

First-passage times of Markovian and non Markovian random walks

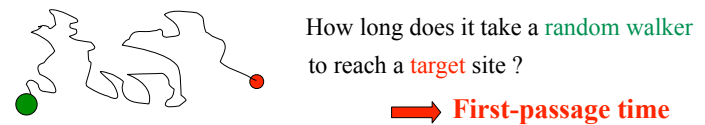
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with:
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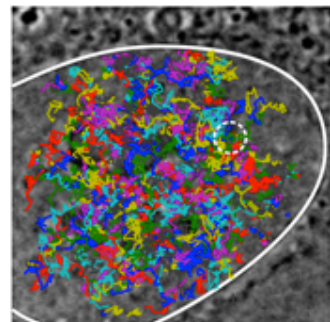
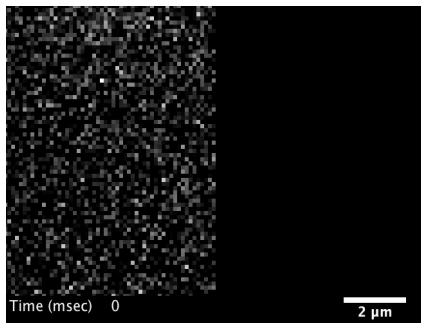
First-passage problems : First definitions



Many physical processes are controlled by **first-encounter** properties

Examples (i): diffusion limited reactions

Transcription kinetics: proteins searching for a specific target sequence on DNA

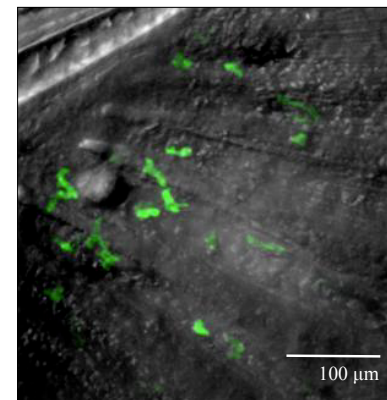


[with D. Normano, M. Dahan, Institut Curie]

- Unicity of a target gene among billions of b.p.
- Specific protein copy number can be < 10

Examples (ii): Immune cells

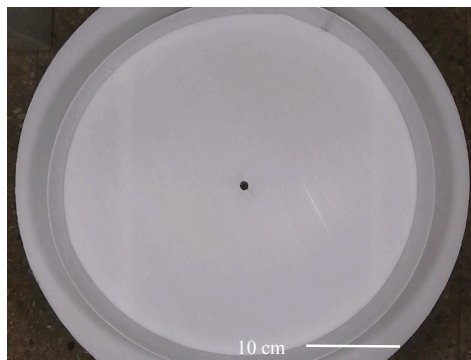
Immune cells searching for pathogens



[with M. Piel, AM Lennon, Institut Curie]

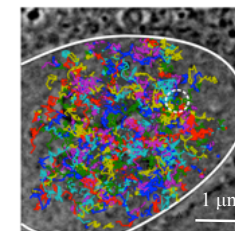
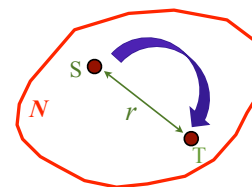
Examples (iii): ants

Ants searching for food



[data O. Feinerman, Weizmann Institute]

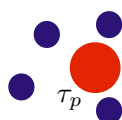
Efficiency of the search process? First-passage time statistics to a target



Main questions:

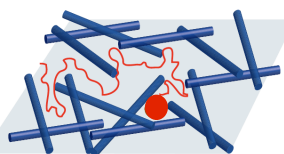
- How does the mean FPT depend on
 1. the **confining volume** N ?
 2. the **distance** r between S and T ?
 3. the **transport process** ? Effects of crowding, active/passive, anomalous transport
- Optimize ?

