



# Diffusion and response of Brownian particles in confined structures (restricted RWs)

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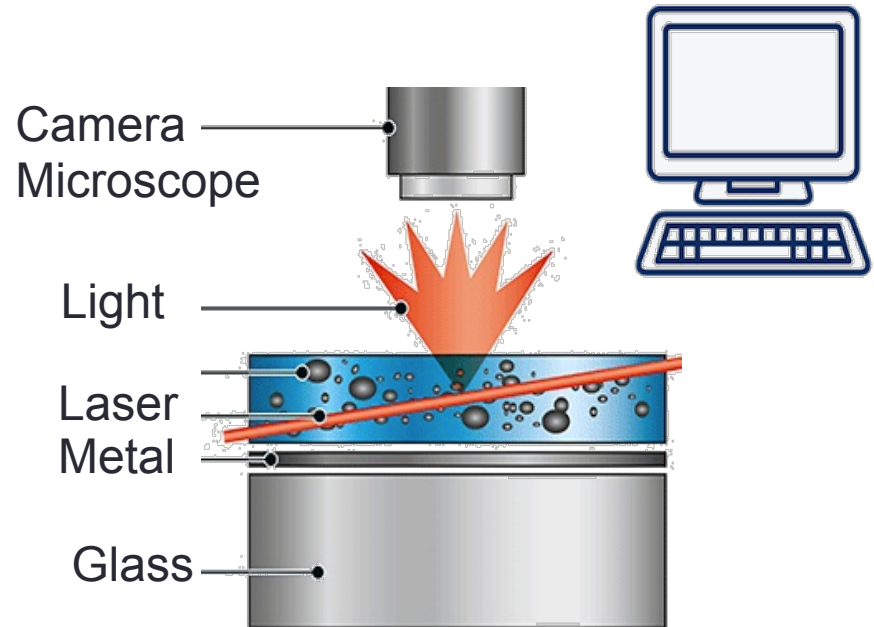
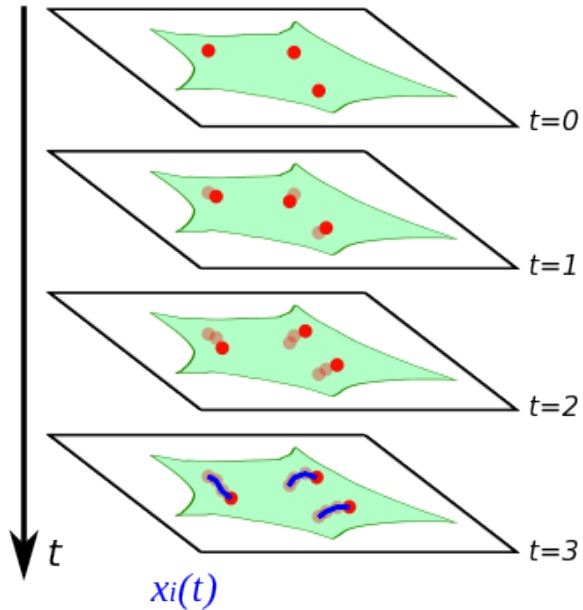
*Meeting Tematico ISC sulla diffusione, Firenze 27-28 sett. 2016*



## Outline of this talk

- Single Particle tracking technique
- Nano-Micro Confinement  
(size and dimensional reduction)
- Confined Random Walks
- Central Limit Theorem: Violations
- Standard versus Anomalous
- Example of Diffusion on:  
Branched structures,  
Fractal Trees, Channels

# Single molecule tracking



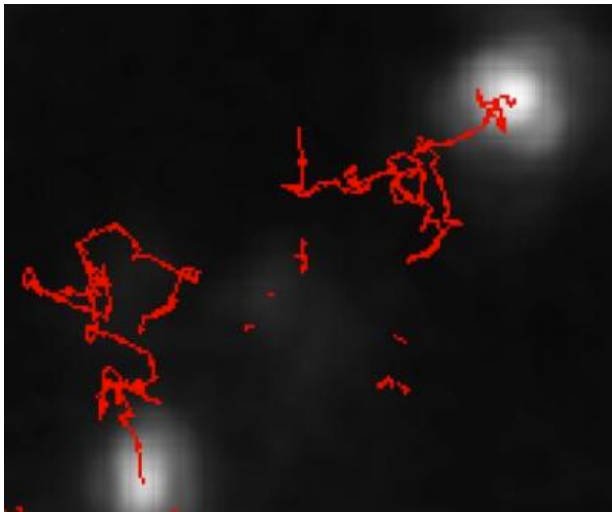
Ultra-Microscope + Laser illumination

CCD camera + Software

**Particle Position**

**method to study**

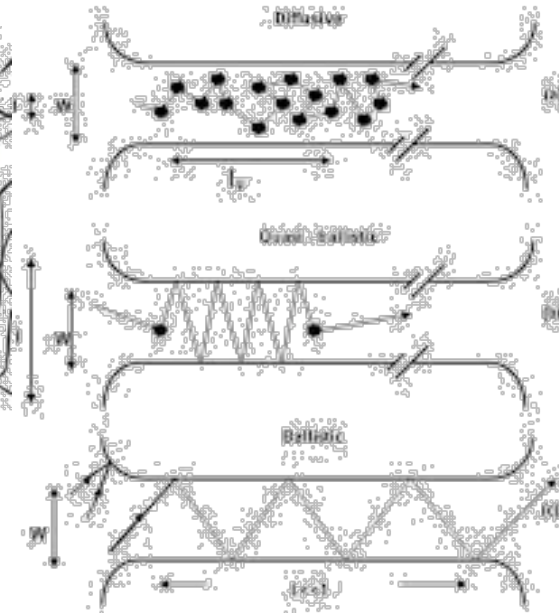
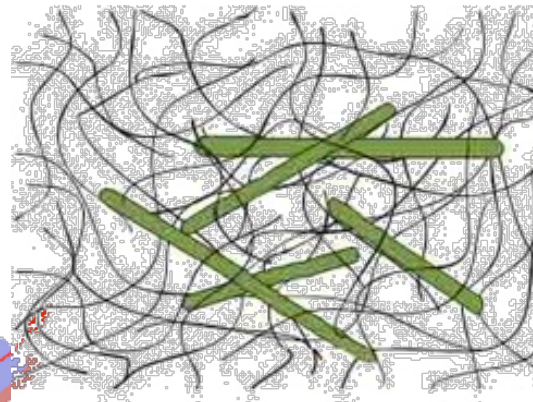
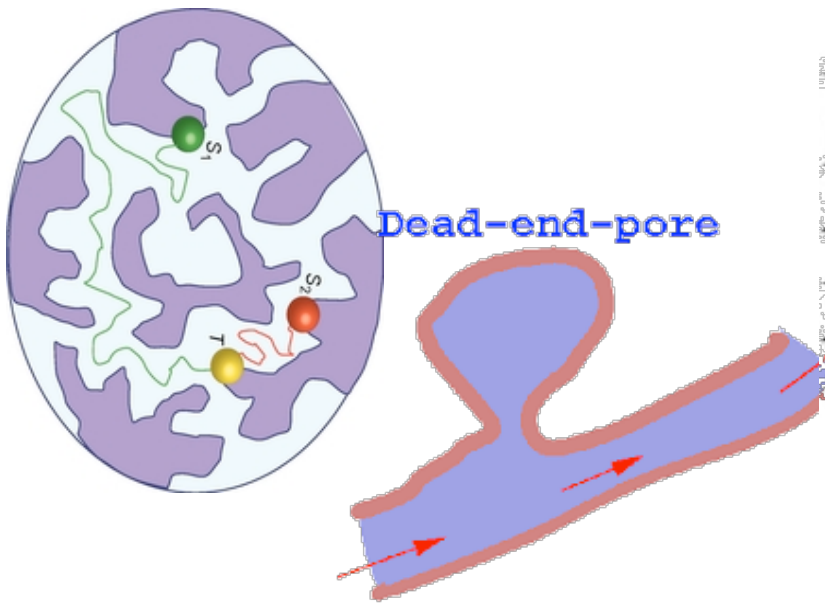
**Nano-Micro Confinement**



