

# Discussion session on Stochastic processes

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# Outlook

## A. Classical

- Anomalous diffusion in biological systems
- Kinetic Ising Models

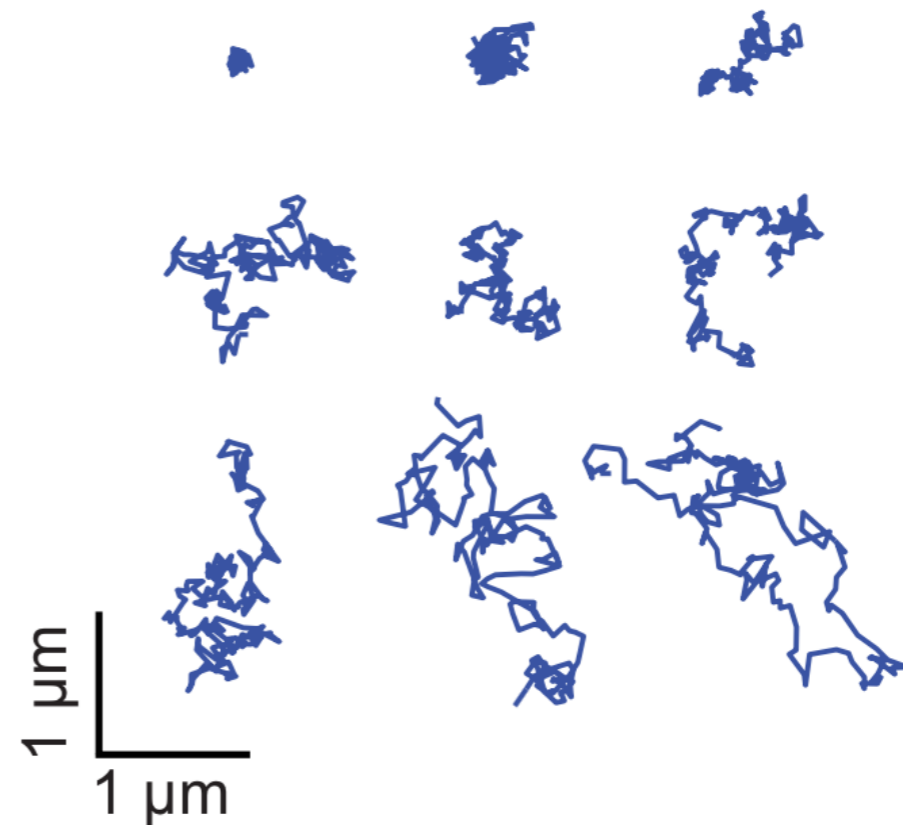
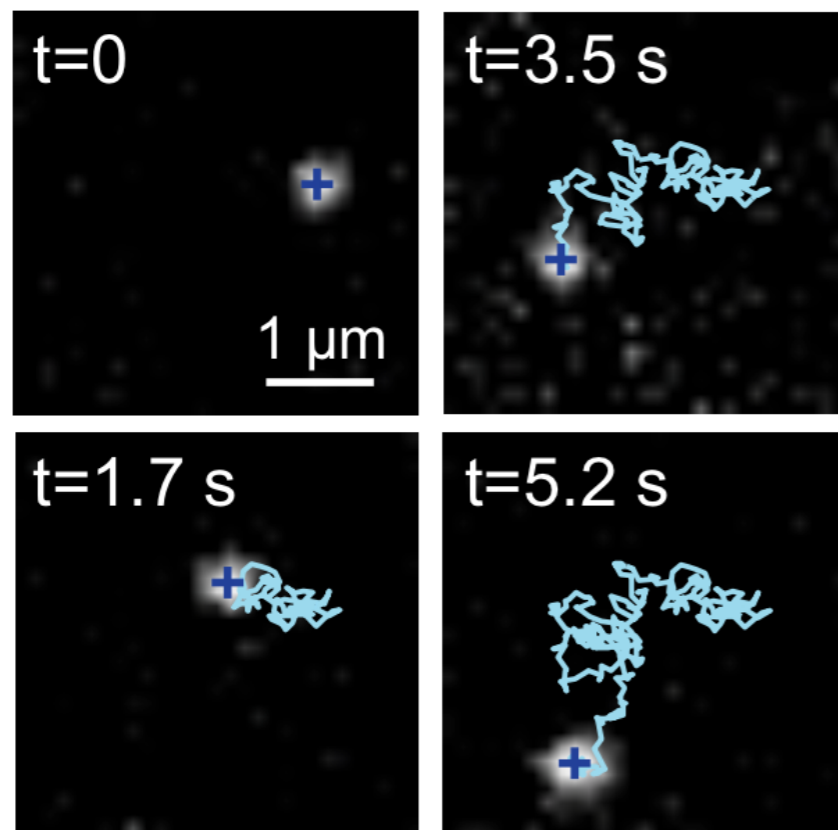
## B. Quantum

