

Diffusion in Biological Systems

class - 14 (16.10.24)

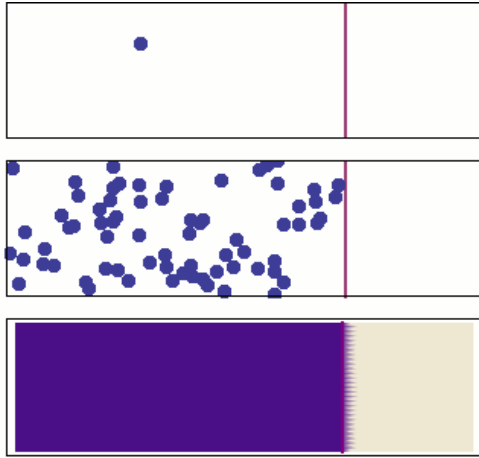
LS2103 (Autumn 2024)

Dr. Neelanjana Sengupta

Associate Professor, DBS

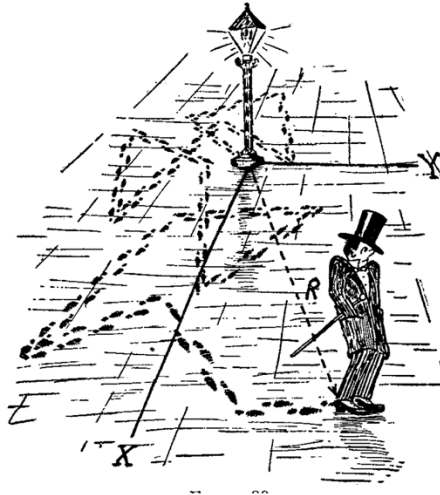
<https://www.iiserkol.ac.in/~n.sengupta/>