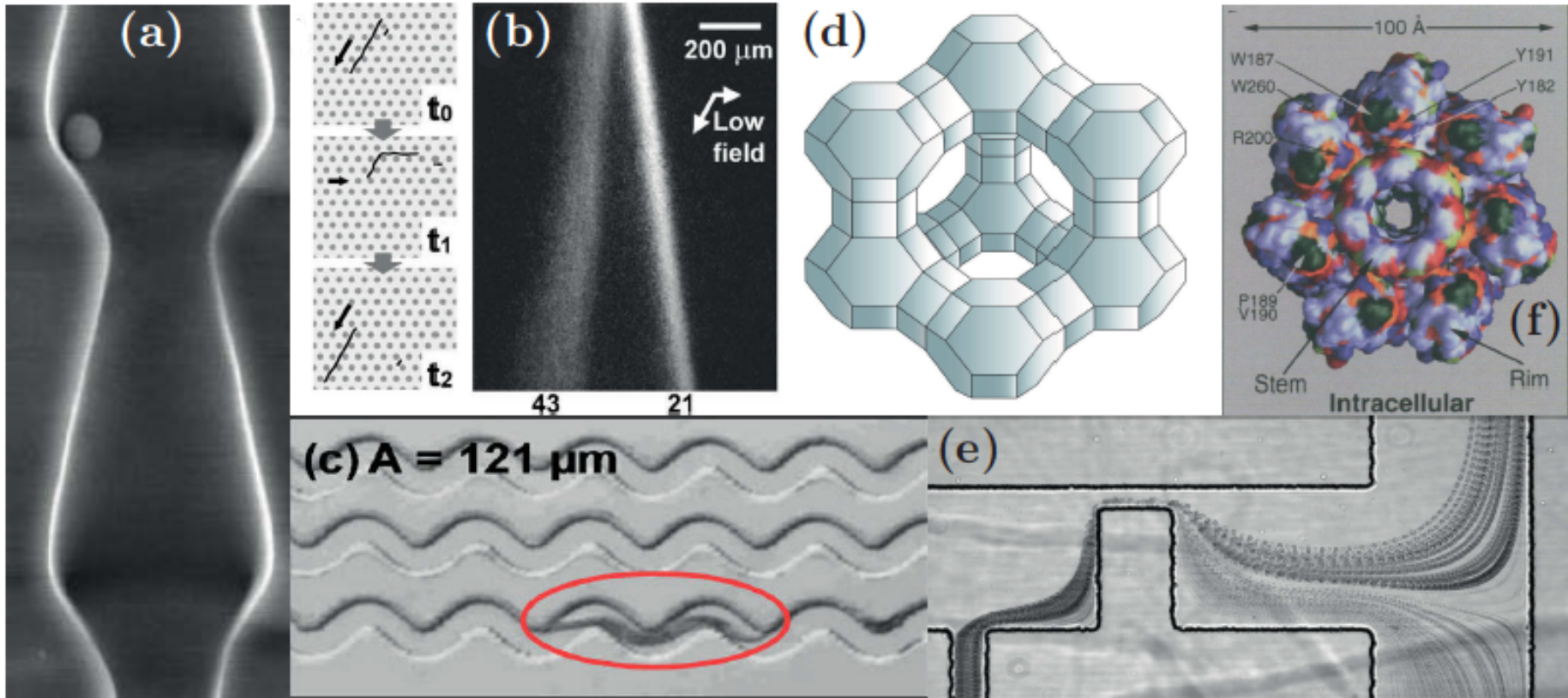
# Nano-Micro confinement

## PHENOMENOLOGY of GEOMETRY CONTROLLED TRANSPORT

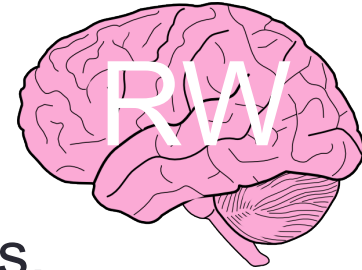
- ◆ **Biology:** transport in tissues, targeting, microtubules of cytoskeleton, cell motility
- ◆ **Chemistry:** controlled reaction, filtering
- ◆ **Physics:** porous media, low-dim transport, microfluidic
- ◆ **Nanotechnology:** nanodevices, chemical delivery
- ◆ **Mathematics:** discrimination normal--anomalous



# Collage from high impact journals

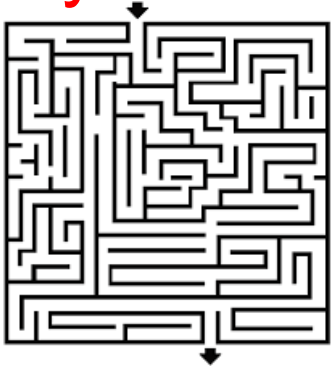


solid-state pore    zeolites (aluminum-silicate)    bio-nanopore  
 c-elegans in micro-channel



**Geometric constraint:** narrow path, compartments, branching, self-similarity (fractals), discontinuity (boundaries)

“labyrinth effect”

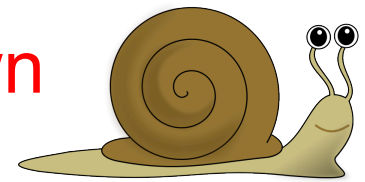


Two-state transport:

**Active** (ON) or **Stalled** (OFF)

Transport

- **hindered or slowed down**  
less: mobility, diffusivity
- **anomalous** sub-diffusion



# Violation of Central Limit Theorem

Sum of displacements

$$\langle |\mathbf{R}_N|^2 \rangle = \sum_{i=1}^N \langle |\delta\mathbf{R}_i|^2 \rangle + 2 \sum_{i>j}^N \langle \delta\mathbf{R}_i \cdot \delta\mathbf{R}_j \rangle$$

does not satisfy C.L.T

Main reasons:

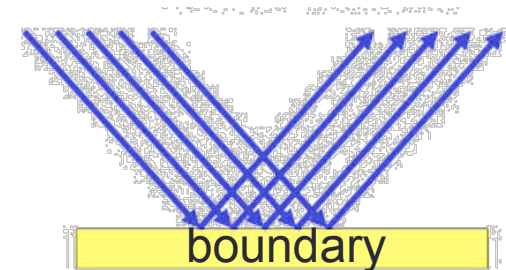
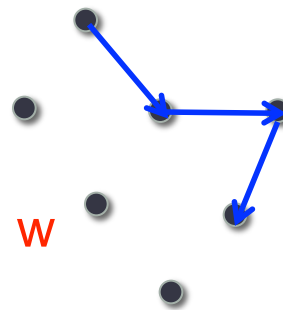
Not negligible **correlation between displacements**

**breaking of symmetry or isotropy**

(spatial correlations)

**Non-asymptotic** regimes

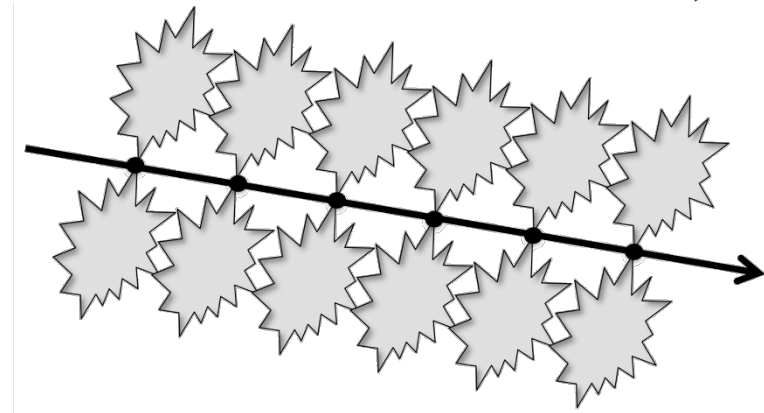
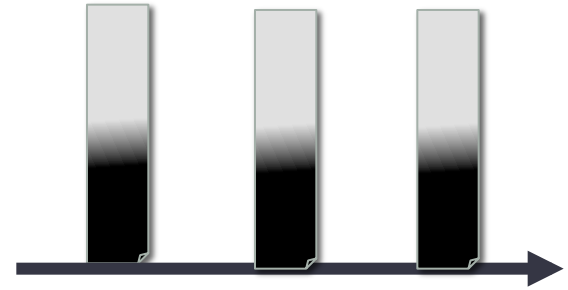
**Mean-free path**  $\sim$  confining section  $w$



