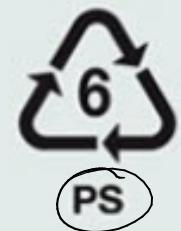


POLYMER

 PETE	 HDPE	 PVC	 LDPE	 PP	 PS	 OTHER
Polyethylene Terephthalate	High-Density Polyethylene	Polyvinyl Chloride	Low-Density Polyethylene	Polypropylene	Polystyrene	Other
Common products: soda & water bottles; cups, jars, trays, clamshells Recycled products: clothing, carpet, clamshells, soda & water bottles	Common products: milk jugs, detergent & shampoo bottles, flower pots, grocery bags Recycled products: detergent bottles, flower pots, crates, pipe, decking	Common products: cleaning supply jugs, pool liners, twine, sheeting, automotive product bottles, sheeting Recycled products: pipe, wall siding, binders, carpet backing, flooring	Common products: bread bags, paper towels & tissue overwrap, squeeze bottles, trash bags, six-pack rings Recycled products: trash bags, plastic lumber, furniture, shipping envelopes, compost bins	Common products: yogurt tubs, cups, juice bottles, straws, hangers, sand & shipping bags Recycled products: paint cans, speed bumps, auto parts, food containers, hangers, plant pots, razor handles	Common products: to-go containers & flatware, hot cups, razors, CD cases, shipping cushion, cartons, trays Recycled products: picture frames, crown molding, rulers, flower pots, hangers, toys, tape dispensers	Common types & products: polycarbonate, nylon, ABS, acrylic, PLA; bottles, safety glasses, CDs, headlight lenses Recycled products: electronic housings, auto parts,
						

Definizione: macromolecola composta da unità molecolari ("monomeri") connesse in forma di catena

Grado di polimerizzazione: $N \sim 10^2 - 10^6$ $p(N)$

Massa molecolare / totale: $M = mN$

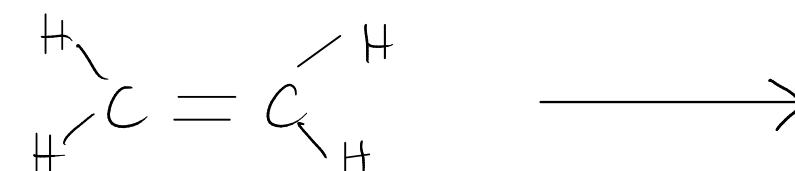
'30 : Kuhn

'71 : Flory (Nobel)

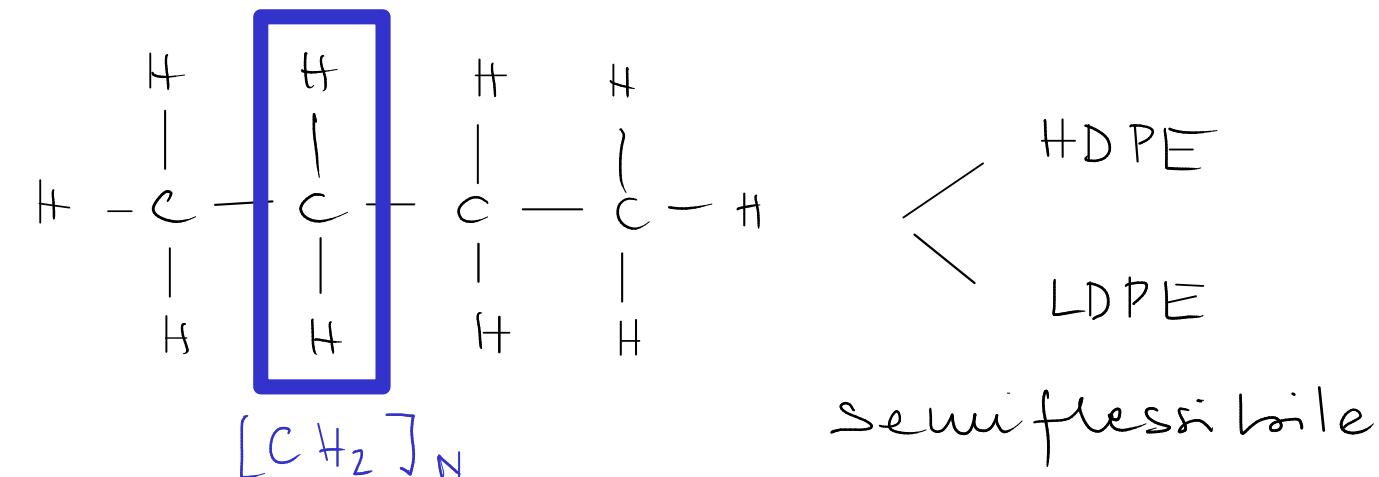
'91 : De Gennes (Nobel)

1) Polimeri sintetici

• Poli etilene (PE)

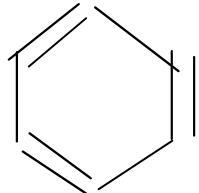


etilene C_2H_4

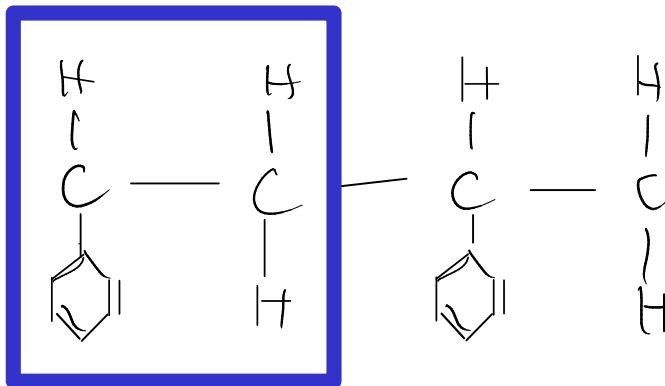


semiflessibile

• Poli(stirene) (PS)



benzeno C_6H_6



Energia di legame: $\epsilon \gg k_B T$ (@Ta)



chimica

regolare \Rightarrow XRAY

casuale \Rightarrow GLASS

flessibile

2) Biopolymen

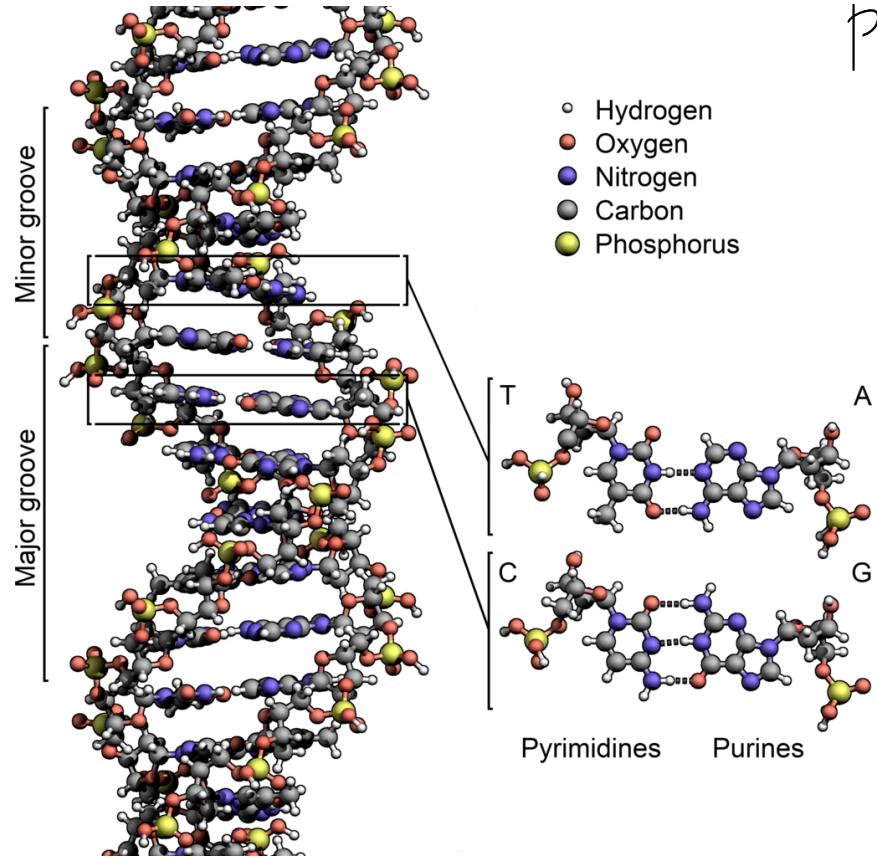
Es: proteine, polisaccandi, ...