In General: Random Biological Processes



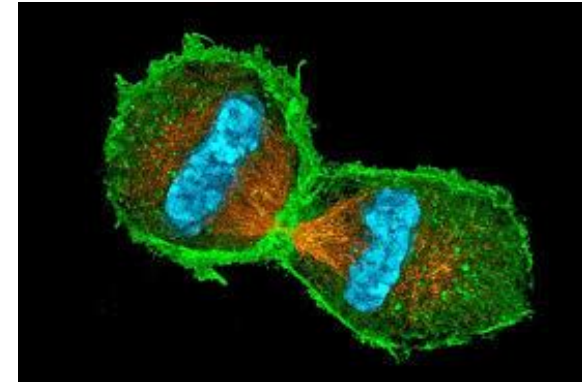
Diffusion

<https://commons.wikimedia.org/w/index.php?curid=8995324>



Random Walk

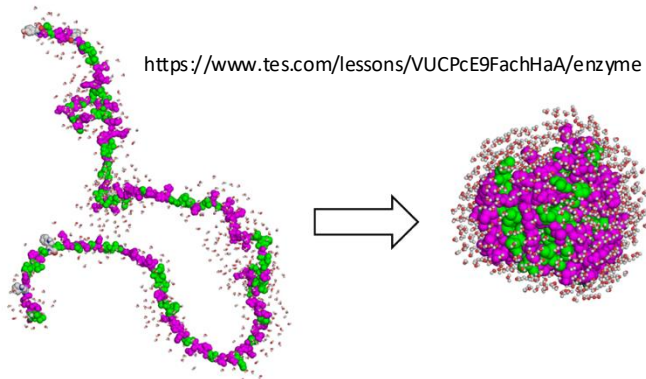
George Gamow, 1961



Cell division

<https://www.thoughtco.com/mitosis-and-cell-division-quiz-4078417>

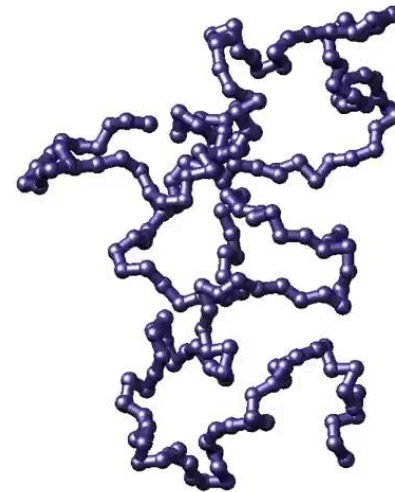
Polypeptide collapse



Unfolded

Folded

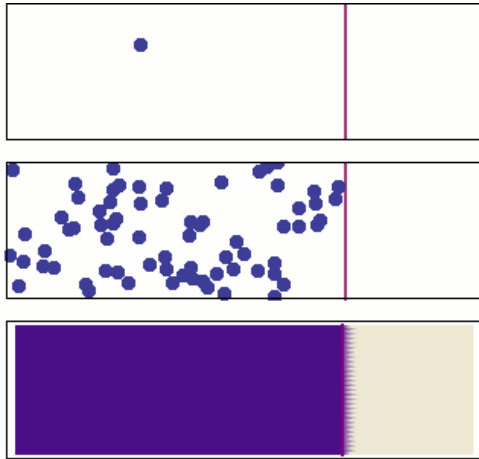
<https://www.tes.com/lessons/VUCPcE9FachHaA/enzyme>



Polymer collapse

https://www.youtube.com/watch?v=ek9vd7_PDjk

Diffusion: unbiased, random movement in space and time



$$\langle r_N^2 \rangle = (d)Na^2$$

τ : time taken for step $(+a)$ or $(-a)$

T: total elapsed time

} No. of steps, $N = \frac{T}{\tau}$

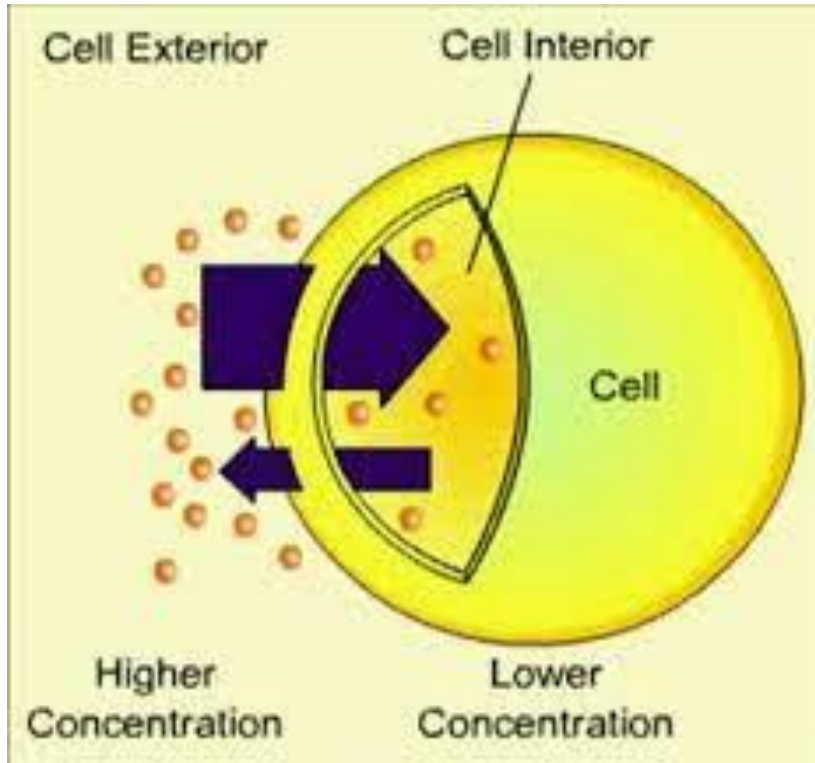
Diffusion Coefficient: $D \equiv \frac{a^2}{2\tau}$

Diffusion Relationship:

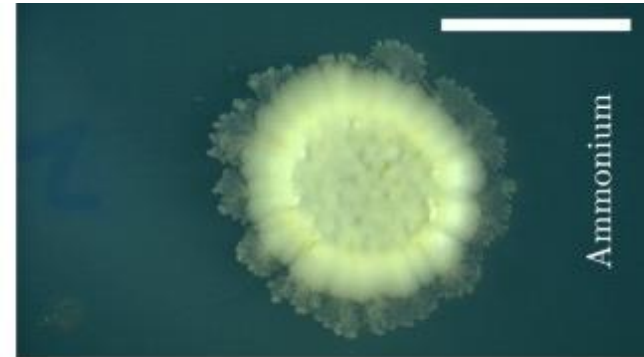
$$\langle r_N^2 \rangle = (2d).D.(elapsed\ time)$$