Violation of CLT  $\rightarrow$  NON STANDARD BEHAVIOUR



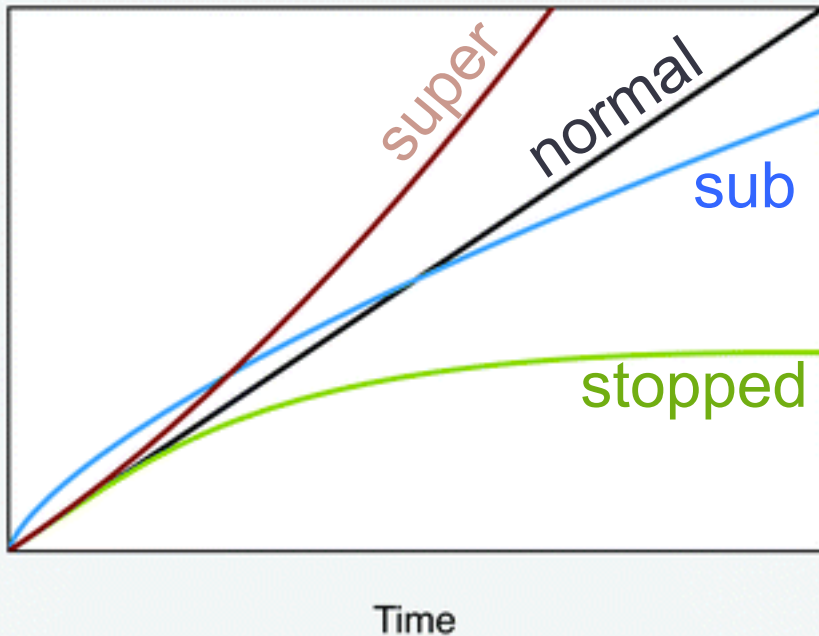
# Strong anisotropy



$$\text{M.S.D.} = \langle |\mathbf{r}(t) - \mathbf{r}(0)|^2 \rangle$$

$$MSD_{||}(t) + MSD_{\perp}(t)$$

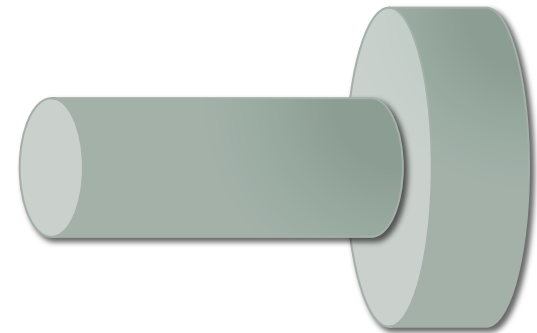
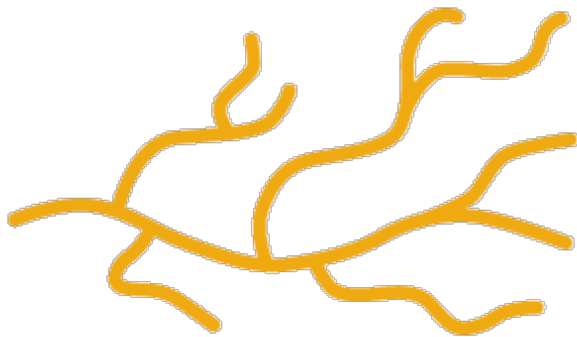
Competition: Backbone  
Sidebranches  
Turnover from:  
Anomalous => Standard





We focus on

BRANCHED STRUCTURES and CHANNELS



**Theory: main difficulty** is taking into account

- Complexity of the support of motion
- Boundary conditions

# Continuous time RW

Natural framework for trapping (waiting time)

$$\mathbf{r}(t) = \mathbf{r}_0 + \sum_{n=0}^{N(t)} \delta \mathbf{r}_n$$

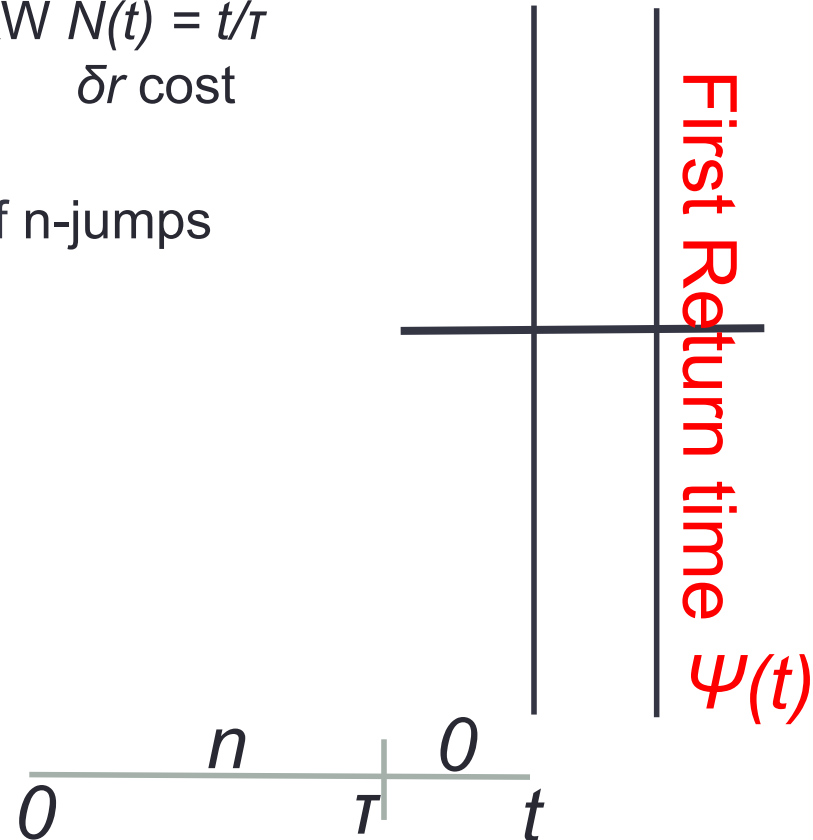
simple RW  $N(t) = t/\tau$   
 $\delta r$  cost

$$p(\mathbf{r}, t) = \sum_{n=0}^{\infty} P_n(t) p_n(\mathbf{r})$$

Pr. of n-jumps

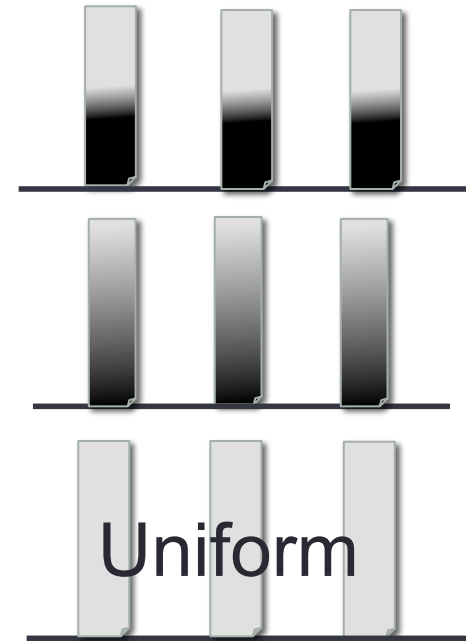
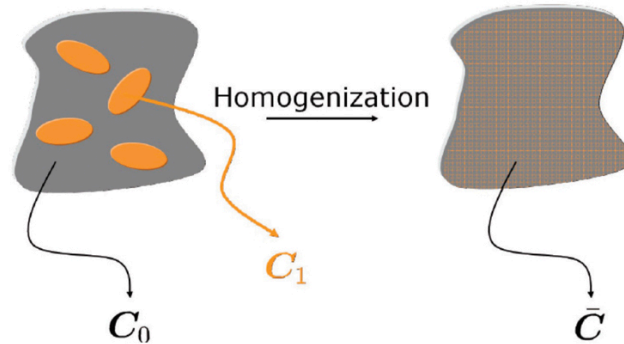
Pr. at  $\mathbf{r}$  after n-jumps

$$P_n(t) = \int_0^t \phi(t - \tau) \psi_n(\tau) d\tau$$



**Simpler method: Invoking HOMOGENIZATION !**

# Invoking Homogenization



Applied to system with strong anisotropy  
= transversal diffusion “saturated”

$$MSD_{//}(t) + MSD_{\perp}(t)$$

1) Asymptotic

Standard  $D_0 \rightarrow D_{eff}$

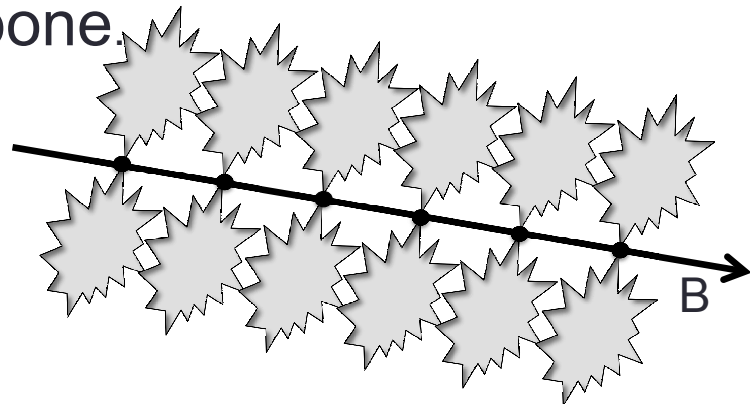
2) Pre-asymptotic

Not Standard, System dependent

# Branched structures: comb-like

Diffusion along the **backbone (B)** depends on **shape & size** of **Side-Branches**. Anomalous regimes arise from their **geometrical importance** over the backbone.