DNA



AMIDO, CELLULOSA

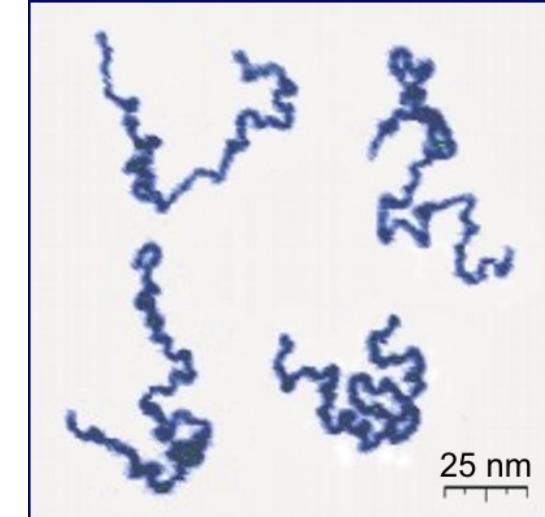
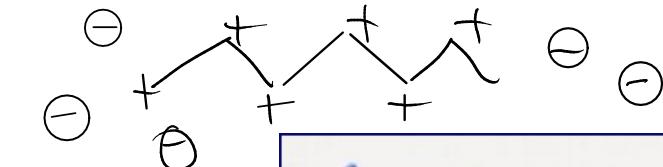


$$p \sim \delta(N - N_0)$$

$$\epsilon \sim K_B T \quad (@ T_a)$$

\downarrow
fisiici

3) Polieletroliti



Roiter, Miuko
J. Am. Chem. Soc. '05

Estensione

Lunghezza di legame : $a \sim 10^{-10} \text{ m}$

Lunghezza totale : $L = N a \sim 10^{-8} \text{ m} - 10^{-4} \text{ m}$

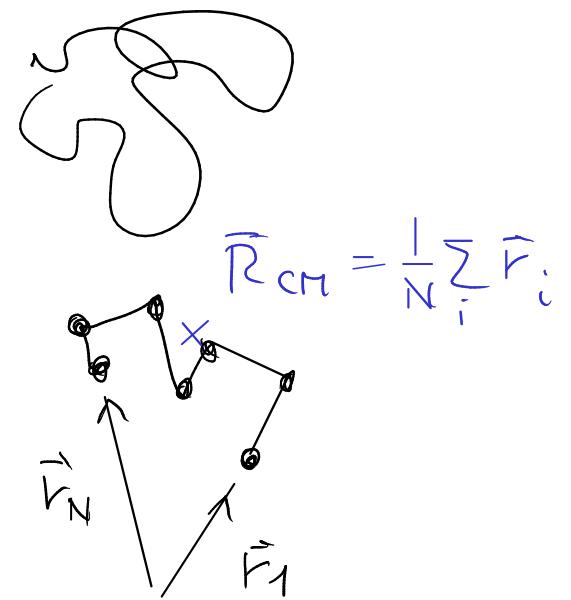
Vettore end-to-end : $\vec{R} = \vec{r}_N - \vec{r}_1$

Distanza $\| \vec{R} \|$

Conformazione : $\{ \vec{r}_1, \dots, \vec{r}_N \}$

$\langle \dots \rangle \leftarrow$ media sulle conformazioni

Raggio di girazione : $R_g = \sqrt{\langle \frac{1}{N} \sum_{i=1}^N | \vec{r}_i - \vec{r}_{CM} |^2 \rangle}$



CATENA IDEALE

24/10

- $N+1$ monomeri
- lunghezza legame $a = \text{cost}$
- orientazioni indipendenti

$$\Rightarrow \langle \bar{a}_i \cdot \bar{a}_j \rangle = a^2 \delta_{ij} = \langle \bar{a}_i \rangle \cdot \langle \bar{a}_j \rangle$$

Vettore end-to-end $\vec{R} = \vec{r}_{N+1} - \vec{r}_1$

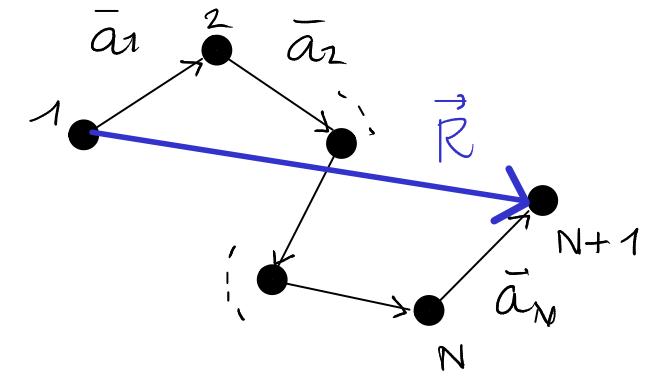
$$\langle \vec{R} \rangle = \left\langle \sum_{i=1}^N \bar{a}_i \right\rangle = \sum_{i=1}^N \langle \bar{a}_i \rangle = \vec{0}$$

Distanza end-to-end

$$\langle |\vec{R}|^2 \rangle = \left\langle \left(\sum_{i=1}^N \bar{a}_i \right) \cdot \left(\sum_{j=1}^N \bar{a}_j \right) \right\rangle = N a^2 + \sum_{i=1}^N \underbrace{\sum_{j>i}^N}_{=0} \underbrace{\langle \bar{a}_i \cdot \bar{a}_j \rangle}_{=0} = a^2 N \sim N$$

Distribuzione

$$p(\vec{R}) = p(R_x) p(R_y) p(R_z)$$



$$\bar{a}_i = \vec{r}_{i+1} - \vec{r}_i$$

Segmenti

$P(\vec{R}) \approx$ gaussiana $N \gtrsim 10$ OK

$$\langle |\vec{R}|^2 \rangle = 3 \langle R_x^2 \rangle \Rightarrow \langle R_x^2 \rangle = \frac{1}{3} a^2 N$$

$$P(\vec{R}) \approx \frac{1}{(2\pi a^2/3 N)^{3/2}} \exp\left(-\frac{3}{2a^2 N} |\vec{R}|^2\right)$$

$$|\vec{R}| \leq L = aN \quad \triangle$$

macro $\rightarrow \vec{R}$

micro $\rightarrow \{\bar{a}_1, \dots, \bar{a}_N\}$

N. conformati per \vec{R}

$$\Omega(\vec{R}) \sim \exp\left(-\frac{3|\vec{R}|^2}{2a^2 N}\right) \Rightarrow S(\vec{R}) = k_B \ln \Omega(\vec{R}) = -\frac{3k_B |\vec{R}|^2}{2a^2 N} + \text{cost}$$

$$F(\vec{R}) = -k_B T \ln \Omega(\vec{R}) = \frac{3k_B T}{2a^2 N} |\vec{R}|^2 + \text{cost} \Rightarrow \text{elasticità entropica}$$

$$= E - TS(\vec{R}) = -TS(\vec{R})$$

CATENA GAUSSIANA

- $M+1$ monomeri
- segmenti \vec{b}_i indipendenti e gaussiani

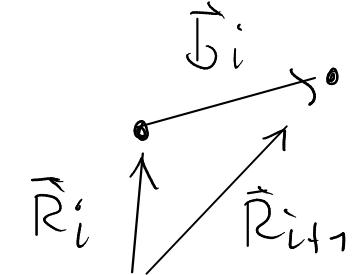
Distribuzione segmenti $\langle \vec{b}_i \rangle = \vec{0}$

$$p(\vec{b}_i) = \frac{1}{(2\pi b^2/3)^{3/2}} \exp\left(-\frac{3|\vec{b}_i|^2}{2b^2}\right)$$

Distribuzione di $\{\vec{R}_1, \dots, \vec{R}_{M+1}\}$

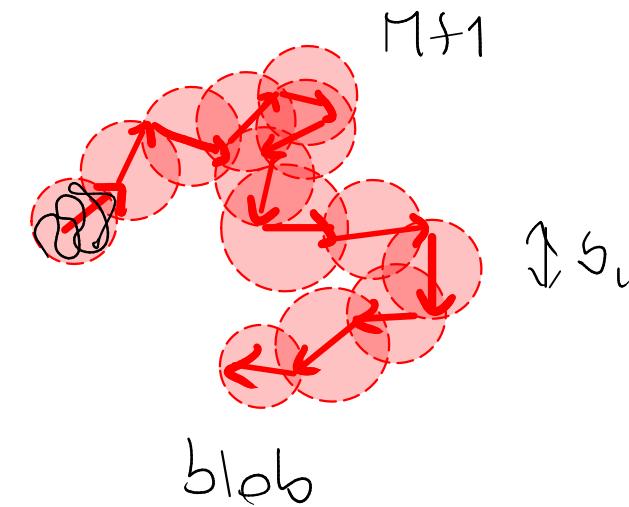
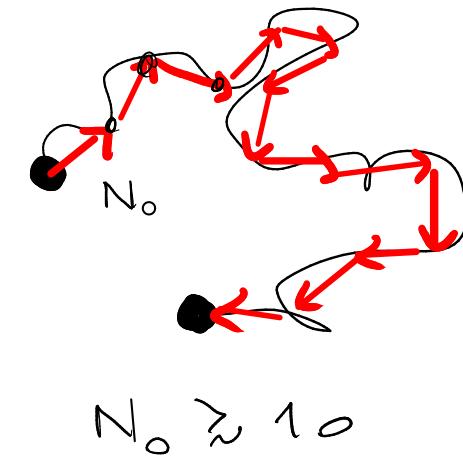
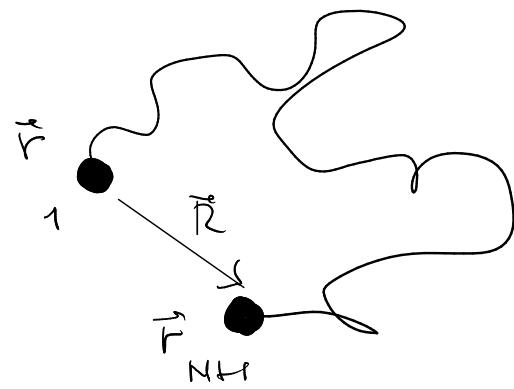
$$p(\vec{R}_1, \dots, \vec{R}_{M+1}) = \left(\frac{3}{2\pi b^2}\right)^{3N/2} \exp\left(-\sum_{i=1}^M \frac{3}{2b^2} |\vec{R}_{i+1} - \vec{R}_i|^2\right)$$

