigma,t) = \frac{1}{2^N} \sum_{\{\sigma'\}} (1 + \sigma_1 \sigma'_1) \dots (1 + \sigma_N \sigma'_N) P(\sigma',t) = \frac{1}{2^N} \left\{ 1 + \sum_i \sigma_i q_i(t) + \sum_{i \neq k} \sigma_i \sigma_k r_{i,k}(t) + \dots \right\}$$

- Recursive system of differential equations: e.g., $\alpha^{-1}\dot{q}_i(t) = -q_i(t) + \gamma/2[q_{i-1}(t) + q_{i+1}(t)]$
- **Exact time-dependent solutions** may be found for: infinite lattice, single spin fixed, (time-delayed) spin correlations, spin systems in a magnetic field, ...

R. Glauber, Time-dependent statistics of the Ising model, J. Math. Phys. (1963)

Kinetic Ising Model for membrane+walker

• Master equation: $\dot{P}(\sigma, x, t) = [\gamma_s \mathcal{L}_s + \gamma_w \mathcal{L}_w] P(\sigma, x, t)$

s: spins (=membrane) w: walker x: position of the walker

- The walker may act as a localized potential for the spins, or it may change the tunneling between the neighboring spins
- If either the spins or the walker dynamics is fast, we may use an adiabatic approximation

• E.g.,
$$\gamma_s \gg \gamma_w \rightarrow \mathcal{L} = \gamma_s \left[\mathcal{L}_s + \frac{\gamma_w}{\gamma_s} \mathcal{L}_w \right]$$
 and $P(\sigma, x, t) \sim P_w(x, t|\sigma) P_s^{(eq)}(\sigma|x)$

- polaronic behavior (rapid formation of a dressing cloud, dragged along by the walker)
- In the opposite limit instead, the walker spreads very fast, and will act as a diffuse potential (mean-field) for the membrane

(Quantum) Kinetic Ising Models

- Ansatz: $P(\sigma,t) = \sqrt{P_{eq}(\sigma)\phi(\sigma,t)}$
- The KIM master equation may be written as $\dot{\phi}(\sigma,t) = -\sum_{\sigma'} H_{\sigma\sigma'}\phi(\sigma',t)$ $H_{\sigma\sigma'} = \sum_{\sigma''} w(\sigma \to \sigma'')\delta_{\sigma\sigma'} - w(\sigma' \to \sigma)\sqrt{\frac{P_{\text{eq}}(\sigma')}{P_{\text{eq}}(\sigma)}}$
- If the KIM satisfies a detailed-balanced condition, then H is a (real) symmetric matrix, so that the KIM-ME may be seen as a Schrödinger equation in imaginary time, which converges exponentially to the thermal equilibrium solution. Diagonalization of H is then equivalent to the complete solution of the KIM-ME
- Classical spins may be promoted to non-commuting Pauli matrices to obtain quantum models
- Quantum states built from thermal states of classical Hamiltonians, e.g., $|\Psi\rangle = \frac{1}{\sqrt{Z_N}} \sum_{\sigma} e^{-\beta H(\sigma)/2} |\sigma\rangle$ fulfill the area law in any dimension, even at criticality, and can be represented efficiently as matrix product states (MPS), or projected entangled pair states (PEPS)

Augusiak, Cucchietti, Haake & Lewenstein, New J. Phys. (2010)

Quantum Brownian Motion

- A small system interacting with a large thermal bath: $H = H_S + H_B + H_I$
- Simplest interaction: $H_I = -\sum_k \kappa_k x_k x_k$

x_k: position of the kth oscillator
κ_k: coupling constant
x: position of the system

- ME for the reduced density matrix: $\dot{\rho}_S(t) = -\frac{1}{\hbar^2} \int_0^t ds \operatorname{Tr}_B[H_{\mathrm{I}}(t), [H_{\mathrm{I}}(s), \rho(s)]]$
- Born approx.: $\rho(t) \simeq \rho_S(t) \otimes \rho_B(0)$
- Markov approx.: the bath evolves on timescales much faster than the system, and as such effectively retains no memory of the system's dynamics
- The spectral density contains all details of the bath-system coupling: $J(\omega) \equiv \sum_{k} \frac{\kappa_k^2}{2m_k\omega_k} \delta(\omega \omega_k)$

Born-Markov QME

- Weak-coupling limit (second-order perturbation in H_I)
- Ohmic spectral density with Lorentz-Drude cut-off: $J(\omega) = \frac{m\gamma\omega}{\pi} \frac{\Lambda^2}{\omega^2 + \Lambda^2}$
- Consider a particle of mass m in a harmonic trap of frequency Ω . In the high-temperature limit $k_B T \gg \hbar \Lambda \gg \hbar \Omega$ one obtains the celebrated Caldeira-Leggett QME:

$$\dot{\rho}_S = -\frac{i}{\hbar} [H_{\rm sys}, \rho_S] - \frac{i\gamma}{2\hbar} [x, \{p, \rho_S\}] - \frac{m\gamma k_B T}{\hbar^2} [x, [x, \rho_S]]$$

momentum damping normal diffusion

 A quantum stochastic process is said to be Markovian only if the underlying noise is white, and if the process is described by a time-independent ME of the Lindblad form

• Note that there exists an *exact* solution of the QBM problem, which however has a timedependent Liouvillian. Strictly-speaking, QBM is not a quantum Markov process.

Impurities in an inhomogeneous 1D BEC



Catani et al., Phys. Rev. A (2012)

How to treat a spatially-dependent profile of the bath? Coupling which is a non-linear function of the system's position:

Aspect ratio of the impurity, $\ln(\delta_p^2/\delta_x^2)$, for a quadratic coupling with the environment:

How to deal with the low temperatures of a quantum gas? Non-Markovian effects will become important here!

And what if the spectral density is not Ohmic? (i.e., the noise is not white)



PM, Lampo, Wehr & Lewenstein, Phys. Rev. A (2015)

Open questions

A. Classical

• Anomalous diffusion in biological systems

Q: mechanisms leading to sub- or super-diffusion, Levy flights, non-stationarity and non-ergodicity?

• Kinetic Ising Models

Q: mechanisms generating interactions between walkers and substrates?

B. Quantum

Quantum Kinetic Ising Models

Q: novel exactly or efficiently solvable quantum Hamiltonians?

Quantum Brownian Motion

Q: importance of non-Markovian effects at low temperatures?

Conclusions

 Walkers on largely heterogeneous substrate can exhibit strongly subdiffusive and non-ergodic behavior, even without transient immobilization

PM, Manzo, Torreno-Pina, García-Parajo, Lewenstein & Lapeyre, Phys. Rev. Lett. (2014) Manzo, Torreno-Pina, PM, Lapeyre, Lewenstein & García-Parajo, Phys. Rev. X (2015)

- We are working towards describing membrane-walkers interactions by means of Kinetic Ising Models (KIMs)
- KIMs may be generalized to the quantum domain
- Quantum Brownian Motion in presence of an inhomogeneous background may be described by couplings which are polynomial functions of the system variables PM, Lampo, Wehr & Lewenstein, Phys. Rev. A (2015)

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