Diffusion in complex environments : Markov vs. non Markov



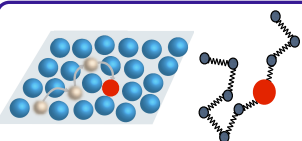
If $\tau_e \ll \tau_p$

Markov process, no memory
ex. Brownian motion



If $\tau_e \gg \tau_p$

Brownian motion in a frozen disordered medium.
Markov process, no memory
ex. rw. on percolation cluster



If $\tau_e \sim \tau_p$

N particles system is Markovian,
but single particle **non Markovian, memory**
ex. monomer of a rouse chain

Note that different from anomalous vs normal diffusion

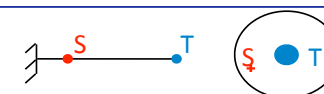
Theoretical context: Markov (almost) only [highly non-exhaustive biblio...]

Infinite space

Persistence exponents (Markov and non Markov) [Majumdar, Bray...]

(quasi) 1D geometries

[see S. Redner's book]



Singular perturbation of the Laplacian

(Brownian motion 2D and 3D) [Ward and Keller, Holcman and Schuss]

Mean return times

[Kac, Hilfer, Blumen...]

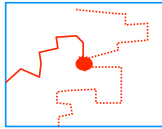
$$\langle \mathbf{T} \rangle_{\text{return}} = 1/P_{\text{stat}} = N$$



MFPT averaged over starting point

No information on the target: average over starting point (GMFPT)

For general scale invariant (d_w, d_f) random walks with $\langle \Delta r^2 \rangle \propto t^{2/d_w}$:



$$\langle \bar{T} \rangle \sim \begin{cases} N & \text{if } d_w < d_f \text{ non compact (transient)} \\ N \ln(N) & \text{if } d_w = d_f \\ N^{d_w/d_f} & \text{if } d_w > d_f \text{ compact (recurrent)} \end{cases}$$

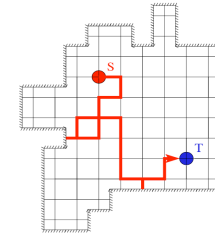
Markov: [Montroll 1965 (Euclidean lattices), ben Avraham 2005 (fractals)...]

➡ **Optimal strategy:**
less compact do better = minimize overlap

Effect of initial position ?

Outline

I/ Reminder :
First-passage time statistics in complex media
(Markovian scale-invariant processes)



Main ingredients

(i) **Markov process** : “Renewal equation” relates

the propagator $P(\mathbf{r}, t | \mathbf{r}')$

the first-passage time density $F(\mathbf{r}, t | \mathbf{r}')$

$$P(\mathbf{r}_T, t | \mathbf{r}_S) = \int_0^t F(\mathbf{r}_T, t' | \mathbf{r}_S) P(\mathbf{r}_T, t - t' | \mathbf{r}_T) dt'$$

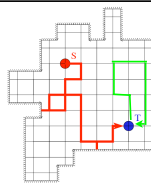
first visit of T at t'
return at T in $t - t'$

Exact expression: $\langle T \rangle = N(H(\mathbf{r}_T | \mathbf{r}_T) - H(\mathbf{r}_T | \mathbf{r}_S))$ [Noh & Rieger (2004)]

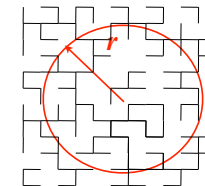
where $H(\mathbf{r} | \mathbf{r}') = \int_0^\infty (P(\mathbf{r}, t | \mathbf{r}') - 1/N) dt$ is the **pseudo-green function**

(ii) **large volume asymptotics** : $\langle T \rangle \simeq N(G_0(0) - G_0(r))$

where $G_0(r) = \int_0^\infty P_0(r, t) dt$ is the **infinite-space Green function**



(iii) Scale invariance



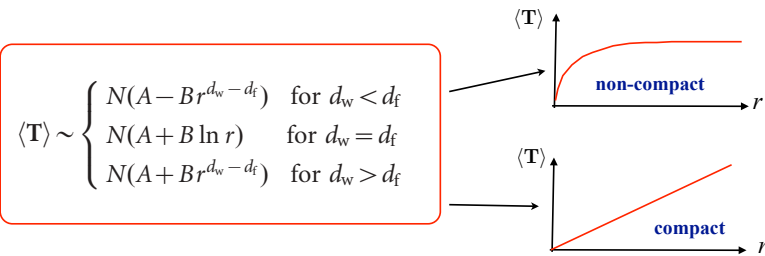
- number of sites enclosed in a **circle of radius r** : $M_r \propto r^{d_f}$
where d_f is the **fractal dimension** of the medium
- mean square displacement $\langle \Delta r^2 \rangle \propto t^{2/d_w}$
where d_w is the **dimension of the walk**
- standard **scaling assumption of the infinite-space propagator** :

$$P_0(\mathbf{r}, t | \mathbf{r}') \sim t^{-d_f/d_w} \Pi\left(\frac{|\mathbf{r} - \mathbf{r}'|}{t^{1/d_w}}\right)$$