Examples: Diffusion within Biological Systems

Molecular passage



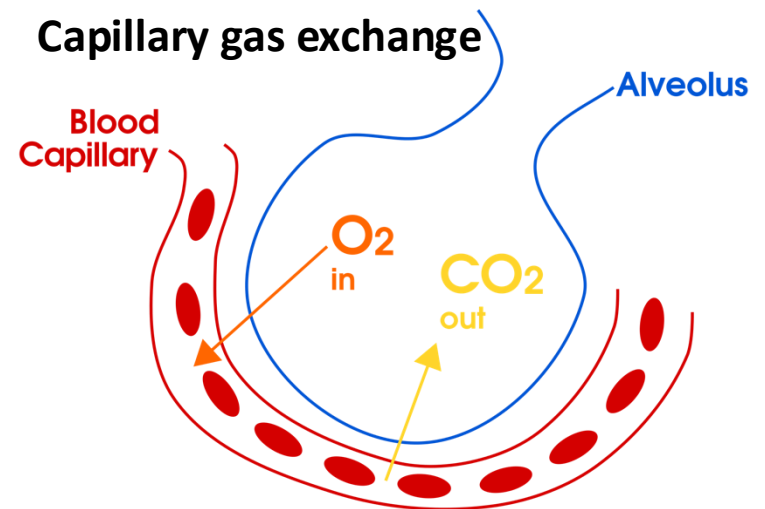
Diffusion limited growth of microbial colonies



(b) SLAD^{-N} with ammonium added

Tronolone et al., Scientific Reports 2018

Capillary gas exchange



Examples: Diffusion within Biological Systems

Apparent Diffusion Coefficient (ADC) as a Cancer Biomarker

OPEN ACCESS Freely available online



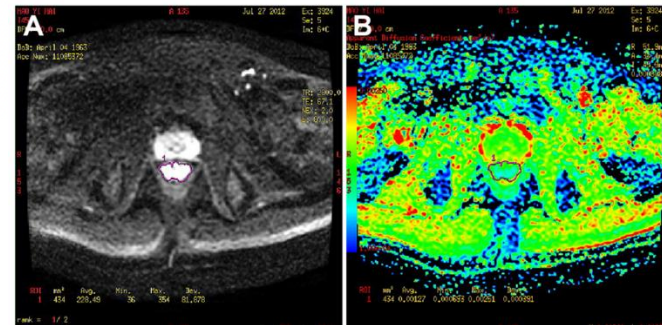
Apparent Diffusion Coefficient (ADC) Value: A Potential Imaging Biomarker That Reflects the Biological Features of Rectal Cancer

Yiqun Sun^{1†}, Tong Tong^{1†}, Sanjun Cai², Rui Bi³, Chao Xin¹, Yajia Gu^{1*}

Int Urol Nephrol (2014) 46:555–561
DOI 10.1007/s11255-013-0557-1

UROLOGY - ORIGINAL PAPER

ADC maps



Apparent diffusion coefficient value as a biomarker reflecting morphological and biological features of prostate cancer