- Quantum Kinetic Ising Models
- Quantum Brownian Motion

# Diffusion in a complex medium

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

(collaboration with M. G. Parajo group @ ICFO)



60 frames/s  
20nm position accuracy

Sampling of  $J=600$  trajectories

$x_j$ : position of the  $j$ -th receptor ( $1 \leq j \leq J$ ) sampled at  $N$  discrete times

time-averaged mean squared displacement:

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

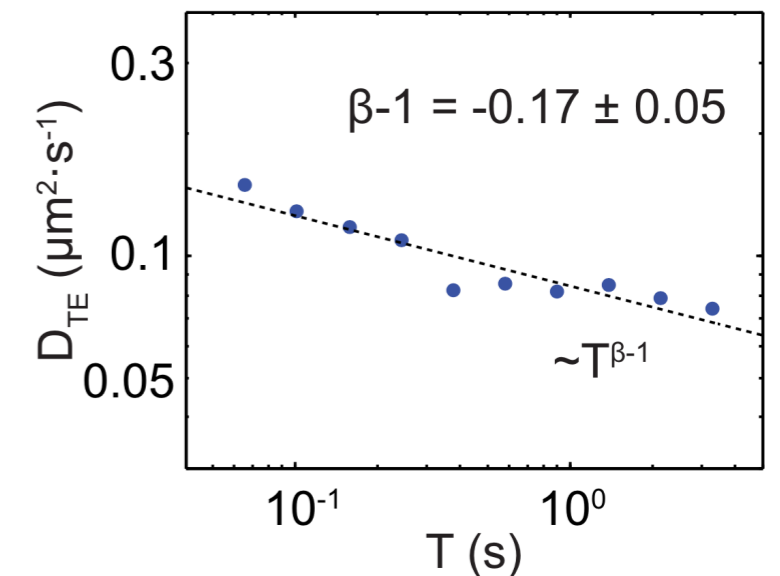
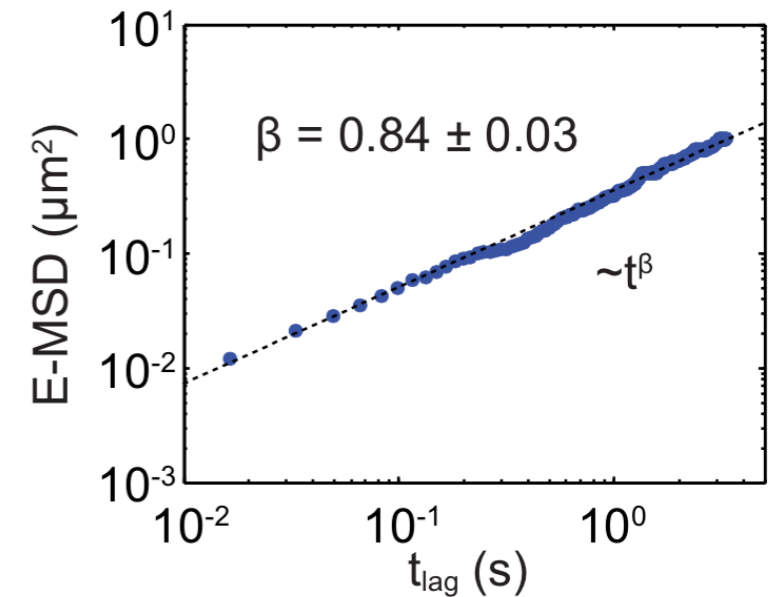
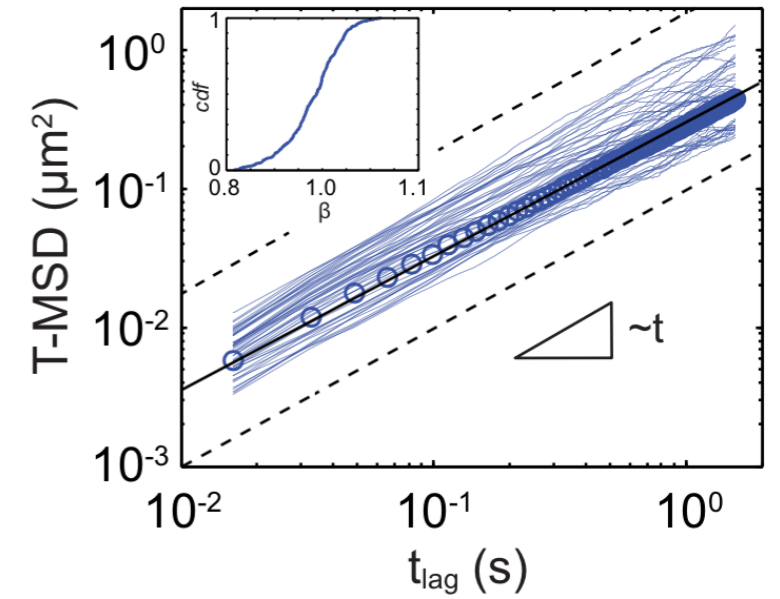
ensemble-averaged mean squared displacement:

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

T-MSD scales differently from E-MSD  $\Rightarrow$  weak ergodicity breaking!

$$\text{TE-MSD}(t_{lag}, T) = \frac{1}{J} \frac{1}{\frac{T}{\Delta t} - m} \sum_{i=1}^{\frac{T}{\Delta t} - m} \sum_{j=1}^J (x_j(t_i + m\Delta t) - x_j(t_i))^2$$

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



# Continuous-Time Random Walk

- CTRW: a fat-tailed distribution of waiting times  $\sim t^{-1-\beta}$  with  $\beta \leq 1$ , so that the average waiting time is infinite, induces non-stationary (thus non-ergodic) subdiffusion, with E-MSD  $\sim t^\beta$ , and TE-MSD scaling as  $D_{TE}(T) \cdot t_{lag}$ , with  $D_{TE} \sim T^{\beta-1}$

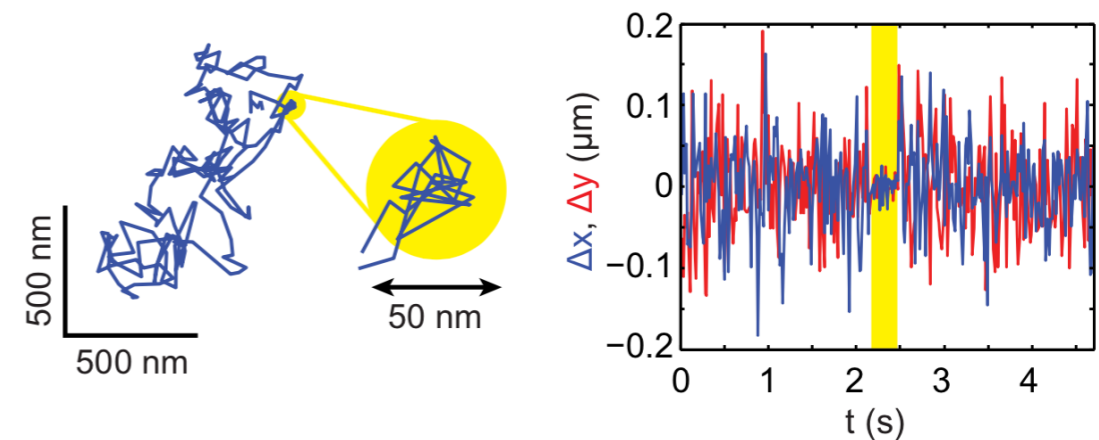
- Widely used model for transport in disordered media (initially developed for amorphous solids)

Montroll & Weiss, 1965; Montroll & Scher, 1973

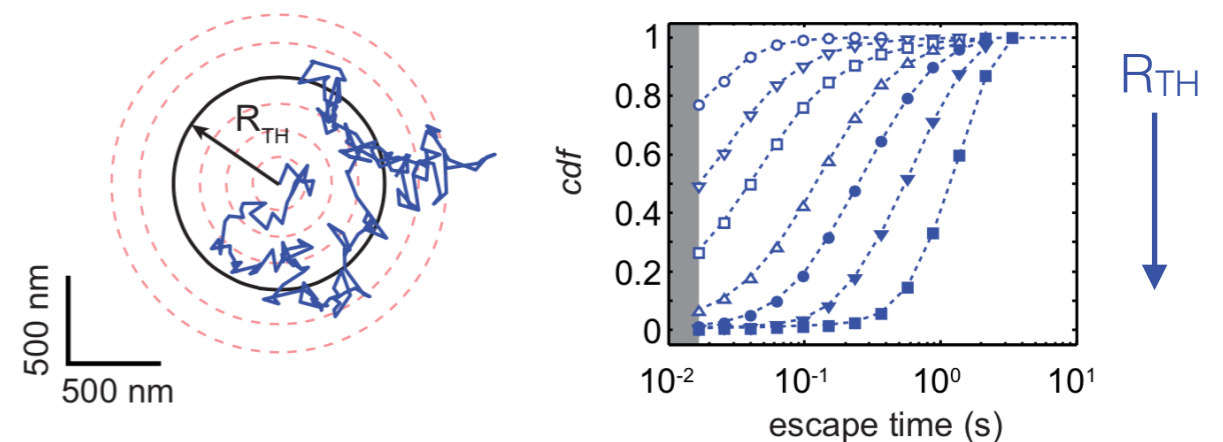
- However, *are trapping events present in the ICFO experiment?*