[ben-Avraham and Havlin, (2000)]

General scaling of the MFPT

[Condamin et al Nature (2007)]

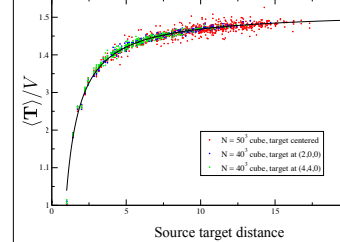


- A and B depend **only** on the infinite-space scaling
- **Linear dependence on the volume N**
- **non-compact** exploration ($d_w < d_f$): memory of the initial position **lost**
- **compact** exploration ($d_w \geq d_f$): the **initial** position **always matters**

Numerics

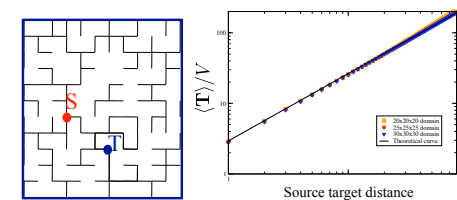
Non compact:

ex: Brownian diffusion in 3D

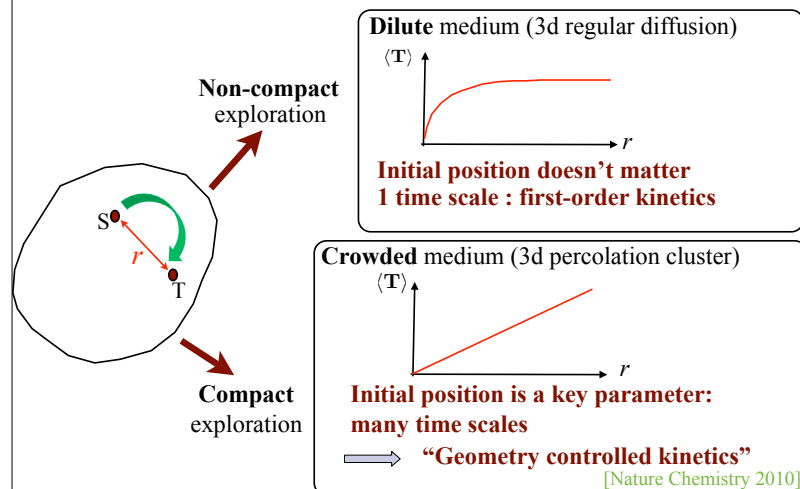


Compact:

ex: Diffusion on percolation cluster



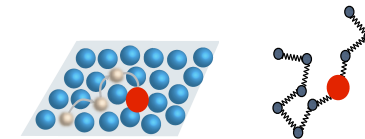
Reaction kinetics in complex media ?



Is intracellular transport in the nucleus compact or non compact ?