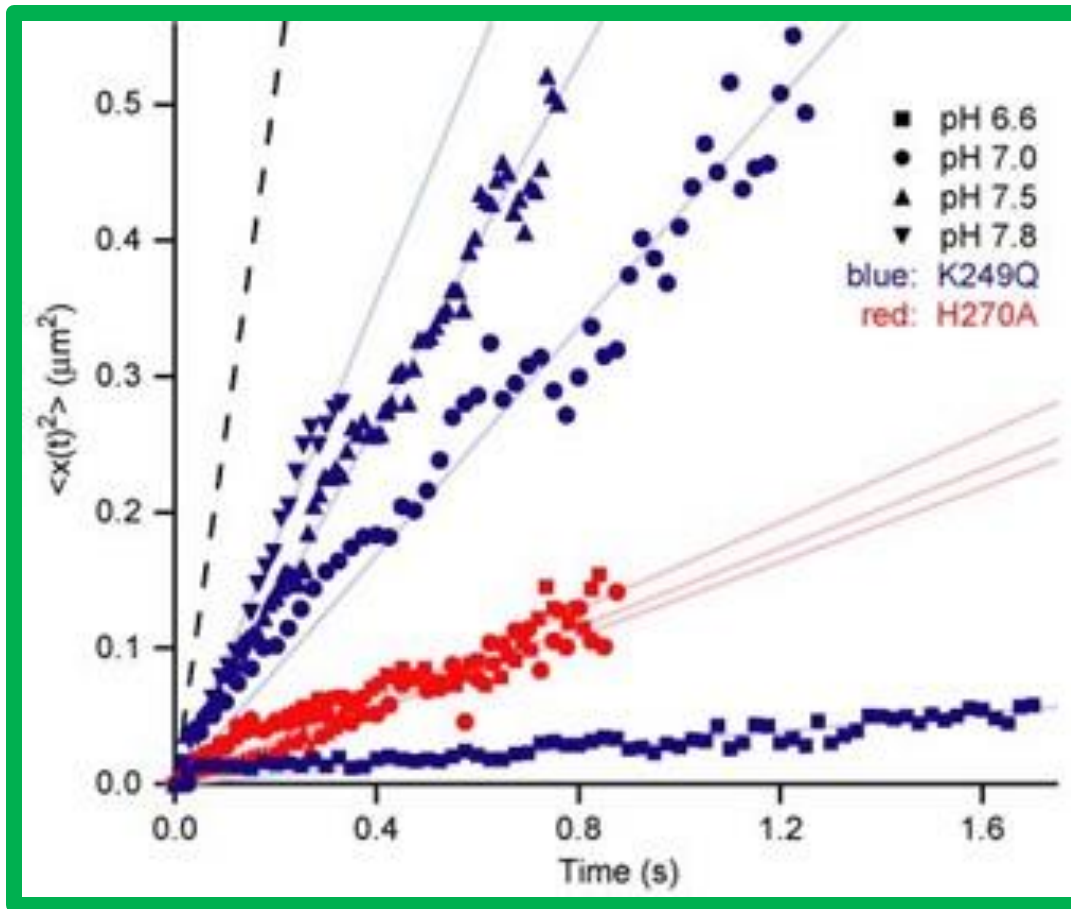
Hyeyeol Bae · Soichiro Yoshida · Yoh Matsuoka · Hiroshi Nakajima ·
Eisaku Ito · Hiroshi Tanaka · Miyako Oya · Takayuki Nakayama ·
Hideki Takeshita · Toshiki Kijima · Junichiro Ishioka · Noboru Numao ·
Fumitaka Koga · Kazutaka Saito · Takumi Akashi · Yasuhisa Fujii ·
Kazunori Kihara

Diffusion Tensor

$$\mathcal{D} = \begin{bmatrix} D_{xx} & D_{xy} & D_{xz} \\ D_{yx} & D_{yy} & D_{yz} \\ D_{zx} & D_{zy} & D_{zz} \end{bmatrix}$$

Estimation of Diffusion Coefficient

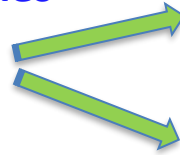
“Single molecule” experiments



Sunny Xie Lab, Harvard Univ.

detour: Binomial Distribution

Only **2 possible outcomes**
of an event with
 N attempts:



Success, probability s

Failure, probability $f = (1 - s)$

Mean success, $\langle n \rangle = s \cdot N$ n = No. of success

Probability of [n success and $(N-n)$ failure] is,

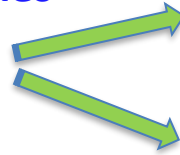
$$P(n, s, N) = \frac{N!}{n! (N-n)!} s^n (1-s)^{N-n}$$

The sum of probabilities for the (n, s, N) is 1.0, ie.

$$\sum_{n=0}^N P(n, s, N) = 1.0$$

detour: Binomial Distribution

Only **2 possible outcomes**
of an event with
 N attempts:



Success, probability s

Failure, probability $f = (1 - s)$

Mean success, $\langle n \rangle = s \cdot N$ n = No. of success

Variance in success, $\sigma^2 = s(1 - s)N$

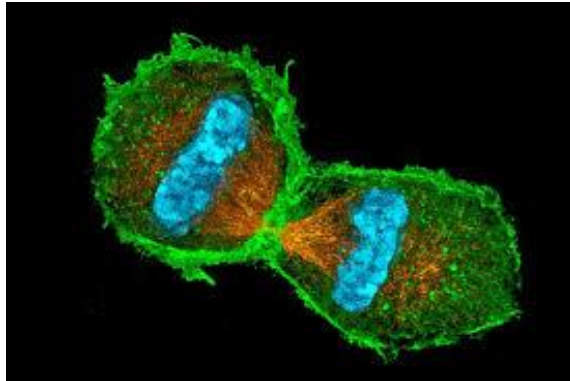
Standard deviation, $\sigma = \sqrt{s(1 - s)N}$

The ratio of std. dev. to mean = $\sqrt{\frac{1-s}{sN}}$ $\propto \frac{1}{\sqrt{N}}$

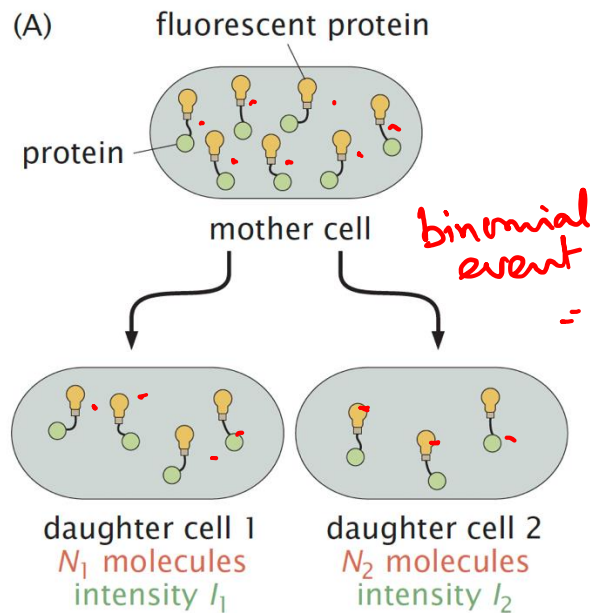
$$r = \frac{\sigma}{\mu}$$

Molecular Partitioning During Cell Division: Binomial Events?