$$\langle \Delta x^2(t) \rangle = \sum_{ij} \langle \delta_i \delta_j \rangle = \sum_i \langle \delta_i^2 \rangle + 2 \sum_i \sum_{j=i+1}^t \langle \delta_i \delta_j \rangle$$



$$\langle \delta^2 \rangle = 1/2$$

**MSD**

$$\langle \Delta x^2(t) \rangle = \frac{t}{2} P_B(t) \quad D \sim \lim_{t \rightarrow \infty} P_B(t)$$

$$\delta_j = \begin{cases} 1 & \mathbf{r}_j \in B \\ 0 & \mathbf{r}_j \notin B \end{cases}$$

$$\langle \delta \rangle_\varepsilon = \varepsilon$$

**DRIFT**

$$\langle \Delta x(t) \rangle_e = \varepsilon t P_B(t)$$

$$R(t) = \frac{\langle \Delta x^2(t) \rangle_0}{\langle \Delta x(t) \rangle_\varepsilon} = \frac{1}{2\varepsilon}$$

exact simplification

**FDR**

# Effective diffusion coefficient

After homogenization

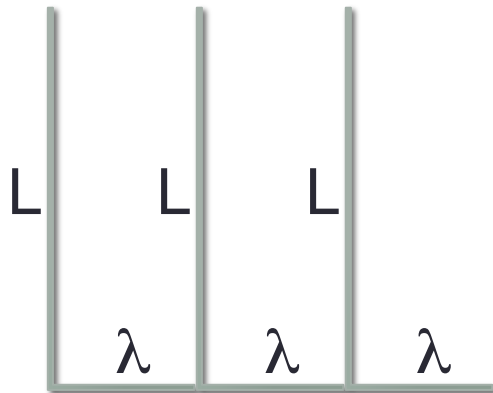
Standard diffusion on the Backbone occurs with a **renormalized coefficient**  $D < D_{free} = D_0$

$$D(L) = D_0 P_B(\infty)$$

$P_B$  asymptotic probability to occupy Backbone sites

**Homogenization:** all the  $M_{sb}$  sites on the SB are equally probable

$$p_s = 1/M_{sb}$$



$$D(L) = D_0 \frac{\lambda}{L + \lambda}$$

$$D(L) = D_0 \frac{\lambda}{L^d + \lambda} \sim L^{-d}$$

$$D(L) = D_0 \frac{M_B(L)}{M_{sb}(L) + M_B(L)}$$

# Homogenization time $t_*(L)$

$t_*(L)$  shortest timescale after which longitudinal diffusion becomes standard.

$t_*(L) \sim$  time taken by RW to visit most of  $M_{sb}(L)$  sites in a single Side-Branch of size  $L$ .

$$t_*(L) = g[M_{sb}(L)]$$

Crossover An. - Std.

$$\langle x^2(t) \rangle_0 \sim \begin{cases} t^{2\nu} & t < t_*(L) \\ D(L) t & t > t_*(L) \end{cases}$$

Matching argument at  $t_*(L)$

$$t_*(L)^{2\nu} = D(L) t_*(L)$$

Anomalous scaling

$$t_*(L) \sim L^u \quad D(L) \sim L^{-d}$$

# Distinct sites on SB visited by RW: $S(t)$

Bouchaud, Georges Phys. Rep. (1990)

$$S_{sb}(t) \sim \begin{cases} \sqrt{t} & d = 1 \\ t / \ln(t) & d = 2 \\ t & d > 2 \end{cases}$$

$$S_{sd}[t_*(L)] \sim M_{sb}(L) \sim L^d$$

$$t_*(L) \sim \begin{cases} L^2 & d = 1 \\ L^2 \ln(L) & d = 2 \\ L^d & d > 2 \end{cases}$$

$$\langle \Delta x^2(t) \rangle_{An} \sim \begin{cases} t^{1/2} & t < t_* & d = 1 \\ \ln(t) & t < t_* & d = 2 \\ c & t < t_* & d = 2 \end{cases}$$