$$\Rightarrow \langle |\vec{R}|^2 \rangle = b^2 M$$



$$\vec{b}_i = \vec{R}_{i+1} - \vec{R}_i$$

Interpretazione 1 : versione "coarse-grained" della catena ideale

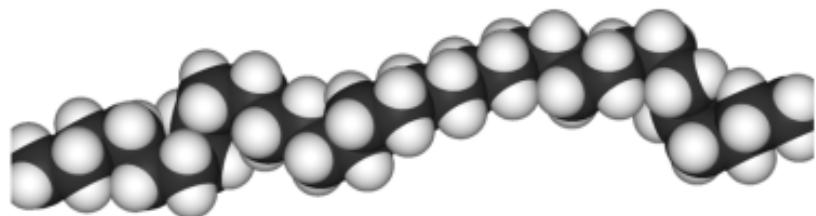


Interpretazione 2 : meccanica !

$$p(\bar{R}_1, \dots, \bar{R}_{N+1}) \sim \exp \left(- \sum_{i=1}^N \frac{3}{2b^2} |\bar{R}_{i+1} - \bar{R}_i|^2 \right) \sim \exp \left(- \frac{H(\bar{R}_1, \dots, \bar{R}_N)}{k_B T} \right)$$

$$H = \sum_{i=1}^N \frac{1}{2} \underbrace{\frac{3k_B T}{b^2}}_{\text{3/2}} |\bar{R}_{i+1} - \bar{R}_i|^2 \Rightarrow \text{catena oscillatori armonici accoppiati'}$$

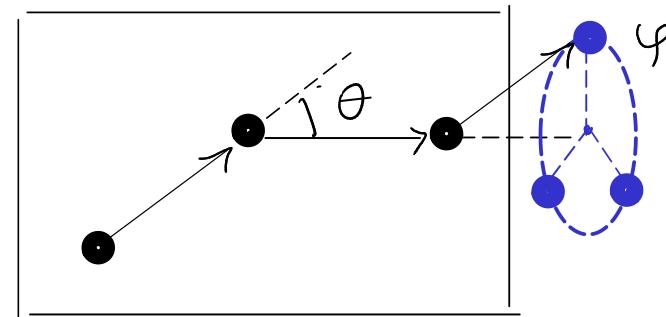
MODELLO DI KRATKY - PORE



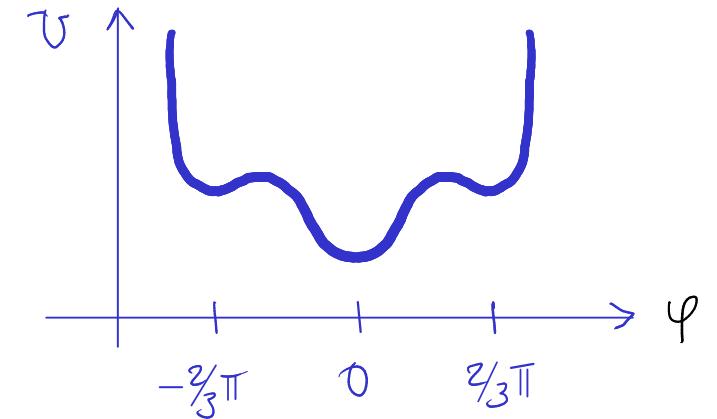
Polietilene $[CH_2]_N$

distanza di legame: $a \approx 1,5 \text{ \AA}$

angolo di legame: $\theta \approx 68^\circ$



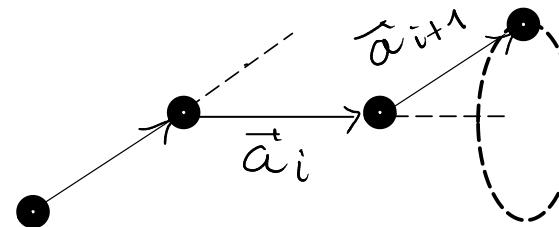
"freely-rotating chain"



- $N+1$ monomeri
- lunghezza segmenti $a = \text{cost}$
- angolo tra segmenti successivi $\theta = \text{cost}$

$$\langle \vec{a}_i \cdot \vec{a}_{i+1} \rangle \approx a^2 \cos \theta$$

$$\langle \vec{a}_i \rangle = \vec{0}$$

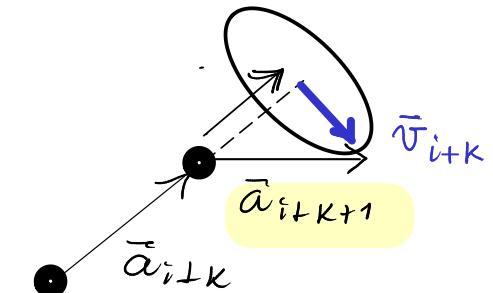


$$\langle \bar{a}_i \cdot \bar{a}_{i+k} \rangle = a^2 (\cos \theta)^k$$

dime. x induzione: $k \Rightarrow k+1$

$$\langle \bar{a}_i \rangle = \vec{0}$$

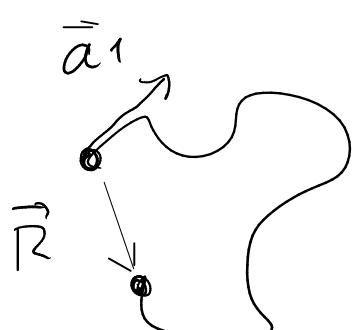
$$\begin{aligned}\langle \bar{a}_i \cdot \bar{a}_{i+k+1} \rangle &= \langle \bar{a}_i \cdot (\cos \theta \bar{a}_{i+k}) \rangle + \langle \bar{a}_i \cdot \bar{v}_{i+k} \rangle \\ &= \cos \theta \langle \bar{a}_i \cdot \bar{a}_{i+k} \rangle + \vec{0} \\ &= a^2 (\cos \theta)^{k+1} \quad \square\end{aligned}$$



Lunghezza di persistenza

$$\cos \theta < 1$$

$$(\text{es.}) \langle \bar{a}_i \cdot \bar{a}_{i+k} \rangle \sim \exp(-\frac{a}{l_p} k) \quad l_p \equiv \text{lunghezza di persistenza}$$



$$\begin{aligned}\langle \bar{R} \cdot \frac{\bar{a}_1}{a} \rangle &= \frac{1}{a} \langle \left(\sum_{i=1}^N \bar{a}_i \right) \cdot \bar{a}_1 \rangle = \frac{1}{a} \sum_{i=1}^N \langle \bar{a}_i \cdot \bar{a}_1 \rangle \\ &= a \sum_{i=1}^N (\cos \theta)^{i-1} = a \frac{1 - \cos \theta^N}{1 - \cos \theta} \xrightarrow{N \rightarrow \infty} \frac{a}{1 - \cos \theta} \\ \sum_{i=0}^{N-1} r^i &= \frac{1 - r^N}{1 - r} \sim \underbrace{\frac{a}{1 - \cos \theta}}_{= l_p}\end{aligned}$$

Distancia end-to-end

$\leftarrow N \Delta$

$$\langle |\vec{R}|^2 \rangle = a^2 N \left[\frac{1+\cos\theta}{1-\cos\theta} - \frac{2\cos\theta}{N} \frac{1-(\cos\theta)^{N+1}}{(1-\cos\theta)^2} \right] \oplus (\text{BH})$$

$$\langle |\vec{R}|^2 \rangle = \sum_{i=1}^N \sum_{j=1}^N \langle \vec{a}_i \cdot \vec{a}_j \rangle = N a^2 + \sum_{i=1}^N \sum_{j=1}^{N-1} a^2 (\cos\theta)^{|j-i|}$$

$$= N a^2 + 2 a^2 \sum_{i=1}^N \sum_{j>i}^N (\cos\theta)^{j-i} = N a^2 + 2 a^2 \sum_{i=1}^N (\cos\theta)^{-i} \sum_{j>i}^N (\cos\theta)^j$$

$$= \dots = \oplus$$

(es.)