Outline

II/ General (Gaussian) non Markovian processes

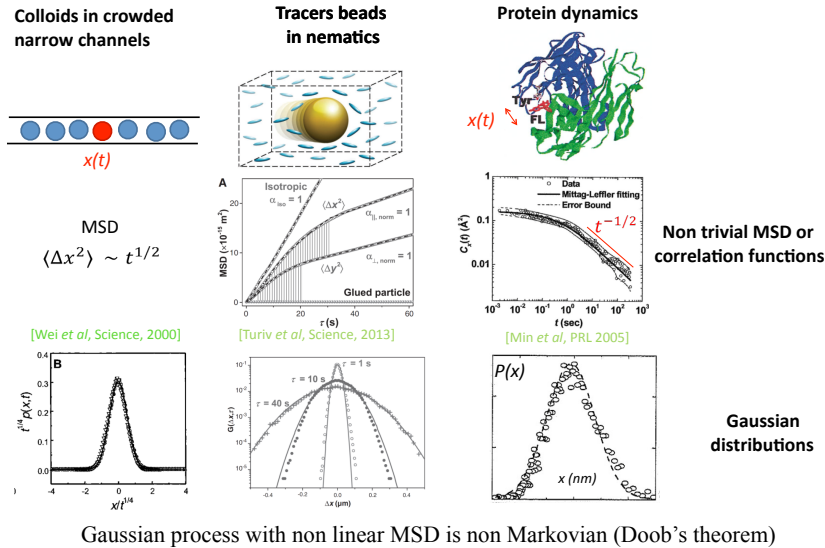


H: Gaussian process $x(t)$ with stationary increments: fully defined by

$$\psi(\tau) = \langle [x(t + \tau) - x(t)]^2 \rangle$$

here assumed given (obtained typically from single particle tracking)

Examples of non Markovian Gaussian processes



Main ingredients (i)

(i) **Generalized Renewal equation** relates

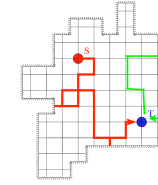
the 1 point pdf $P(x, t)$

the first-passage time density $F(\tau)$

$$p(T, t) = \int_0^t d\tau F(\tau) p(T, t | \text{FPT} = \tau)$$

first visit of T at τ

return at T in $t - \tau$

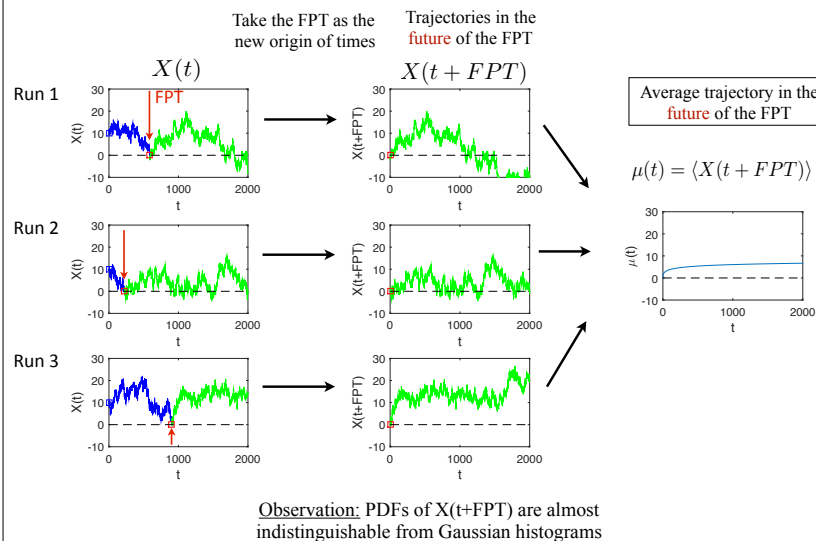


Exact expression: $\frac{\langle T \rangle}{V} = \int_0^\infty dt [q_\pi(t) - p(T, t)]$

with the conditional 1 point pdf q_π of the process in the future of the FPT

$$q_\pi(t) = \int_0^\infty d\tau p(T, t + \tau | \text{FPT} = \tau) F(\tau)$$

Process in the future of the FPT



Main ingredients (ii)+(iii)

(ii) **Large volume asymptotics**

(iii) **Gaussian approximation of q_π**

$$\langle T \rangle = V \int_0^\infty dt \frac{e^{-\mu(t)^2/2\psi(t)} - e^{-x_0^2/2\psi(t)}}{[2\pi\psi(t)]^{1/2}}$$

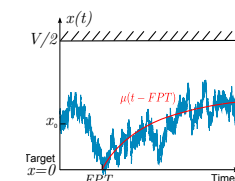
where $\mu(t)$ is the mean trajectory in the future of the FPT, defined by

$$\int_0^\infty \frac{dt}{\sqrt{\psi(t)}} \left\{ [\mu(t + \tau) - \mu(t)] K(t, \tau) e^{-\mu(t)^2/2\psi(t)} - x_0 [1 - K(t, \tau)] e^{-x_0^2/2\psi(t)} \right\} = 0,$$

$$K(t, \tau) = [\psi(t + \tau) + \psi(t) - \psi(\tau)] / [2\psi(t)]$$

Can be solved (at least numerically) + exact asymptotics for simple $\psi(t)$

- mean **future** trajectory matters
- defined by MSD function only
- Markov approx. $\mu(t) = 0$ **overestimates** MFPT



Results: Fractional Brownian motion

MSD

$$\langle [x(t) - x(t')]^2 \rangle = K(t - t')^{2H} \quad 0 < H < 1$$

Used in finance, biophysics...