$$S(t) \sim t^{d_s/2} \quad d_s = 2d / d_w$$

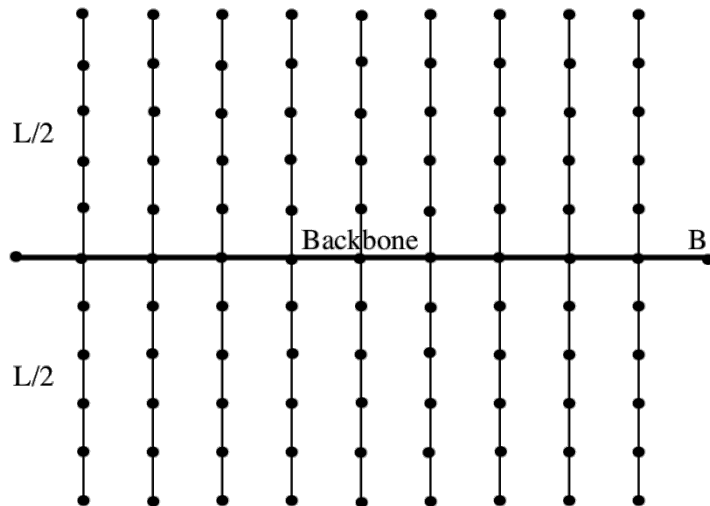
fractals

$$t_*(L) \sim L^{2d/d_s} \quad 2\nu = 1 - d_s / 2$$

$$\langle \Delta x^2(t) \rangle \sim t^{1-d_s/2}$$

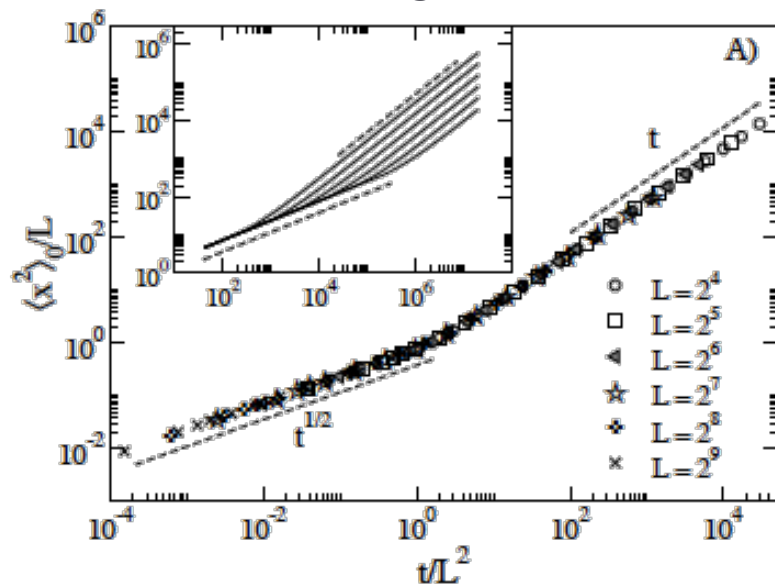


# Side-branch $d = 1$

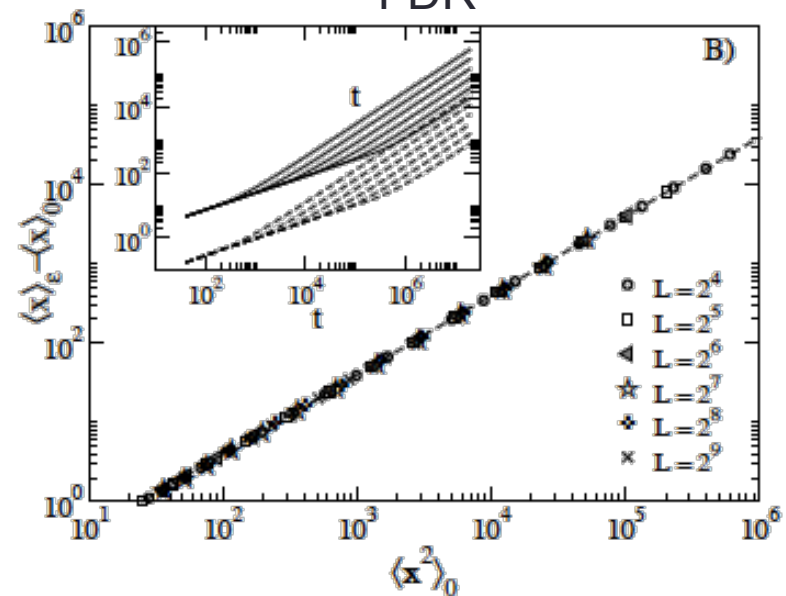


Simple comb lattice

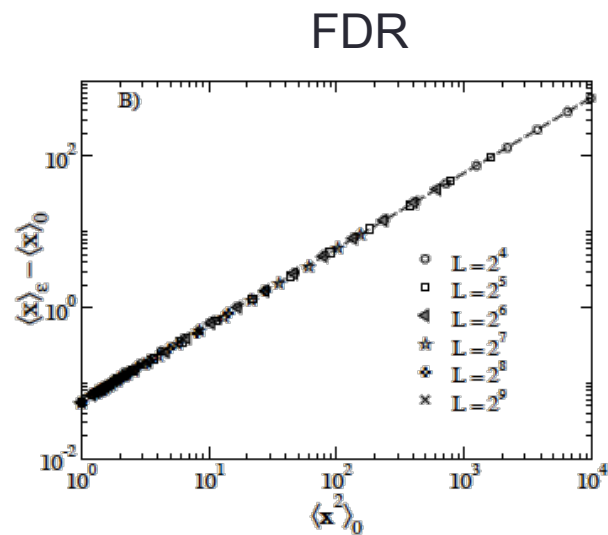
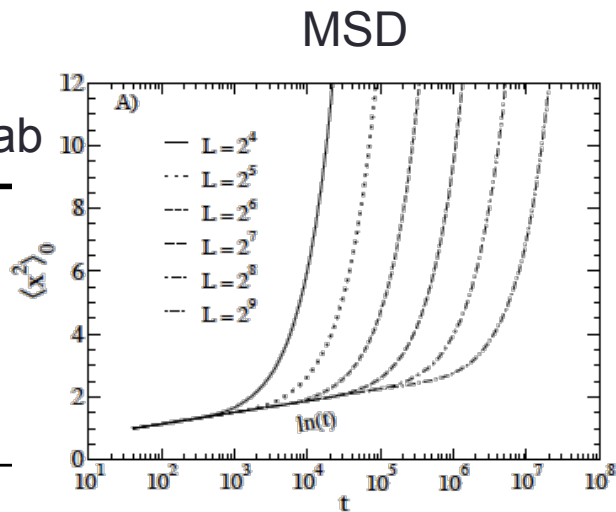
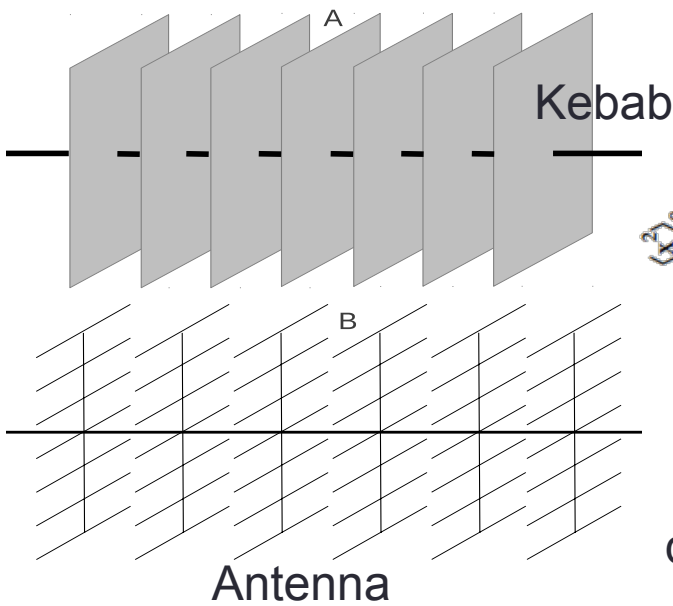
MSD



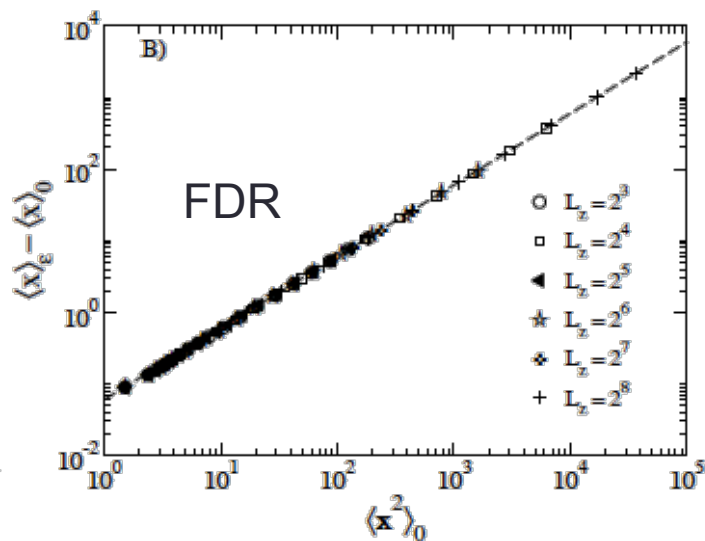
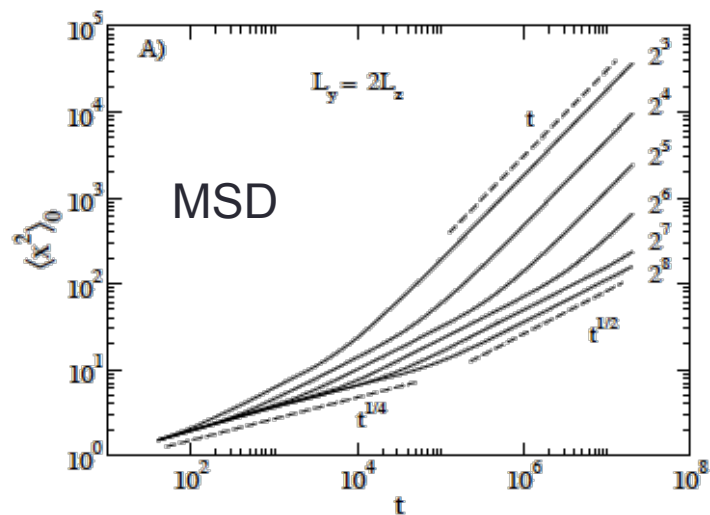
FDR



