

Polymer Simulations

Report

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1 The random walk model:Gaussian Chain

Consider a linear polymer to be a freely-jointed chain with N beads, length of each bond is b , that occupy zero volume. The path of the chains is like a 'random walk' in three dimensions, limited only by the constraint that each segment must be joined to its neighbors.

Consider the 'end to end' vector \mathbf{R} joining one end of the polymer to the other, the average value $\langle \mathbf{R} \rangle$ is zero, since the probability of \mathbf{R} equals $-\mathbf{R}$. Therefore we will calculate $\langle \mathbf{R}^2 \rangle$

$$\langle \mathbf{R}^2 \rangle = \sum_{n=1}^N \sum_{m=1}^N \langle r_n \cdot r_m \rangle \quad (1)$$

We consider that there is no correlation between bead n and m , therefore we find :

$$\langle \mathbf{R}^2 \rangle = \sum_{n=1}^N \langle r_n^2 \rangle = Nb^2 \quad (2)$$

The probability distribution of \mathbf{R} is:

$$P(\mathbf{R}, N) = \left(\frac{3}{2\pi Nb^2}\right)^{3/2} \exp\left(-\frac{3\mathbf{R}^2}{2Nb^2}\right) \quad (3)$$

The probability distribution is Gaussian, so we also call it Gaussian Chain.

1.1 Random walk Simulation

```
%This program is used to simulate the brownian motion
classdef FirstTry<handle

properties
    dimension%dimension=1,2 or 3
    numParticles%number of particles in polymers;
    dt%pas de temps
    numSteps% number of motion
    diffusionConst %constante diffusion
    paths %the paths of polymer;
end

methods

%class constructor
function obj=FirstTry(dimension,numParticles,dt,diffusionConst,numSteps)
    obj.dimension=dimension;
    obj.numParticles=numParticles;
    obj.dt=dt;
    obj.numSteps=numSteps;
    obj.diffusionConst=diffusionConst;
    obj.paths = zeros(obj.numParticles,3,obj.numSteps);
```

```

end

function Calculate(obj)

for j=1
noise = [zeros(1,obj.dimension);...
sqrt(2*obj.diffusionConst*obj.dt)*randn(obj.numParticles-1,obj.dimension)];
obj.paths(:,1:obj.dimension,j)=cumsum(noise);

end

for j=2:obj.numSteps
noise = [sqrt(2*obj.diffusionConst*obj.dt)*randn(obj.numParticles,obj.dimension)];
obj.paths(:,1:obj.dimension,j)=obj.paths(:,1:obj.dimension,j-1)+noise;

end

end %'random walk 'simulation

function Plot(obj)
f=figure;
b =[-20 20];
a= axes('Parent',f,'NextPlot','replaceChildren','XLim',b,'YLim',b,'ZLim',b);
c=rand(1,3);

x = obj.paths(:,1,1);
y = obj.paths(:,2,1);
z = obj.paths(:,3,1);
%      end
l=line('XData',x,'YData',y,'ZData',z,'Color',c,'linestyle','-','Marker','o','markersize',10,'Parent',f);
for i=2:obj.numSteps

set(l,'XData',obj.paths(:,1,i),'YData',obj.paths(:,2,i),'ZData',obj.paths(:,3,i));

pause(1)

drawnow

end
end

end
end

```

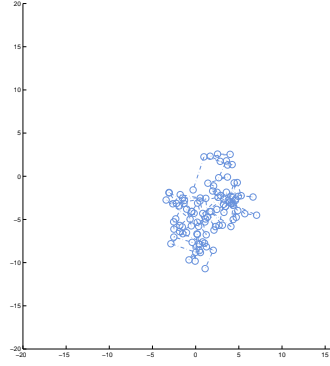


Figure 1: initial position

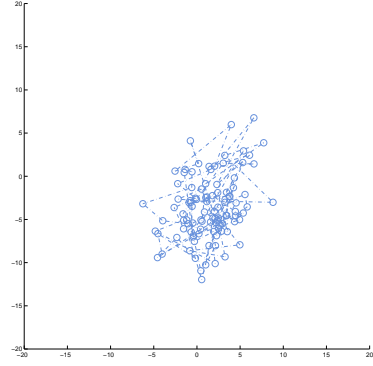


Figure 2: final position

1.2 Probability distribution function of R

In order to verify that the PDF of R is Gaussian, we calculate the 'end to end distance' R for each simulation, then we plot it with histogram by coordinate (x, y, z) respectively.