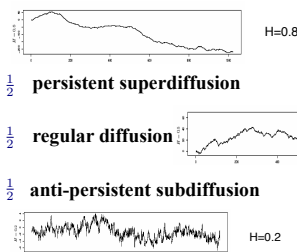
Limit of single file, Rouse, Edwards-Wilkinson

Model with **long-range correlated** increments :

Take the derivatives with respect to t and t' :

$$\langle \dot{x}(t) \dot{x}(t') \rangle = K 2H(2H - 1) \frac{1}{(t - t')^{2-2H}}$$

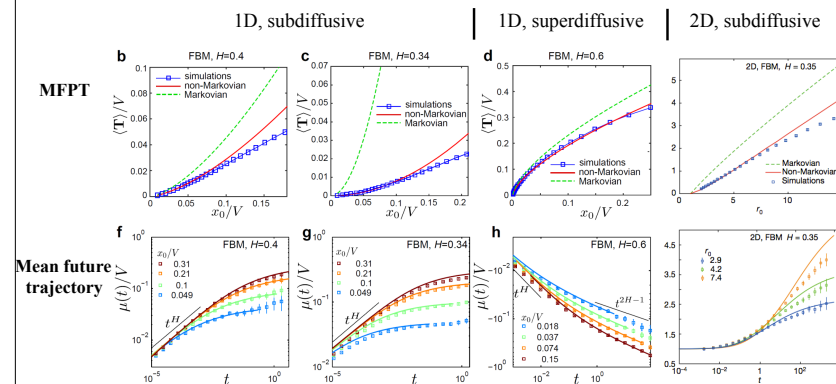
$\begin{cases} > 0 \text{ if } H > \frac{1}{2} & \text{persistent superdiffusion} \\ = 0 \text{ if } H = \frac{1}{2} & \text{regular diffusion} \\ < 0 \text{ if } H < \frac{1}{2} & \text{anti-persistent subdiffusion} \end{cases}$



Results: Fractional Brownian motion

[Nature 2016]

$$\langle [x(t) - x(0)]^2 \rangle = K t^{2H}$$



Results: diffusion in viscoelastic fluids

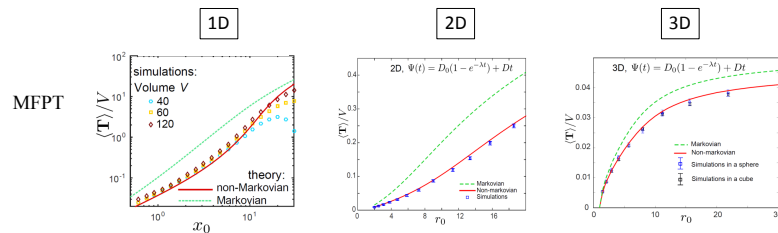
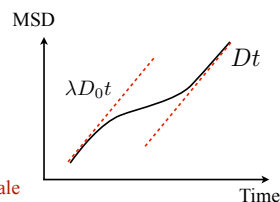
[Nature 2016]

$$\langle [x(t) - x(0)]^2 \rangle \simeq D_0(1 - e^{-\lambda t}) + Dt$$

Simple model for

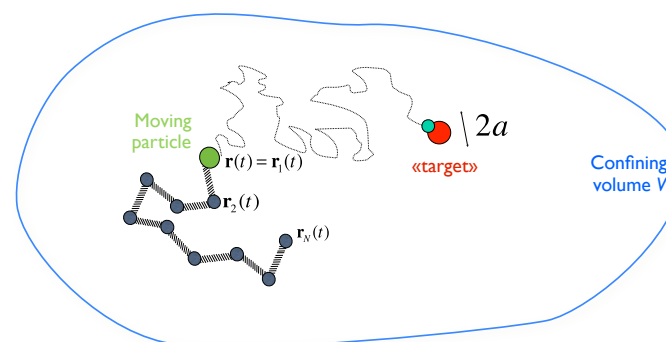
- Tracers in nematics [Turiv, 2013]
- diffusion in Maxwell fluids [Grimm et al, Soft Matt 2011]
- diffusion in polymer solutions [Ochab-Marcinek 2011]