Límite $N \rightarrow \infty$

$$\langle |\vec{R}|^2 \rangle = a^2 \frac{1+\cos\theta}{1-\cos\theta} N \approx b^2 M$$

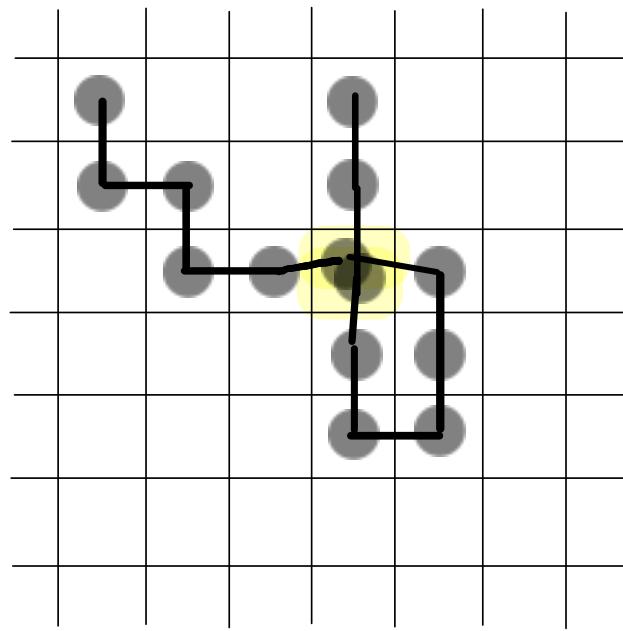
Límite $\theta \rightarrow 0$

$$\langle |\vec{R}|^2 \rangle \approx a^2 \frac{2 - \frac{\theta^2}{2}}{\theta^2/2} N \approx a^2 \frac{4}{\theta^2} N = \left(a \frac{4}{\theta^2} \right)^2 \frac{\theta^2}{4} N$$

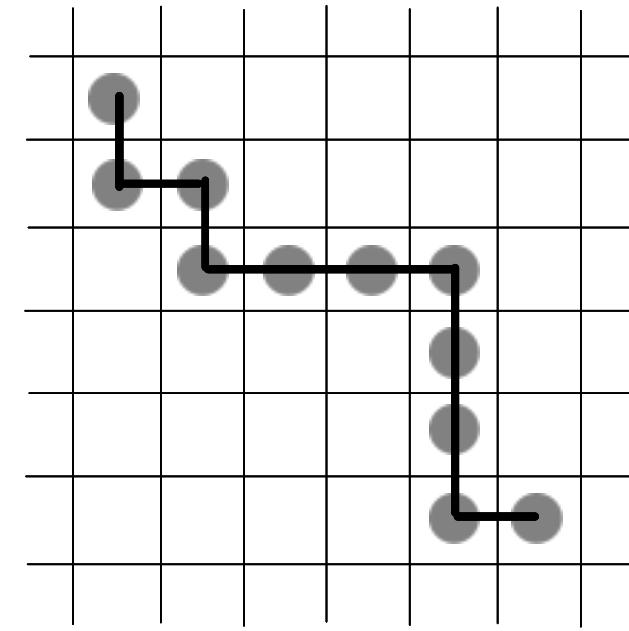
$$\underbrace{b^2}_{\sim b^2} \quad \underbrace{M}_{\sim M} \quad l_p = \frac{2}{\theta^2} = \frac{b}{2}$$

$$b = 2l_p$$

MODelli su RETI COLO



RW



SAW

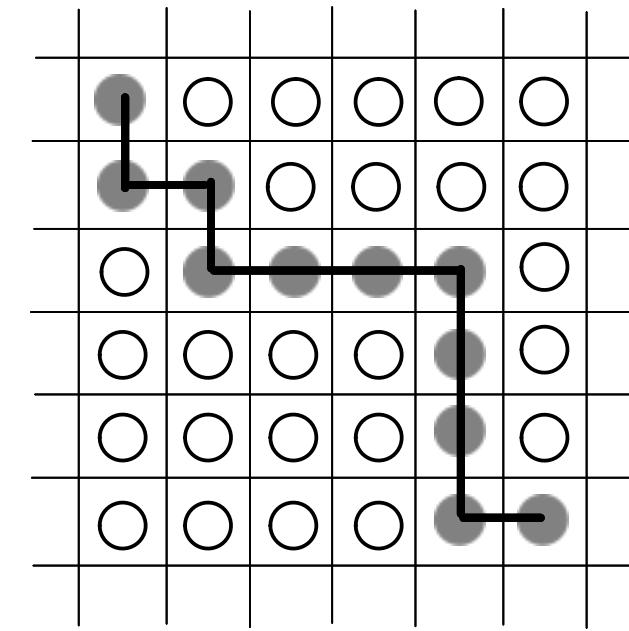
Self-avoiding
walk

$$\langle |\vec{R}|^2 \rangle \sim N$$

$$N \sim R_*^{df}$$

$$R_* \sim N^{1/2}$$

$$R_* \sim N^{3/8}$$



effetti energetici

Ruolo del solvente

$$R_* \sim N^\nu$$

$\nu = \frac{3}{5}$ BUONO
 $\nu = \frac{1}{2}$ θ
 $\nu = \frac{1}{3}$ CATTIVO

SAW

- N monomeri con $N \gg 1$
- reticolato : connettività ≥ 2 , volume cella $v = a^3$
- **limite diluito** : monomeri indipendenti

$$g = \frac{N}{R^3} \sim \frac{N}{N^{3/2}} \sim \frac{1}{N^{1/2}} \quad R \sim N^{1/2}$$

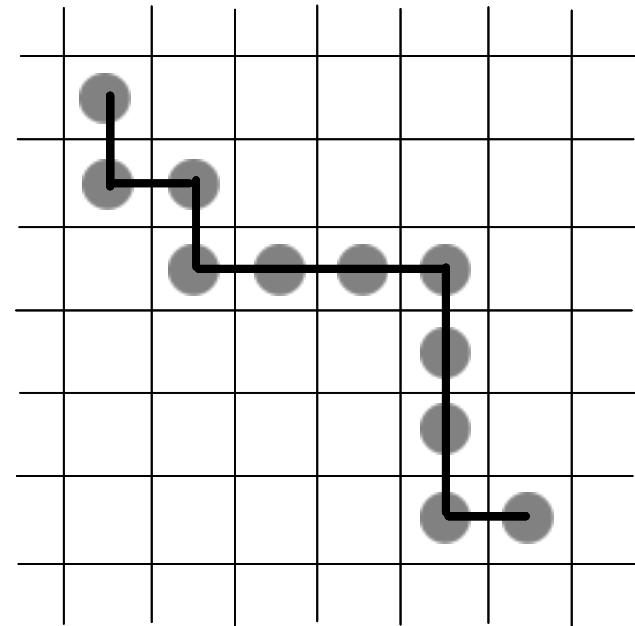
Goal: n. di conformazioni compatibili con volume escluso

Catena ideale:

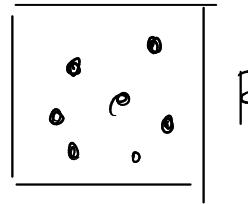
$$p_{id}(\tilde{R}) = \frac{1}{(2\pi a^2/3N)^{3/2}} \exp\left(-\frac{3}{2a^2 N} |\tilde{R}|^2\right) = \frac{1}{Z(N)} \exp\left(-\frac{3|\tilde{R}|^2}{2a^2 N}\right)$$

N. conformazioni con distanza E2E tra R e $R + dR$

$$\Omega_{id}(R) = 4\pi R^2 dR p_{id}(R) \cdot Z(N) \quad (N \gg 1)$$



volume escluso



monomeri indipendenti

$$n. \text{ celle} : \frac{R^3}{V} = \frac{R^3}{a^3}$$

$$\text{Prob. assenza overlap } N=2 : \left(1 - \frac{a^3}{R^3} \right)$$

Prob. assenza overlap per N monomeri:

$$p(R) = \left(1 - \frac{a^3}{R^3} \right)^{\frac{N(N-1)}{2}} = \exp \left[\frac{N(N-1)}{2} \log \left(1 - \frac{a^3}{R^3} \right) \right]$$

$$\text{Approx: } N \gg 1, \frac{a}{R} \ll 1$$

$$p(R) \approx \exp \left[\frac{N^2}{2} \left(-\frac{a^3}{R^3} \right) \right]$$

N. conformati compatti con volume escluso

$$\Omega(R) = \Omega_{\text{id}}(R) \cdot p(R) \sim R^2 \exp \left(-\frac{3R^2}{2a^2 N} - \frac{N^2 a^3}{2R^3} \right)$$