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

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

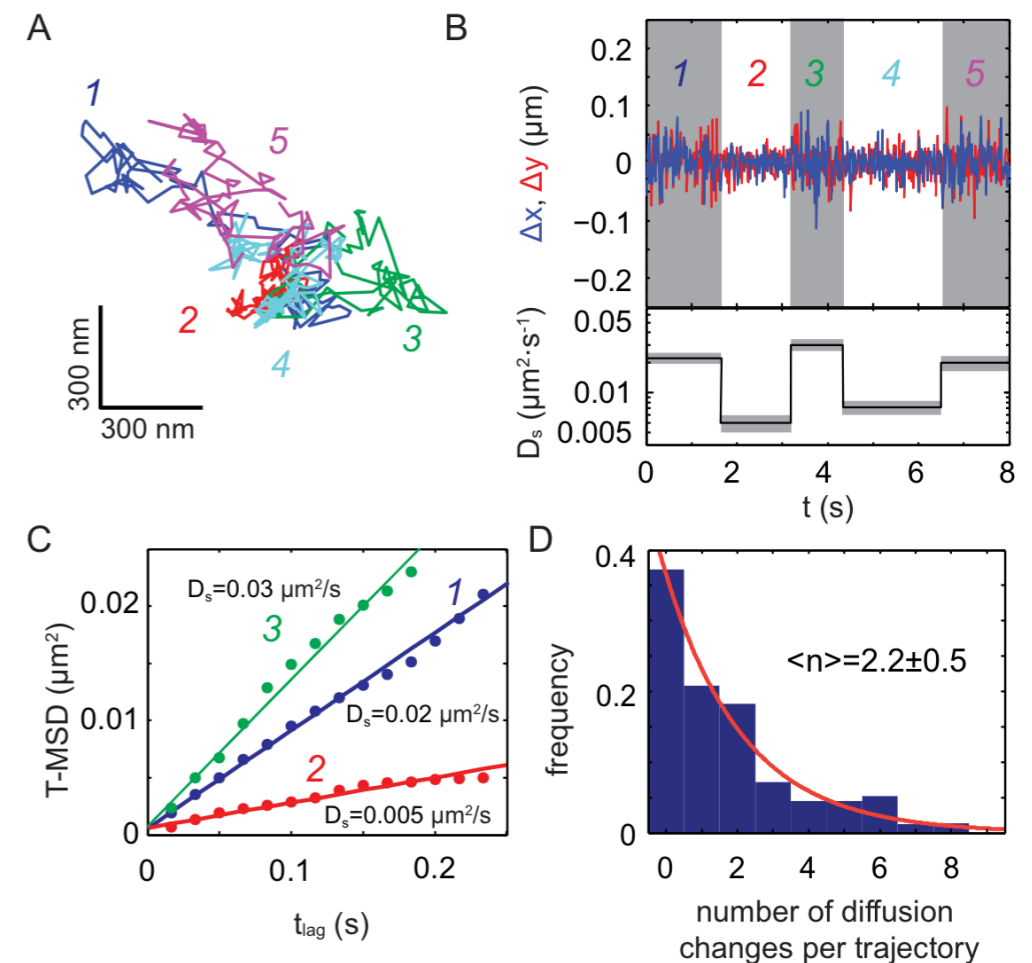
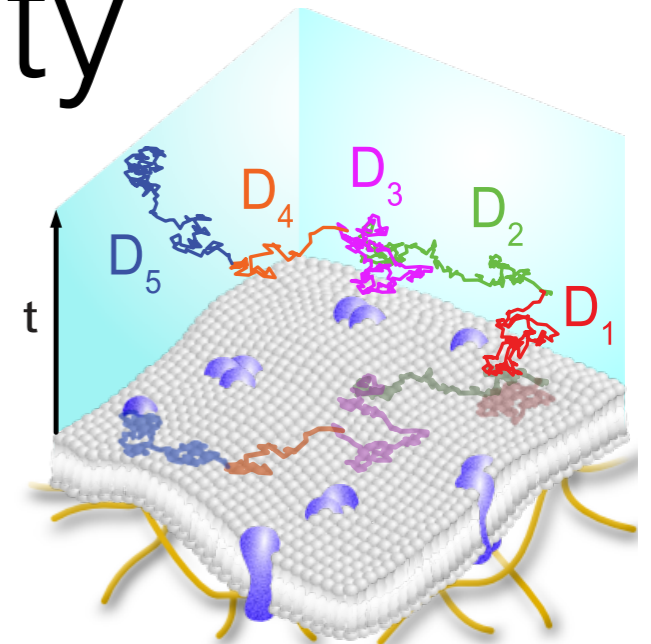