Diffusive at short times, diffusive at long times, single time scale



➡ Even in the case of a MSD with only **1 relaxation time**, **memory** effects are **large**

Physical picture: the Rouse chain



The evolution of $\mathbf{r}_1(t)$ depends on the whole configuration $\{\mathbf{r}_1(t), \mathbf{r}_2(t), \dots, \mathbf{r}_N(t)\}$

➡ $\mathbf{r}_1(t)$ is a **non-Markovian variable**

amenable to Markovian problem (but higher dimensional)

Non Markovian theory

[Nature Chemistry 2012]

$$\langle T \rangle = V \int_0^\infty dt \left[P(x_1 = 0, t | \pi, 0) - P(x_1 = 0, t | \text{ini}, 0) \right]$$

$$\pi(\{x_i\})$$

Probability density of the configuration $\{x_i\}$
at the very instant of reaction

How can we calculate $\pi(\{x_i\})$?

- **Wilemski-Fixman approx**: replace π by the **equilibrium** distribution

$$\pi(\{x_i\}) = P_{\text{stat}}(\{x_i\} | x_1 = 0)$$

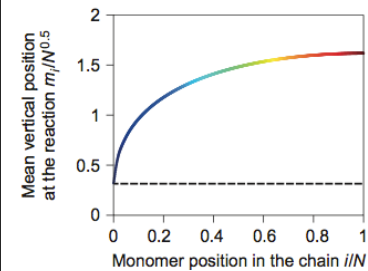
- **Our approach**: find the **Gaussian** distribution that is the closest from π

Solve closure equations for :

m_i average positions of monomers at first contact

Configurations at first contact

[Nature Chemistry 2012]



Configurations at first contact
(non-Markovian theory)

Equilibrium conformations



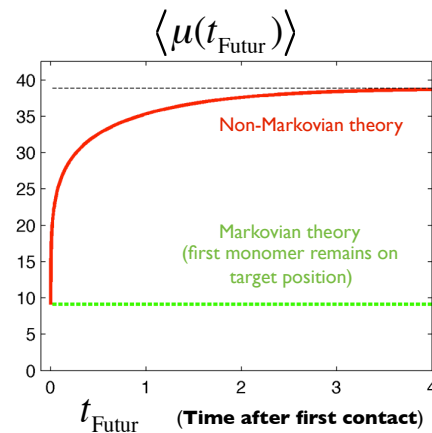
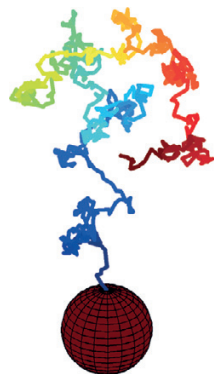
Elongated shape at first contact \rightarrow faster kinetics / Markovian theory

2 equivalent descriptions

[Nature Chemistry 2012]

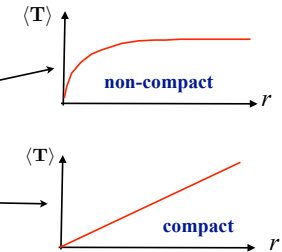
Positions of all the non-reactive
monomers at first contact

Average path after first contact



Finally : General scaling of the MFPT

$$\langle T \rangle \sim \begin{cases} N(A - Br^{d_w - d_f}) & \text{for } d_w < d_f \\ N(A + B \ln r) & \text{for } d_w = d_f \\ N(A + Br^{d_w - d_f}) & \text{for } d_w > d_f \end{cases}$$



True for Markov and non Markov

Non trivial even if it looks trivial:

- the Markov approx. $\mu = 0$ can be very wrong (scaling wise)
- 2 or 3 length scales in the problem
- prefactor can depend a lot on non Markovian effects

Finally

- if no information on the target : less compact walks do better
- if the target is close : compact walks can yield short time scales

Thanks

O. Bénichou, T. Guerin, N. Levernier