Distanza E2E tipica R_* = più probabile

$$F(R) = -k_B T \ln[\Omega(R)] = -T S(R)$$

$$= F_0(N) - 2k_B T \ln R + \frac{3k_B T}{2a^2 N} R^2 + \frac{k_B T N^2 a^3}{2 R^3}$$

$F_{id}(R)$ $F_{ex}(R)$

Caso ideale:

$$\frac{dF}{dR} = -\frac{2k_B T}{R} + \frac{3k_B T}{a^2 N} R \Rightarrow \frac{2k_B T}{R_*} = \frac{3k_B T R_*}{a^2 N} \Rightarrow R_*^2 = \frac{2}{3} \frac{a^2 N}{\langle \bar{R}^2 \rangle}$$

SAW: argomento Flory

$$\frac{2}{R_*} = \frac{3R_*}{a^2 N} \sim \frac{3N^2 a^3}{2R_*^4} \quad \frac{2}{3} = \frac{R_*^2}{a^2 N} - \frac{N^2 a^3}{2R_*^3} \quad O(1) \sim \frac{R_*^2}{N} - \frac{N^2}{R_*^3}$$

$$R_* \sim N^\nu \quad N^{2\nu-1} \quad N^{2-3\nu}$$

Trascuriamo $\frac{2}{3}$

$$\frac{R_*^2}{a^2 N} = \frac{N^2 a^3}{2 R_*^3} \Rightarrow R_*^5 \sim N^3 \Rightarrow R_* \sim N^{3/5}$$

$$R_* \sim N^\nu \quad \leftarrow \nu \equiv \text{esponente di Flory}$$

d dimensioni : $R_* \sim N^{\frac{3}{d+2}}$

Argomento di Flory : $\nu \sim 0.6$

Gruppo di rinormalizzazione : $\nu = 0.588$

EFFETTI ENERGETICI

- SAW
- Interazioni :
 - monomer - monomer ϵ_{mm}
 - monomer - solvente ϵ_{ms}
 - solvente - solvente $\epsilon_{ss} = 0$
- Approssimazione campo medio : trascurare fluttuazioni

Energia interazione per conformazione α

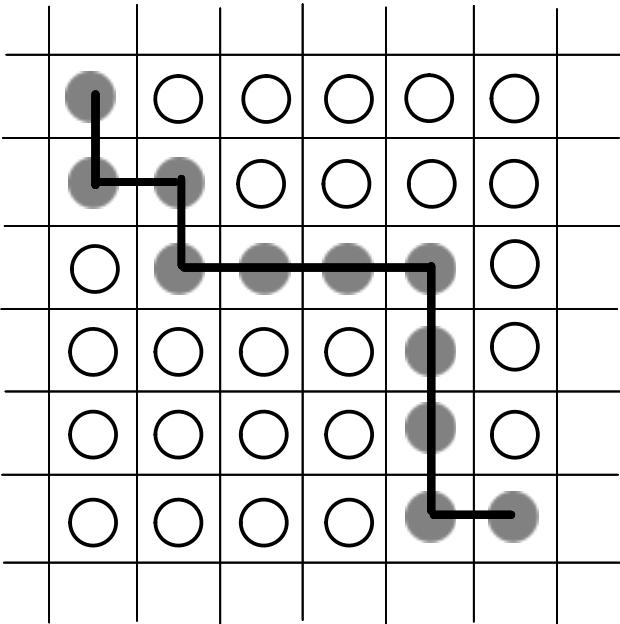
$$U_\alpha = \frac{1}{2} \sum_{i=1}^N \epsilon_{mm} n_{mm}^{(i)} + \sum_{i=1}^N \epsilon_{ms} n_{ms}^{(i)}$$

Energia media per R data

$$U(R) = \frac{1}{2} \sum_{i=1}^N \epsilon_{mm} \langle n_{mm}^{(i)} \rangle + \sum_{i=1}^N \epsilon_{ms} \langle n_{ms}^{(i)} \rangle$$

Energia libera

$$F(R) = F_{\text{SAW}}(R) + U(R) \leftarrow \text{MF}$$

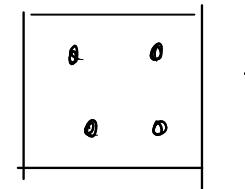


$$= F_{\text{id}}(R) + F_{\text{ex}}(R) + U(R)$$

Approx. campo medio

$$U(R) = \frac{1}{2} N \epsilon_{mm} \langle n_{mm} \rangle + N \epsilon_{ms} \langle n_{ms} \rangle$$

Bassa densità



$$\text{fractione siti occupati : } \frac{a^3 N}{R^3} = \phi$$

$$\begin{aligned} \langle n_{mm} \rangle &= z\phi \\ \langle n_{ms} \rangle &\approx z(1-\phi) \end{aligned} \quad \Rightarrow \quad U(R) = \frac{1}{2} N \epsilon_{mm} z \frac{a^3 N}{R^3} + N \epsilon_{ms} z \left(1 - \frac{a^3 N}{R^3}\right)$$

$$U(R) = \frac{1}{2} z \underbrace{(\epsilon_{mm} - 2\epsilon_{ms})}_{-\epsilon} \frac{a^3}{R^3} N^2 + \underbrace{\epsilon_{ms} z N}_{\sim \sim \text{cost}(N)}$$