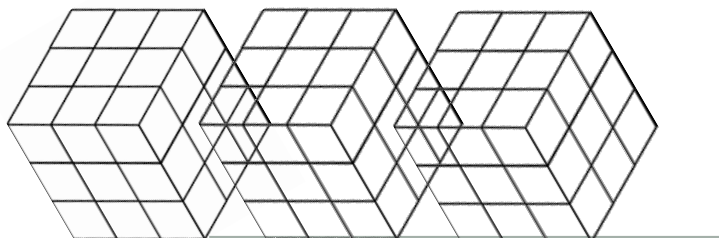
# Side-Branch $d = 2$



$d_s = 3/2$

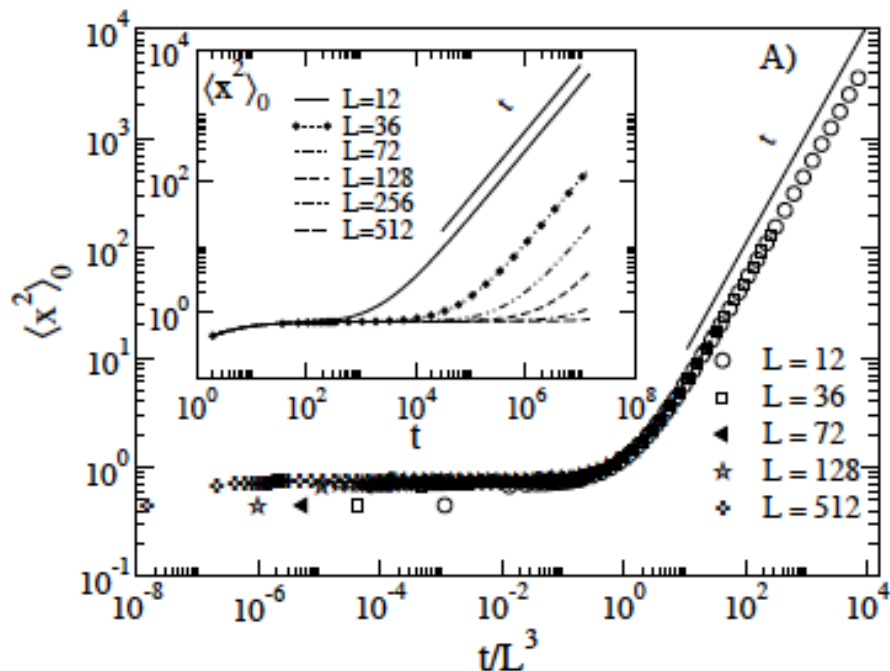


# Side-branch $d = 3$

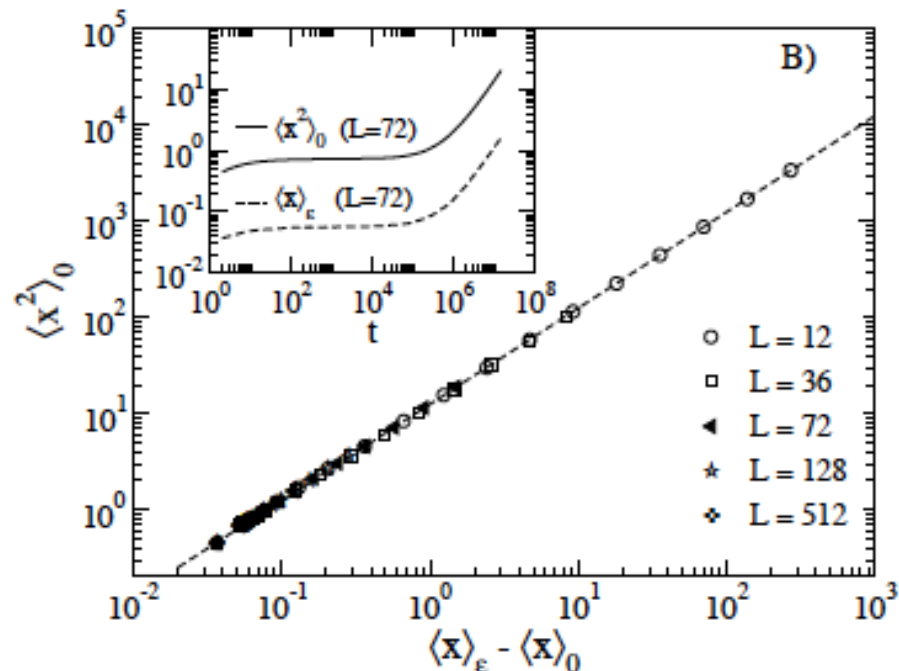


SB = CUBES along the Backbone

MSD



FDR



## More complex scenario

Discrimination standard - diffusion  
requires **spectrum of all moments**

$$\langle x^q(t) \rangle$$

a) Simple scaling (collapse)

$$P(x,t) = \frac{1}{\lambda(t)} F\left(\frac{x}{\lambda(t)}\right)$$

$$\langle x^q(t) \rangle \sim \lambda(t)^q = t^{\nu q}$$

b) Strong anomalous diffusion (Multi-scaling)

$$\langle x^q(t) \rangle \sim t^{q\nu(q)}$$

# RW on Fractal branched structures

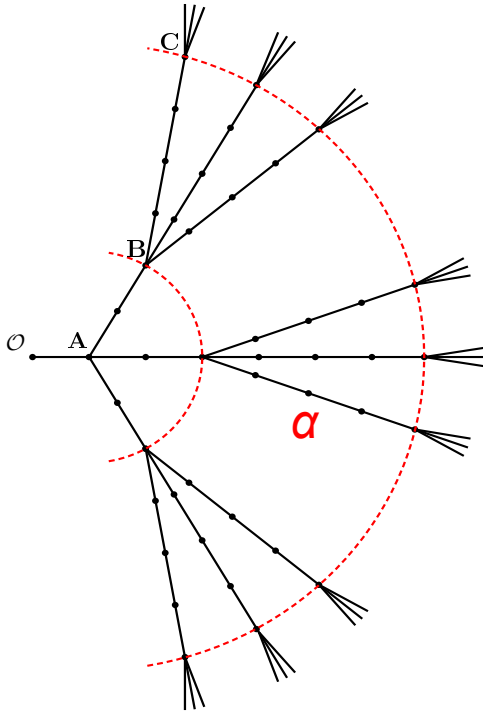
Example of a complex graph  $NT_k$   
**RW exhibits standard MSD**

$$d_f = d_s = 1 + \frac{\ln(k)}{\ln(2)}$$

$$\langle x^2(t) \rangle \sim t^{d_s/d_f} = t$$

What about the  $Q_t(\alpha, x)$  ??

Write a Master Equation for  $Q_t(\alpha, x)$   
 probability RW occupies the **Site  $x$**  in the  
**Branch  $\alpha$**        $x = \text{distance from } O$



# Distance from the origin

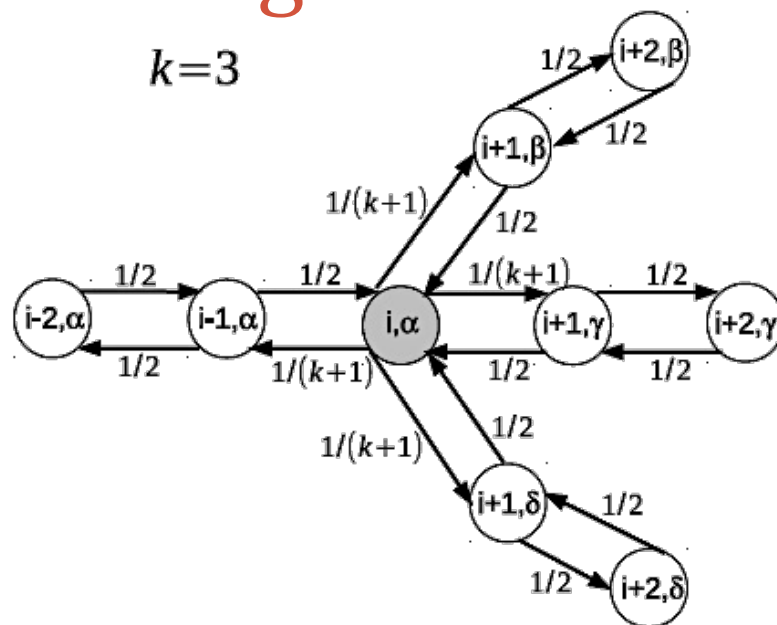
A walker on a **branching point**

$X_{t+1} = X_t + 1$  in  $k$  possibilities

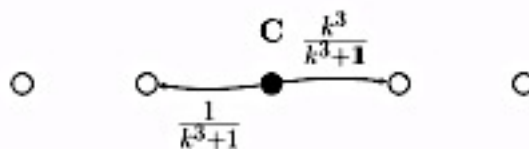
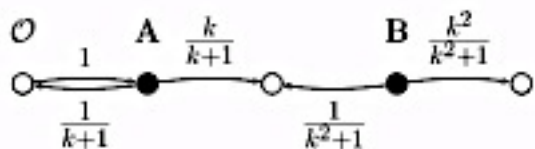
$X_{t+1} = X_t - 1$  in 1 possibility

$$w(x+1|x) = \begin{cases} \frac{k}{k+1}, & \text{if } x = 2^n - 1 \\ 1/2, & \text{elsewhere} \end{cases}$$

$$w(x-1|x) = \begin{cases} \frac{1}{k+1}, & \text{if } x = 2^n - 1 \\ 1/2, & \text{elsewhere} \end{cases}$$



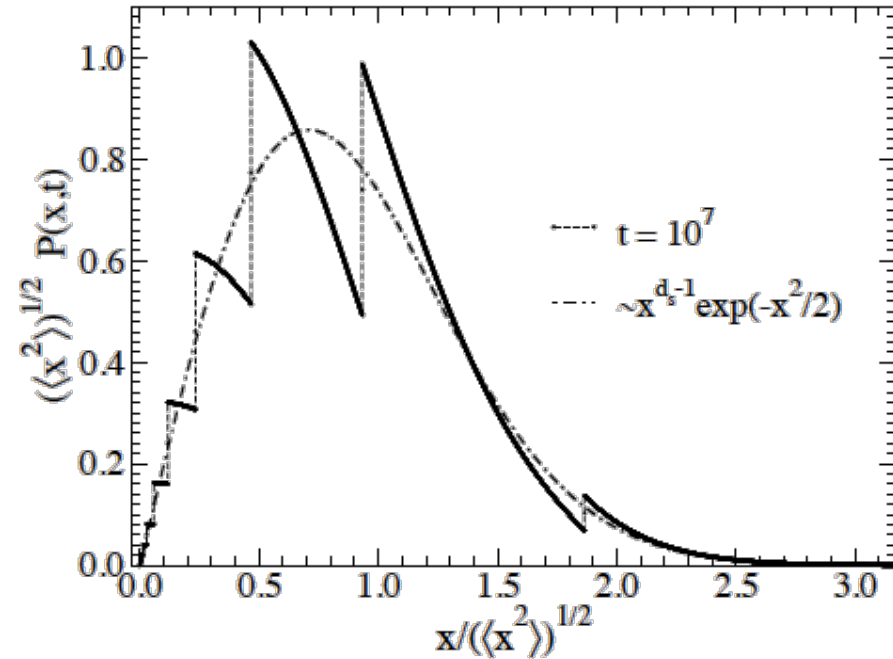
The RW on  $NT_k$  is reduced to **1D RW with DEFECTS** spaced of  $2^n$  (branch. points)