- For CTRW, the long-time dynamics is dominated by anomalous trapping events, so that the escape-time distribution should be independent of the trapping radius  $R_{TH}$

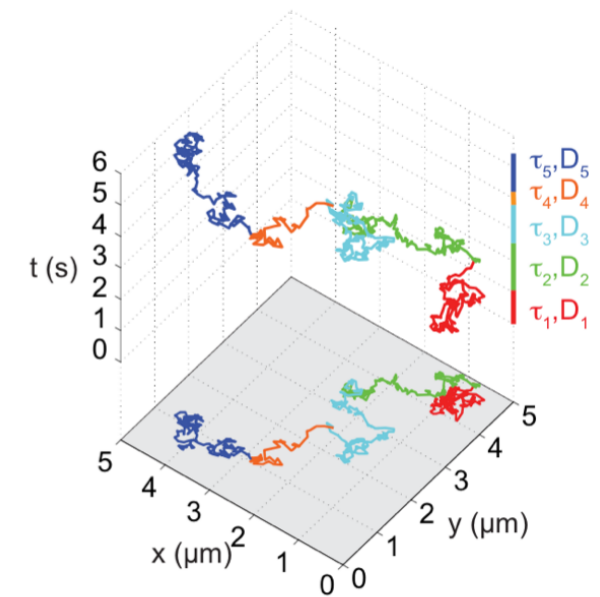


# Strongly varying diffusivity

- 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



# Theoretical model



- Let's then consider 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  $P_D(D) \sim D^{\sigma-1} e^{-D/b}$   
and a conditional distribution of transit times  $P_\tau(\tau|D) \sim D^\gamma e^{-\tau D^\gamma/k}$

power law behaviour at small  $D$   
fast decay at large  $D$

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

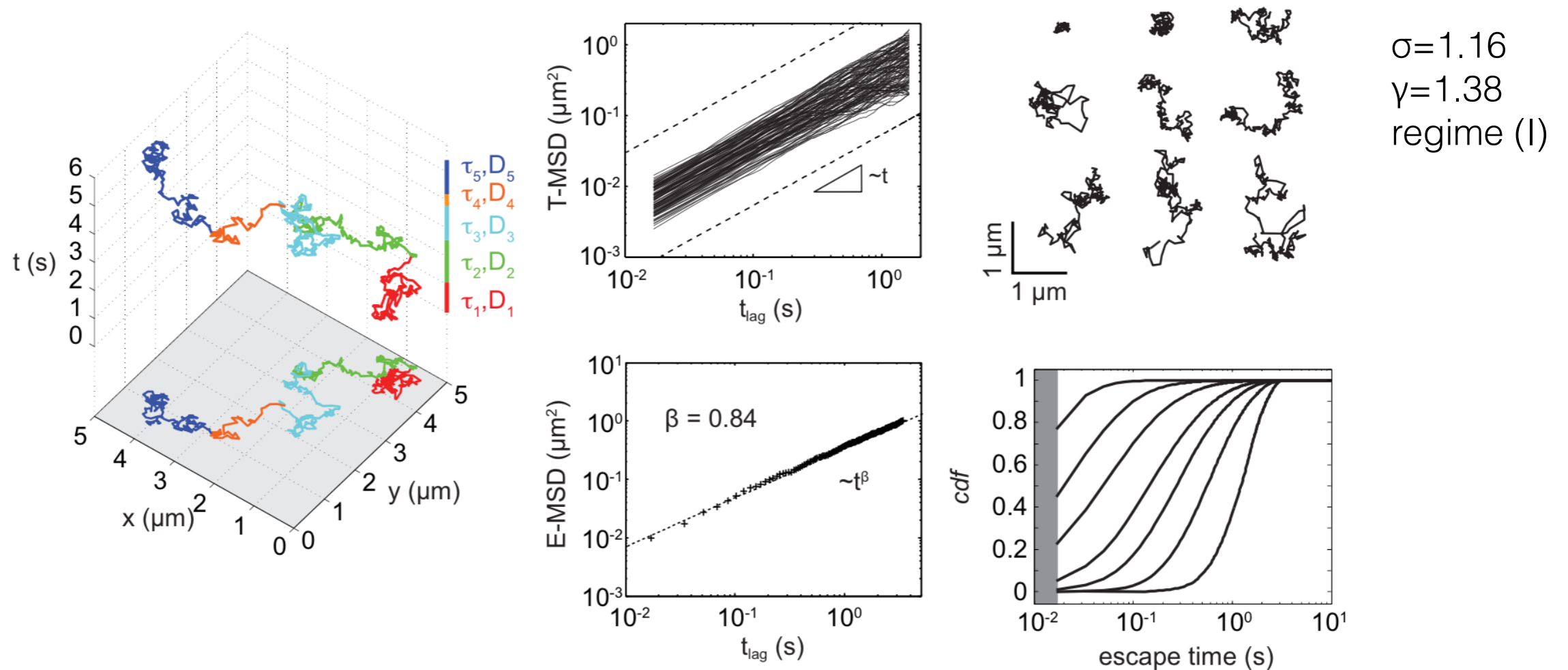
- Three possible regimes:

(0) for  $\gamma < \sigma$ , the long-time dynamics is compatible with regular Brownian motion, and  $\beta = 1$

(I) for  $\sigma < \gamma < \sigma + 1$ , the average transit-time diverges: non-ergodic sub-diffusion with  $\beta = \sigma/\gamma$

(II) for  $\gamma > \sigma + 1$ , both  $\langle \tau \rangle$  and  $\langle (r_{\text{area}})^2 \rangle$  diverge: non-ergodic sub-diffusion with  $\beta = 1 - 1/\gamma$

# Comparison with experiments



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

Transport properties strongly linked to molecular function

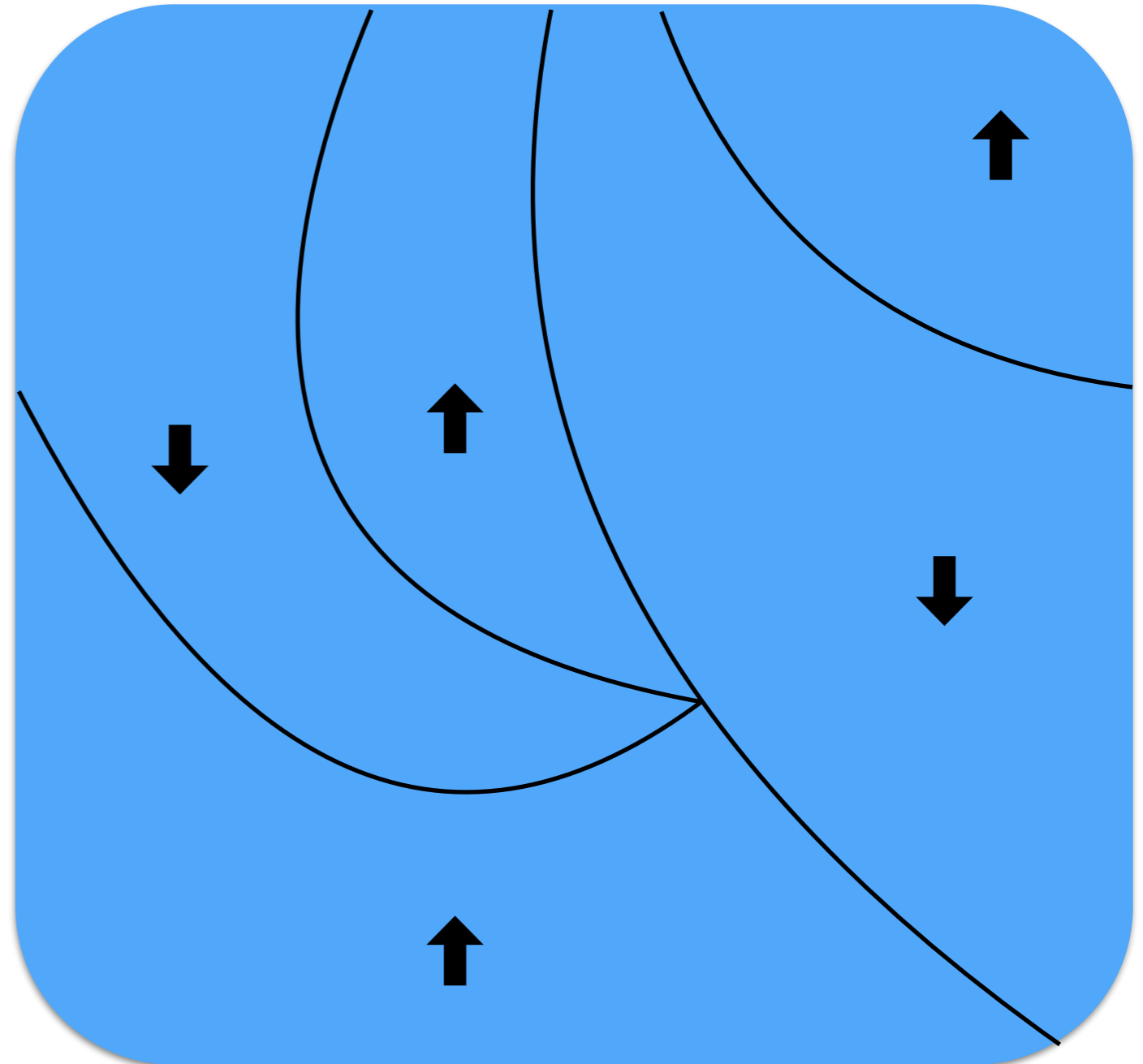


# Include receptor/membrane interactions?

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

Link diffusivity to local membrane state  
(e.g. faster diffusion on an  $\uparrow$  background)

Allow walker to locally modify  
the membrane state



# Kinetic Ising Model for the membrane

- Kinetic Ising Model:  $\dot{P}(\sigma, t) = \sum_{\sigma'} [w(\sigma' \rightarrow \sigma)P(\sigma', t) - w(\sigma \rightarrow \sigma')P(\sigma, t)]$  w: transition rate
- Interacting membrane:  $w(\sigma_i \rightarrow -\sigma_i) = \frac{\alpha}{2} \left[ 1 - \frac{\gamma}{2} \sigma_i (\sigma_{i-1} + \sigma_{i+1}) \right]$  
 $\alpha/2$ : single spin-flip rate  
 $\gamma > 0$ : IFM  
 $\gamma < 0$ : AFM
- Detailed balance at equilibrium:  $P_{\text{eq}}(\sigma_1, \dots, \sigma_i, \dots, \sigma_n) w(\sigma_i \rightarrow -\sigma_i) = P_{\text{eq}}(\sigma_1, \dots, -\sigma_i, \dots, \sigma_n) w(-\sigma_i \rightarrow \sigma_i)$