$$-V(R) = -\frac{1}{2} \epsilon z \frac{a^3}{R^3} N^2 + \text{const}(N)$$

$$F(R) = F_0(N) - 2k_B T \ln R + \frac{2k_B T}{3a^2 N} R^2 + \frac{1}{2} k_B T \frac{a^3}{R^3} \left(1 - \frac{\epsilon z}{k_B T}\right) N^2$$

~~~~~  
↓  
interazioni

$$N_{\text{eff}} = a^3 \left(1 - \frac{\epsilon z}{k_B T}\right)$$

$$F(R) = F_0(N) - 2k_B T \ln R + \frac{2k_B T}{3a^2 N} R^2 + \frac{1}{2} k_B T \frac{a^3}{R^3} \left(1 - \frac{\epsilon z}{k_B T}\right) N^2$$

## Ruolo del solvente

$$v_{\text{eff}} = v \left( 1 - \frac{z\epsilon}{k_B T} \right)$$

$$\epsilon = -\underbrace{\epsilon_{mm}}_{>0} + 2\underbrace{\epsilon_{ms}}_{>0} > 0$$

1)  $k_B T \gg z\epsilon$   $v_{\text{eff}} > 0$

SAW rismalizzato

$$R_f \sim N^{3/5}$$

BUON SOLVENTE

2)  $k_B T = z\epsilon$   $v_{\text{eff}} = 0$

Catena ideale

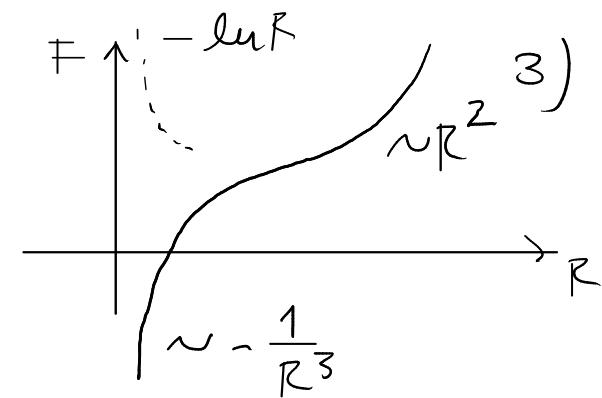
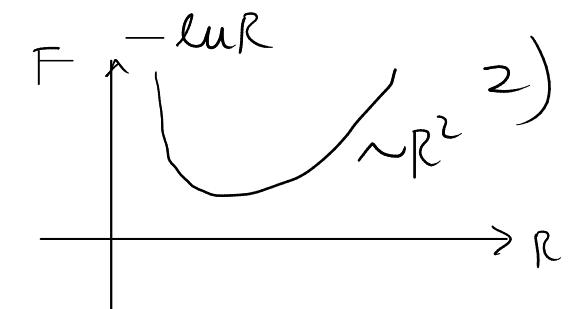
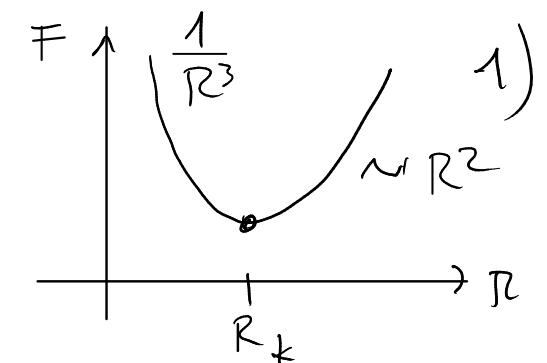
$$R_f \sim N^{1/2}$$

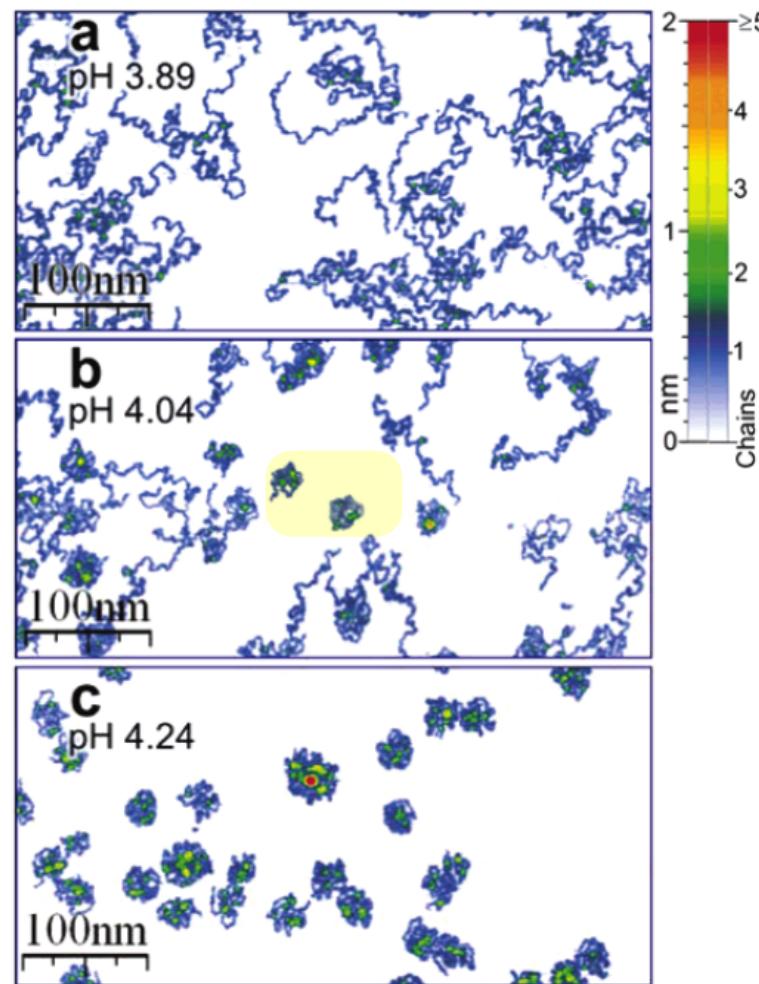
SOLVENTE  $\theta$

3)  $k_B T \ll z\epsilon$

Collasso catena

→ "globule"





COIL - GLOBULE  
transition

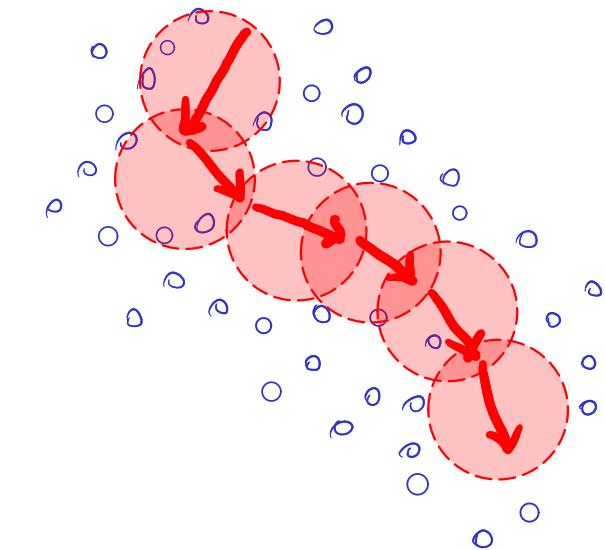
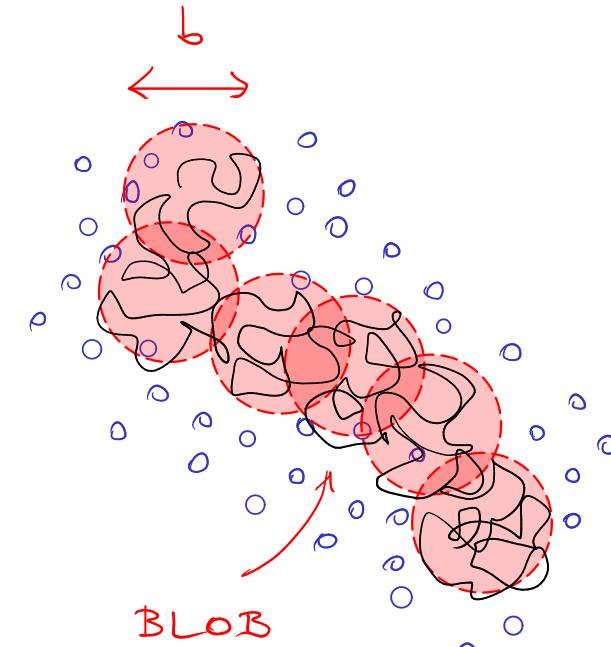
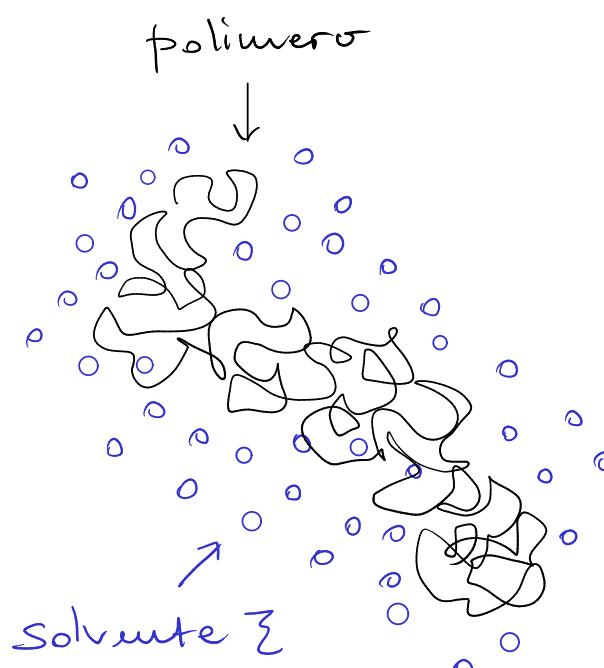
**Figure 2.** AFM-visualized conformations of adsorbed P2VP molecules:  
 (a) pH 3.89, extended coils; (b) pH 4.04, intermediate state; (c) pH 4.24,  
 compact coils. Z-scale bar shows a number of superposed chains assuming  
 the height increment of 0.4 nm.

Roiter Minko JACS 2005

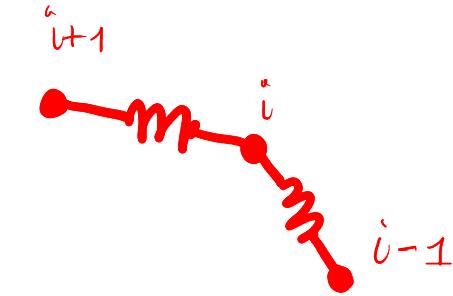
## MODELLO DI ROUSE

31/10

Dinamica di una catena gaussiana in un solvente



$$\frac{1}{2} \frac{3K_B T}{b^2} |\vec{R}_{i+1} - \vec{R}_i|^2$$



$$\sum \frac{\partial \vec{R}_i}{\partial t} = \vec{F}_i + \vec{\theta}_i(t)$$

$$\langle \vec{\theta}_i \rangle = \vec{0}$$

$$\langle \vec{\theta}_{\alpha i}(t) \vec{\theta}_{\beta j}(t') \rangle = 2 \underbrace{K_B T}_{\theta_\alpha} \sum \delta_{\alpha\beta} \delta_{ij} \underbrace{\delta(t-t')}_{\delta(t-t')}$$

- $M+1$  monomeri effettivi
- no volume escluso / attrazione
- catena gaussiana
- forze stocastiche indipendenti

$$\sum \frac{\partial \vec{R}_i}{\partial t} = \frac{3K_B T}{b^2} (\vec{R}_{i+1} - \vec{R}_i) + \frac{3K_B T}{b^2} (\vec{R}_{i-1} - \vec{R}_i) + \vec{\theta}_i(t)$$

$0 < i < M$

$$\left\{ \begin{array}{l} \bar{R}_{-1} = \bar{R}_0 \\ \bar{R}_{M+1} = \bar{R}_M \end{array} \right. \Rightarrow \quad \begin{array}{c} \text{Diagram showing a discrete sequence } \bar{R}_i(t) \text{ from } i=0 \text{ to } M. \\ \text{Curly arrows indicate shifts: } s \rightarrow s+ds \text{ and } s \rightarrow s-ds. \end{array} \quad \begin{array}{l} \bar{R}(s,t) \equiv \bar{R}_i(t) \\ \bar{R}(s+ds,t) \equiv \bar{R}_{i+1}(t) \quad \bar{R}(s-ds,t) \equiv \bar{R}_{i-1}(t) \end{array}$$

$$\sum \frac{\partial \bar{R}}{\partial t} = \frac{3k_B T}{b^2} \left[ \bar{R}(s+ds,t) + \bar{R}(s-ds,t) - 2\bar{R}(s,t) \right] + \vec{\theta}(s,t)$$

$$\sum \frac{\partial \bar{R}}{\partial t} \approx \frac{3k_B T}{b^2} \frac{\partial^2 \bar{R}}{\partial s^2} + \vec{\theta}(s,t) \quad \text{B.C.:} \quad \frac{\partial \bar{R}}{\partial s} \Big|_0 = \frac{\partial \bar{R}}{\partial s} \Big|_M = \vec{0}$$

Soluzione in spazio di Fourier:

$$\bar{R}(s,t) = \bar{x}_0(t) + 2 \sum_{p=1}^{\infty} \cos\left(\frac{p\pi}{M}s\right) \bar{x}_p(t) + \cancel{\sum_{q \neq 0} \left[ \int ds \cos\left(\frac{q\pi}{M}s\right) \dots \right]}$$

$$\bar{x}_p(t) = \frac{1}{M} \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \bar{R}(s,t)$$

$$\int_0^\pi dx \cos(px) \cos(qx) = \frac{\pi}{2} \delta_{pq} (1 + \delta_{p0})$$

$$\frac{1}{2} \left\{ \frac{\sin[(p-q)x]}{p-q} + \frac{\sin[(p+q)x]}{p+q} \right\}$$

## Equazioni del moto

$$\frac{\partial \vec{x}_p}{\partial t} = \frac{1}{M} \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \frac{\partial \vec{r}}{\partial t} = \frac{3k_B T}{3\pi b^2} \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \frac{\partial^2 \vec{r}}{\partial s^2} \quad (1)$$

$$+ \frac{1}{M^2} \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \vec{\theta}(s, t) \quad (2)$$