$$P_{t+1}(x-1) = \frac{1}{2}P_t(x-2) + \frac{1}{k+1}P_t(x)$$

$$P_{t+1}(x) = \frac{1}{2}P_t(x-1) + \frac{1}{2}P_t(x+1)$$

$$P_{t+1}(x+1) = \frac{k}{k+1}P_t(x) + \frac{1}{2}P_t(x+2)$$



## Approximation

$$\mathcal{F}_t(x) = \frac{2}{\Gamma(d_s/2)(2t)^{d_s/2}} x^{d_s-1} e^{-x^2/2t}$$

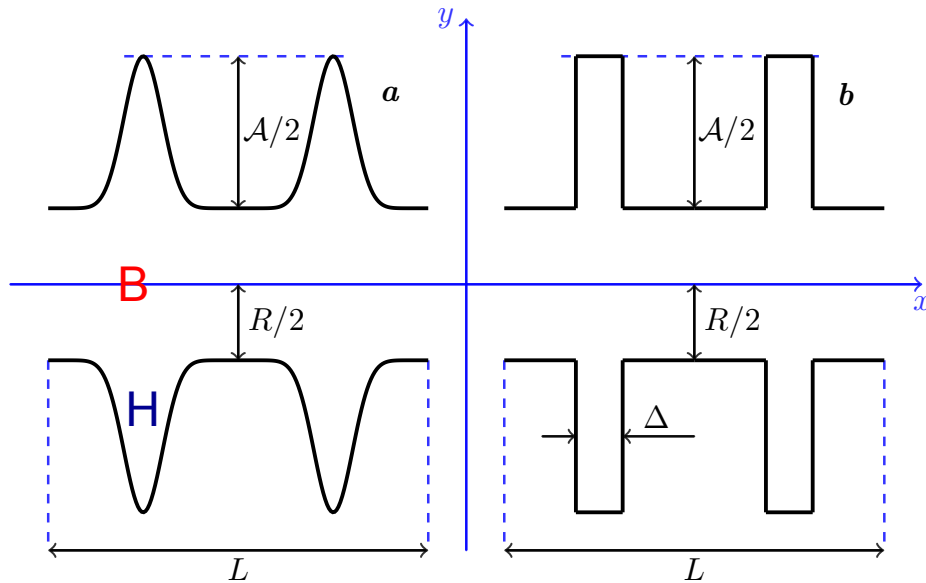
$$\langle x^q(t) \rangle = \int_0^\infty dx x^q \mathcal{F}_t(x) \sim t^{q/2}$$

“Radial” Gaussian distribution

Gaussian Scaling of moments

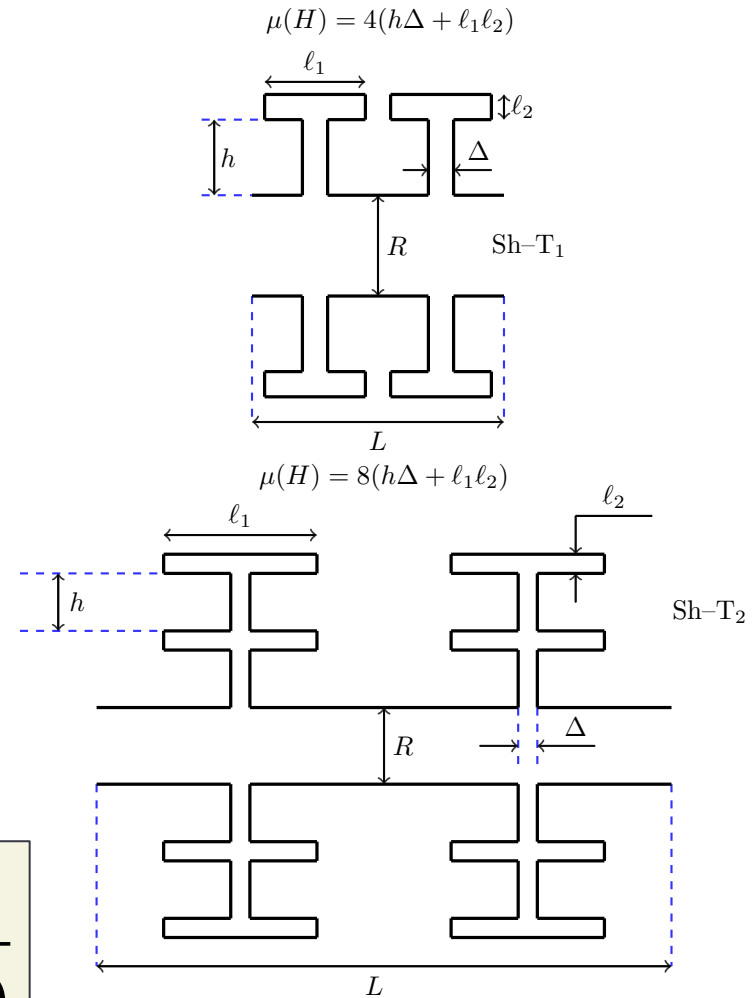


# Simpler approach: homogenization

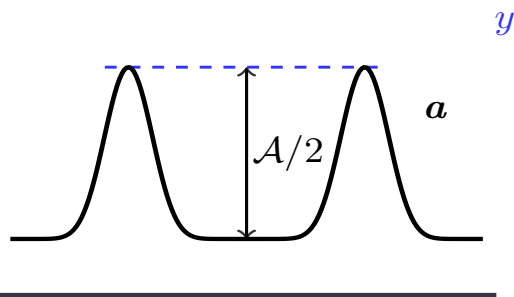


## Test of homogenization formula

$$D = D_0 P_B(\infty) = D_0 \frac{\mu(B)}{\mu(H) + \mu(B)}$$



# Periodically corrugated channels

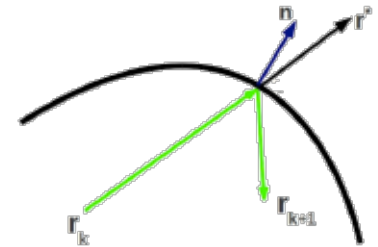


$$\frac{\partial P}{\partial t} = D_0 \left( \frac{\partial^2 P}{\partial^2 x} + \frac{\partial^2 P}{\partial^2 y} \right)$$