- $P_{\text{eq}}(\sigma_1, \dots, \sigma_i, \dots, \sigma_n) \propto \exp[\beta J \sigma_i (\sigma_{i-1} + \sigma_{i+1})] = \cosh[\beta J (\sigma_{i-1} + \sigma_{i+1})] \left\{ 1 + \frac{\tanh(2\beta J)}{2} \sigma_i (\sigma_{i-1} + \sigma_{i+1}) \right\}$   
 $\exp(\pm x) = \cosh(x) \pm \sinh(x)$
- $P(\sigma, t) = \frac{1}{2^N} \sum_{\{\sigma'\}} (1 + \sigma_1 \sigma'_1) \dots (1 + \sigma_N \sigma'_N) P(\sigma', t) = \frac{1}{2^N} \left\{ 1 + \sum_i \sigma_i q_i(t) + \sum_{i \neq k} \sigma_i \sigma_k r_{i,k}(t) + \dots \right\}$   
 $q_i(t) = \langle \sigma_i(t) \rangle$   
 $r_{i,k}(t) = \langle \sigma_i(t) \sigma_k(t) \rangle$
- Recursive system of differential equations:  $\alpha^{-1} \dot{q}_i(t) = -q_i(t) + \gamma/2 [q_{i-1}(t) + q_{i+1}(t)]$
- 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  $P(\sigma, x, t) \sim P_w(x, t|\sigma)P_s^{(\text{eq})}(\sigma|x)$ 
  - polaronic behavior (rapid formation of a dressing cloud, dragged along by the walker)
- In the opposite limit instead, the walker spreads very fast, and should act as some kind of diffuse potential (mean-field) for the membrane

# (Quantum) Kinetic Ising Models

- Ansatz:  $P(\sigma, t) = \sqrt{P_{\text{eq}}(\sigma)}\phi(\sigma, t)$
- The KIM master equation may be written as  $\dot{\phi}(\sigma, t) = - \sum_{\sigma'} H_{\sigma\sigma'}\phi(\sigma', t)$   

$$H_{\sigma\sigma'} = \sum_{\sigma''} w(\sigma \rightarrow \sigma'')\delta_{\sigma\sigma'} - w(\sigma' \rightarrow \sigma)\sqrt{\frac{P_{\text{eq}}(\sigma')}{P_{\text{eq}}(\sigma)}}$$
- 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:  $H_I = - \sum_k \kappa_k x_k x$