$$(1) \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \frac{\partial^2 \vec{r}}{\partial s^2} = \frac{p\pi}{M} \int_0^{p\pi} dt \cos t \frac{\partial^2 \vec{r}}{\partial t^2} = \frac{p\pi}{M} \left\{ \underbrace{\left[ \sin t \frac{\partial^2 \vec{r}}{\partial t^2} \right]_0^{p\pi}}_{=0} - \int_0^{p\pi} dt \sin t \frac{\partial^2 \vec{r}}{\partial t^2} \right\}$$

$$+ = \frac{p\pi}{M} s ; \frac{\partial^2 \vec{r}}{\partial s^2} = \left(\frac{p\pi}{M}\right)^2 \frac{\partial^2 \vec{r}}{\partial s^2}$$

$$= \frac{p\pi}{M} \left\{ \underbrace{\left[ \cos t \frac{\partial \vec{r}}{\partial t} \right]_0^{p\pi}}_{=0} - \int_0^{p\pi} dt \cos t \vec{r}(t) \right\} = - \left(\frac{p\pi}{M}\right)^2 \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \vec{r}(s)$$

perché  $\frac{\partial \vec{r}}{\partial t} = 0 \Leftrightarrow t = p\pi \quad (s=M)$   
 $\quad \quad \quad e \quad t=0 \quad (s=0)$

$$\rightarrow - \frac{3k_B T}{3\pi b^2} \frac{(p\pi)^2}{M} \vec{x}_p(t)$$

$$\textcircled{2} \quad \bar{\theta}_p(t) \equiv \frac{1}{M} \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \bar{\theta}(s, t)$$

$$\begin{cases} \langle \bar{\theta}_p(t) \rangle = 0 \\ \langle \theta_{\alpha p}(t) \theta_{\beta q}(t') \rangle = \frac{1}{M^2} \int_0^M ds \int_0^M ds' \cos\left(\frac{p\pi s}{M}\right) \cos\left(\frac{q\pi s'}{M}\right) \langle \theta_{\alpha p}(s, t) \theta_{\alpha q}(s', t') \rangle \\ = \frac{2k_B T \varepsilon}{M^2} \underbrace{\int_0^M ds \cos\left(\frac{p\pi s}{M}\right) \cos\left(\frac{q\pi s}{M}\right)}_{\frac{M}{\pi} \frac{\pi}{2} \delta_{qp}} \delta(t-t') \delta_{\alpha\beta} \\ = \frac{k_B T \varepsilon}{M} \delta_{qp} (1 + \delta_{p0}) \delta(t-t') \delta_{\alpha\beta} \end{cases}$$

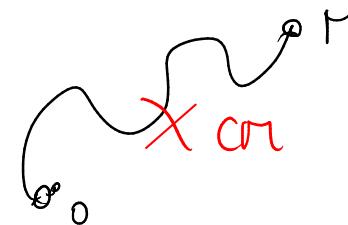
$$\frac{\partial \bar{x}_p}{\partial t} = - \frac{3k_B T}{\varepsilon \pi b^2} \frac{p^2 \pi^2}{M} \bar{x}_p(t) + \frac{1}{\varepsilon} \bar{\theta}_p(t)$$

Modi di Rouse :

$$\vec{x}_p(t) = \frac{1}{M} \int_0^M ds \cos\left(\frac{p\pi}{M}s\right) \vec{R}(s,t)$$

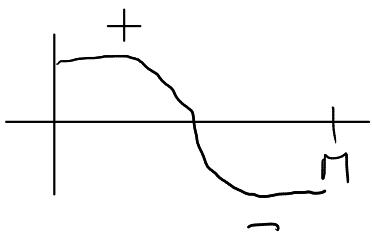
$p=0$  :

$$\vec{x}_0(t) = \frac{1}{M} \int_0^M ds \vec{R}(s,t)$$



$$\frac{1}{M} \sum_{i=0}^{M-1} \vec{R}_i(t) = c M$$

$p=1$  :



$$\frac{\partial \vec{x}_p}{\partial t} = - \frac{3 k_B T}{\varepsilon \pi b^2} \frac{p^2 \pi^2}{M} \vec{x}_p(t) + \frac{1}{\varepsilon} \vec{\theta}_p(t) = - \frac{1}{\tau_p} \vec{x}_p(t) + \frac{1}{\varepsilon} \vec{\theta}_p(t)$$

$$\langle \theta_{\alpha p}(t) \theta_{\beta q}(t') \rangle = \frac{k_B T \varepsilon}{M} \delta_{qp} (1 + \delta_{pq}) \delta(t-t') \delta_{\alpha p}$$

Noto del CM:  $\dot{P} = 0$

$$\frac{\partial \vec{x}_p}{\partial t} = \frac{1}{\zeta} \bar{\theta}_o(t) \Rightarrow \vec{x}_o(t) = \bar{x}_o(0) + \frac{1}{\zeta} \int_0^t dt' \bar{\theta}_o(t')$$

$$\langle |\vec{x}_o(t) - \bar{x}_o(0)|^2 \rangle = \frac{1}{\zeta^2} \int_0^t dt' \int_0^t dt'' \langle \bar{\theta}_o(t') \cdot \bar{\theta}_o(t'') \rangle$$

$$= 6 \frac{1}{\zeta^2} \frac{K_B T \zeta}{M} t = 6 \frac{K_B T}{\zeta M} t \sim t \quad D_{CM} \sim \frac{1}{M}$$


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Dinamica dei modi di Rouse ( $p \geq 1$ )

$$\langle \vec{x}_p(t) \cdot \vec{x}_p(0) \rangle = \dots = \langle |\vec{x}_p(0)|^2 \rangle e^{-t/\tau_p}$$

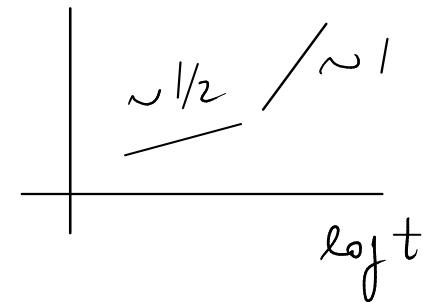
$$\tau_p \equiv \frac{\zeta M^2 b^2}{3 K_B T p^2 \pi^2} \quad \text{tempo di correlazione di Rouse}$$

$$\tau_p \sim M^2 \sim \frac{1}{p^2}$$

Dinamica segmentale  $\rightarrow \bar{R}(s,t) \rightarrow \bar{R}_i(t)$   $\log y < \Delta r^2$

$$t \gg \tau_1 \quad \langle |\bar{R}_i(t) - \bar{R}_i(0)|^2 \rangle \sim t \quad (\text{es.})$$

$$t \ll \tau_1 \quad \langle |\bar{R}_i(t) - \bar{R}_i(0)|^2 \rangle \sim t^{1/2} \quad \text{sotto-diffusione}$$



Dinamica rotazionale

$$\vec{R}(t) = \vec{R}(M,t) - \vec{R}(0,t) \quad \text{vettore end-to-end}$$

$$\langle \vec{R}(t) \cdot \vec{R}(0) \rangle = 16 \sum_{p=1,3,\dots}^{\infty} \langle \vec{x}_p(t) \cdot \vec{x}_p(0) \rangle \quad \tau_R \sim \tau_1 \sim M^2$$

$$\begin{aligned} \vec{R}(t) &= \vec{x}_0 + 2 \sum_{p=1}^{\infty} \cos(p\pi) \vec{x}_p(t) - \vec{x}_0 - 2 \sum_{p=1}^{\infty} \vec{x}_p(t) \\ &= 2 \sum_{p=1}^{\infty} [\cos(p\pi) - 1] \vec{x}_p(t) = -4 \sum_{p=1,3,5,\dots}^{\infty} \vec{x}_p(t) \end{aligned}$$