Fick-Jacobs  
Theory

$$Q(x, t) = \int_0^{w(x)} dy P(x, y, t)$$

No flux boundaries



Substitute in original Equation

+ factorization  $P(x, y, t) = Q(x, t) R(y, t|x)$

+ **homogenization**  $R(y, t|x) \sim 1/w(x)$

+ no-flux boundary conditions

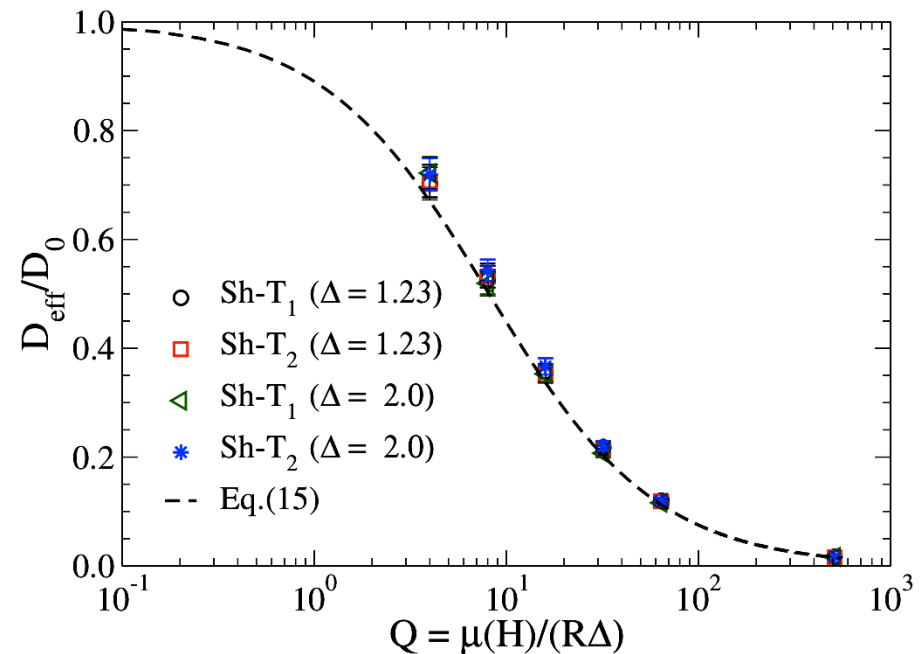
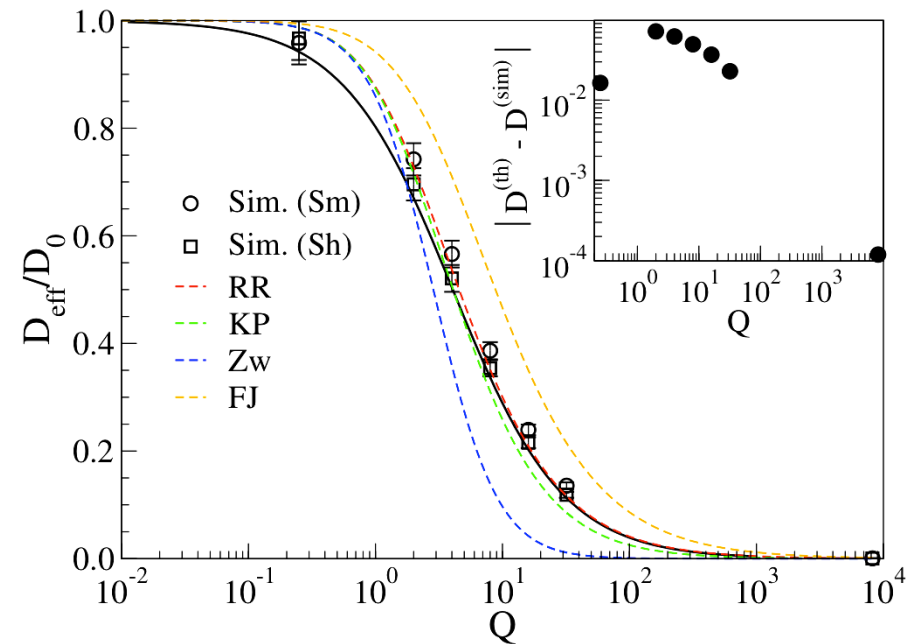
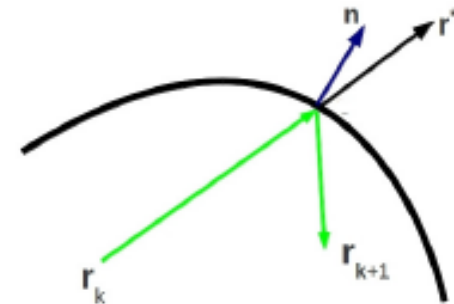
$$\frac{\partial Q}{\partial t} = D_0 \frac{\partial}{\partial x} \left( w(x) \frac{\partial Q}{\partial x} \frac{1}{w(x)} \right)$$

1-d

# Asymptotic diffusion in PCC

No flux boundaries

$$\frac{d\mathbf{r}}{dt} = -\frac{\nabla V(\mathbf{r})}{\eta} + \sqrt{2D_0}\xi_t$$



## 2 State Model H,B

$$\langle \Delta x^2(t) \rangle = 2D_0 \int_0^t du P_B(u)$$

$$\frac{dP_B(t)}{dt} = -k_B(t)P_B(t) + k_H(t) [1 - P_B(t)]$$

$$k_B(t) \sim a/\sqrt{t} \quad k_H(t) \sim b/\sqrt{t}$$

### Solution

$$P_B(t) = P_s(0) + e^{-\sqrt{t/t_0}} + \frac{D}{D_0} \left( 1 - e^{-\sqrt{t/t_0}} \right)$$

$$\langle \Delta x^2(t) \rangle = 2Dt + C \left[ 1 - e^{-\sqrt{t/t_0}} (\sqrt{t/t_0} + 1) \right]$$

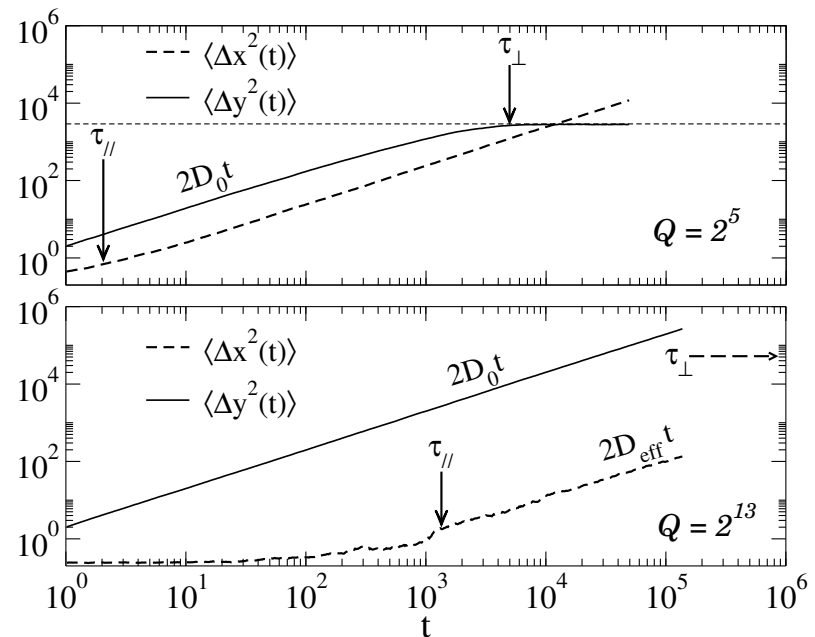
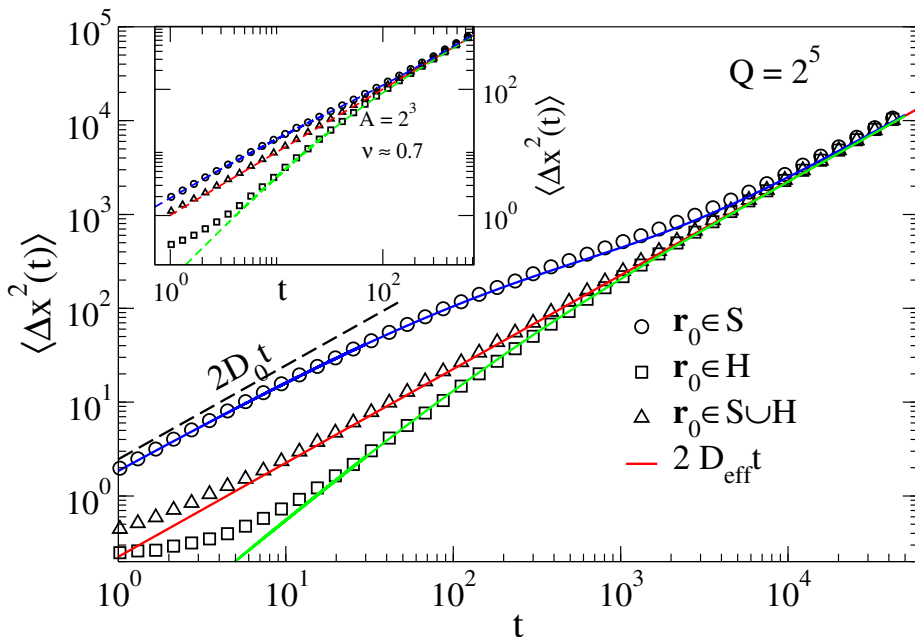
$$C = 4D_0 t_0 [P_B(0) - D/D_0]$$

Dependence on I.C.: constant  $C$

- Backbone  $P_B(0)=1$
- Humps  $P_B(0)=0$
- Uniform  $P_B(0)=D/D_0$

# Pre-asymptotic behavior

- Backbone  $P_B(0) = 1$   $C > 0$
- Humps  $P_B(0) = 0$   $C < 0$
- Uniform  $P_B(0) = D/D_0$   $C = 0$





## Conclusions

- **CONFINED RW** on branched structures and corrugated channels
- **HOMOGENIZATION** meaning and implications
- **CROSSOVER** from Anomalous to Standard
- **MATCHING ARGUMENT** → Anomalous Scaling
- **GENERALIZATION of FDR** (standard and anomalous)
- **CHECK** formula for  $D_{eff}$  and FDR in channels