$x_k$ : position of the  $k^{\text{th}}$  oscillator  
 $\kappa_k$ : coupling constant  
 $x$ : position of the system

- ME for the reduced density matrix:  $\dot{\rho}_S(t) = -\frac{1}{\hbar^2} \int_0^t ds \text{Tr}_B [H_I(t), [H_I(s), \rho(s)]]$

- 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:  $J(\omega) \equiv \sum_k \frac{\kappa_k^2}{2m_k \omega_k} \delta(\omega - \omega_k)$

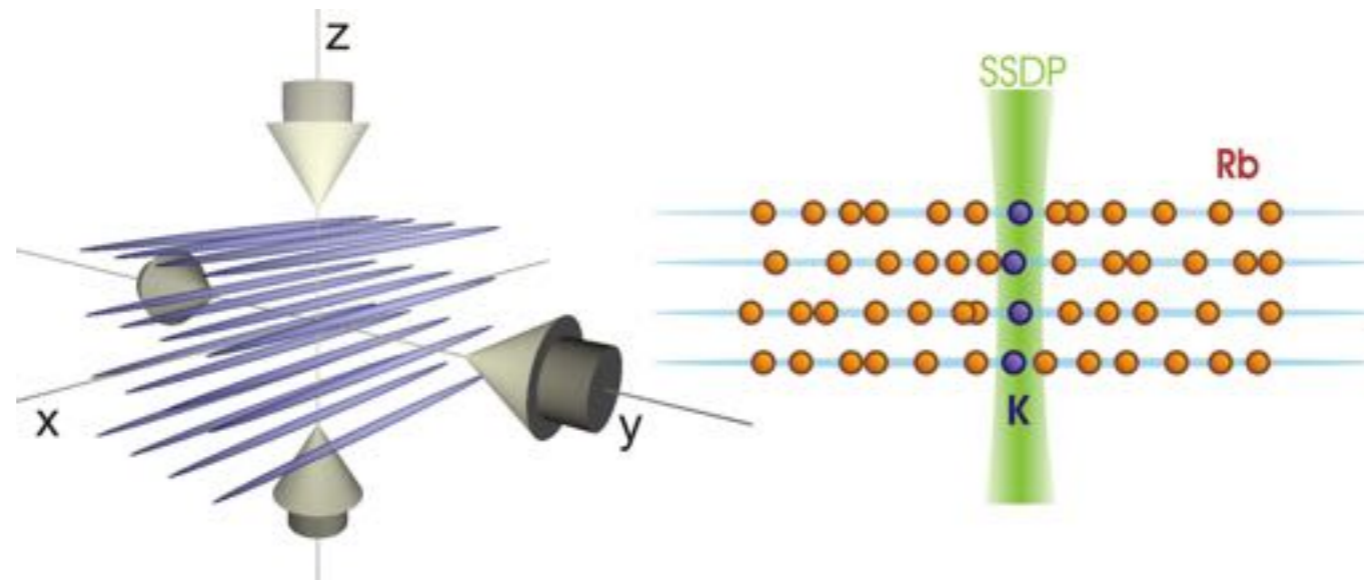
# Born-Markov QME

- 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  $\Omega$ . In the high-temperature limit  $k_B T \gg \hbar\Lambda \gg \hbar\Omega$  one obtains the celebrated Caldeira-Leggett QME:

$$\dot{\rho}_S = -\frac{i}{\hbar}[H_{\text{sys}}, \rho_S] - \underbrace{\frac{i\gamma}{2\hbar}[x, \{p, \rho_S\}]}_{\text{momentum damping}} - \underbrace{\frac{m\gamma k_B T}{\hbar^2}[x, [x, \rho_S]]}_{\text{normal diffusion}}$$

- 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 time-dependent Liouvillian. Strictly-speaking, QBM is not a quantum Markov process.

# Impurities in an inhomogeneous 1D BEC



Catani et al., PRA (2012)

# How to treat a spatially-dependent profile of the bath?

Employ a coupling which is a non-linear function of the system's position: 
$$H_I = - \sum_k \kappa_k x_k f(x)$$

PM, Lampo, Wehr & Lewenstein, PRA (2015, in press)

# How to deal with the low temperatures of a quantum gas?

Non-Markovian effects may become important here!

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

# Outlook

## 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: models for quantum diffusion in complex or inhomogeneous media?  
importance of non-Markovian effects at low temperatures?