- 2 disp-  
 0 pari

## Modello Rouse

$$\left\{ \begin{array}{l} D_{CM} \sim \frac{1}{M} \\ \tau_R \sim M^2 \end{array} \right.$$

## Esperimenti

$$\left\{ \begin{array}{l} D_{CM} \sim \frac{1}{M^\nu} \\ \tau_R \sim M^{2\nu+1} \end{array} \right.$$

TABLE 2: Experimental Values of  $\alpha$  and  $d_F = 1/\alpha$  As Found in the Literature or in This Study

| molecule family                                   | $\alpha$ | $d_F$ | range                    | source                     |
|---------------------------------------------------|----------|-------|--------------------------|----------------------------|
| globular proteins                                 | 0.39     | 2.56  | 2.04                     | PDB <sup>13</sup>          |
| globular proteins                                 | 0.39     | 2.56  | 1.46                     | this work                  |
| PS in toluene                                     | 0.41     | 2.45  | 2.93                     | this work                  |
| PMMA in acetone below 30 kD                       | 0.46     | 2.17  | 1.68                     | this work                  |
| PS in acetone                                     | 0.47     | 2.15  | 1.72                     | this work                  |
| PS in CDCl <sub>3</sub> below 20 kD               | 0.47     | 2.12  | 1.62                     | this work                  |
| PMMA in CDCl <sub>3</sub> below 30 kD             | 0.48     | 2.07  | 1.68                     | this work                  |
| oligosaccharides <sup>a</sup>                     | 0.48     | 2.07  | 2.17 (3.40) <sup>b</sup> | NMR <sup>3</sup>           |
| PS in THF below 20 kD                             | 0.50     | 2.01  | 1.72                     | this work                  |
| PEO in D <sub>2</sub> O                           | 0.54     | 1.86  | 3.90                     | this work                  |
| small molecules in D <sub>2</sub> O <sup>a</sup>  | 0.54     | 1.84  | 1.39                     | NMR <sup>5</sup>           |
| PMMA in acetone above 25 kD                       | 0.54     | 1.84  | 1.81                     | this work                  |
| PEO in water                                      | 0.55     | 1.82  | 2.80                     | NMR <sup>6</sup>           |
| small molecules in CDCl <sub>3</sub> <sup>a</sup> | 0.56     | 1.77  | 1.60                     | NMR <sup>5</sup>           |
| DNA                                               | 0.57     | 1.75  | 1.69                     | fluorescence <sup>12</sup> |
| PEO in CDCl <sub>3</sub>                          | 0.58     | 1.73  | 4.13                     | this work                  |
| denatured peptide <sup>a,c</sup>                  | 0.58     | 1.71  | 1.15                     | NMR <sup>2</sup>           |
| PMMA in CDCl <sub>3</sub> above 25 kD             | 0.61     | 1.65  | 1.81                     | this work                  |
| PS in CDCl <sub>3</sub> above 20 kD               | 0.61     | 1.63  | 2.63                     | this work                  |
| PS in THF above 20 kD                             | 0.62     | 1.61  | 2.00                     | this work                  |
| Linear alkanes                                    | 0.71     | 1.41  | 0.50 (C8-C26)            | this work                  |

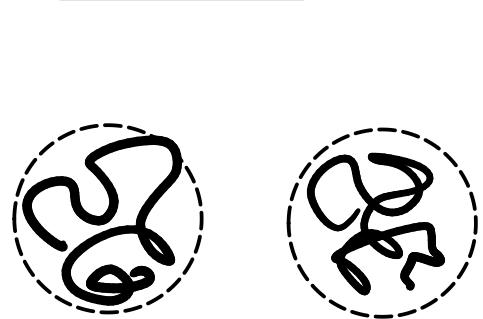
## Modelli di Zimm

→ MF interazioni idrodinamiche

$$D_{CM} \sim \frac{1}{M^\alpha} \quad \sim \frac{1}{M^{1/2}}$$

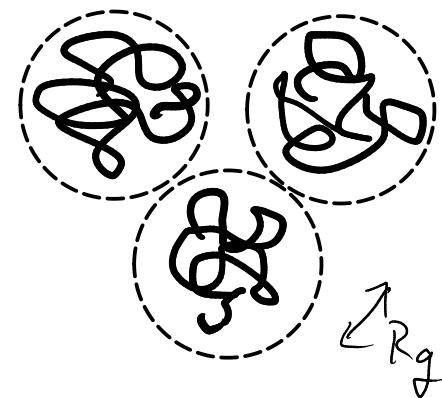
## REGIMI DI DENSITÀ

Diluito



$$\bar{g} \approx \bar{g}^*$$

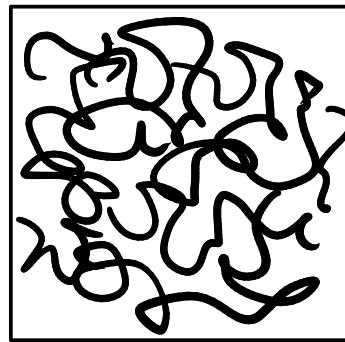
Semi-diluito



$$\bar{g} \approx \bar{g}^*$$

$$\bar{g} \gtrsim \bar{g}^*$$

Concentrato



$$\bar{g} \gg \bar{g}^*$$

MELT

De Gennes

"Reptation"

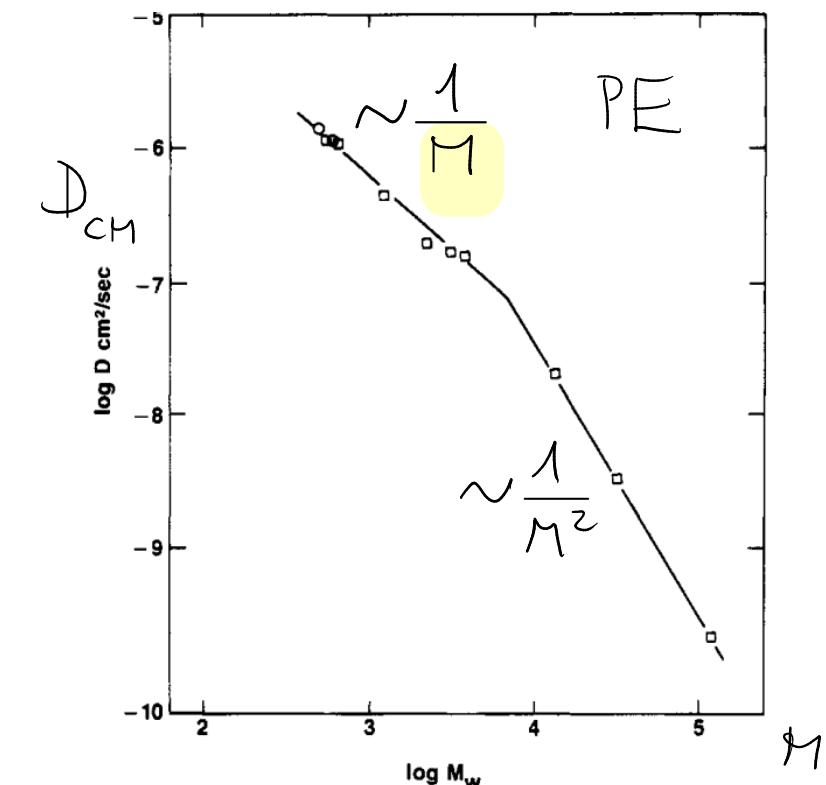


Figure 11. Self-diffusion coefficient corrected to the temperature at which the friction factor for viscosity equals  $2.3 \times 10^{-9}$  dyn·s/cm. The temperatures are listed in Table III. Symbols are same as Figure

Pearson et al.  
Macromolecules