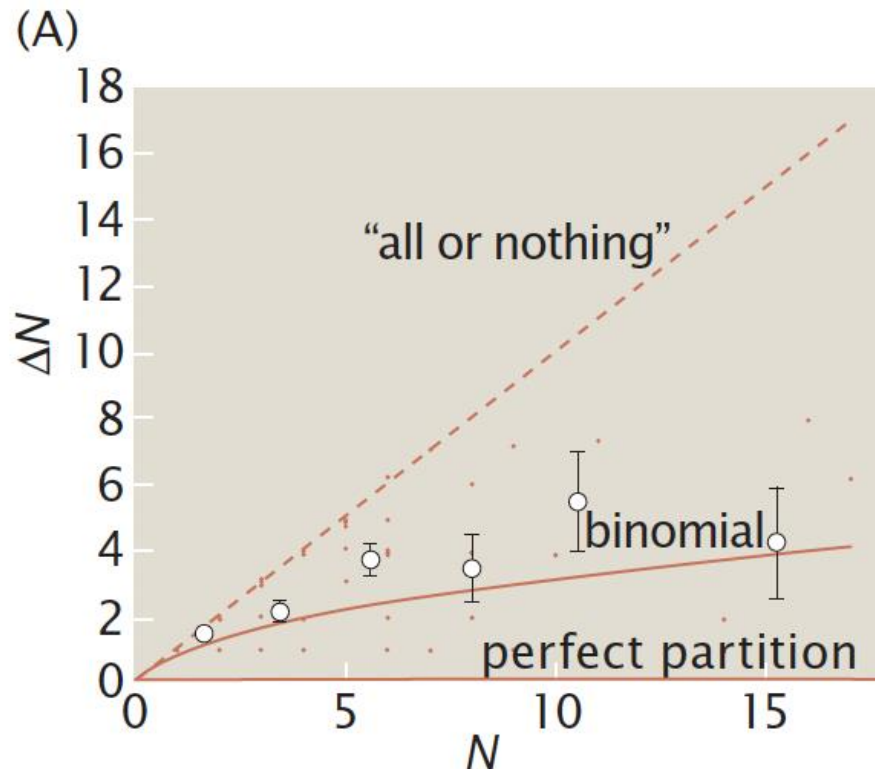
“Count” fluorescent proteins



$$\Delta N = |N_{d(1)} - N_{d(2)}|$$

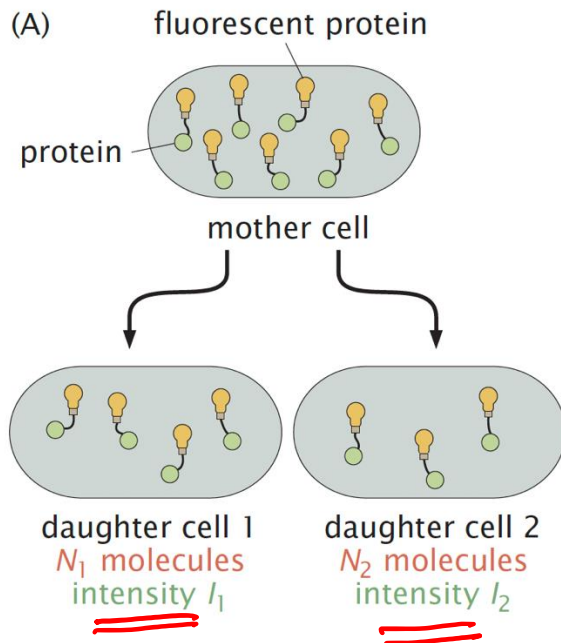
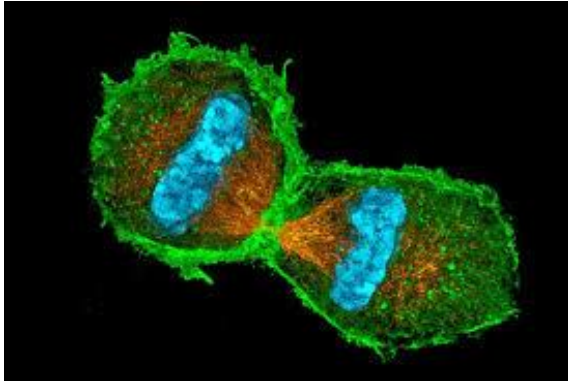


$$N_1 + N_2 = N$$

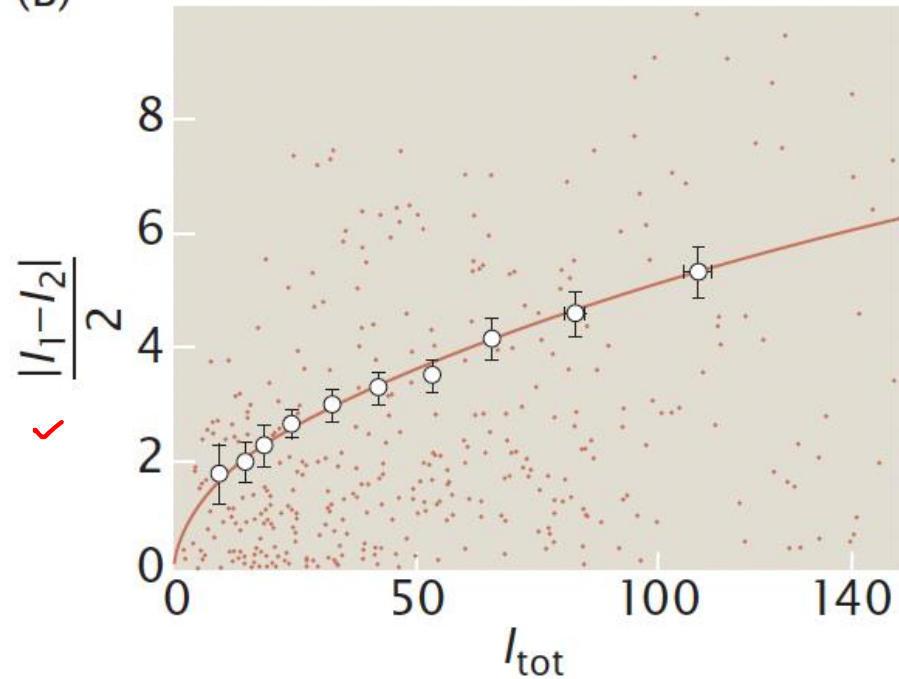


Molecular Partitioning During Cell Division: Binomial Events?

“Count” fluorescent proteins

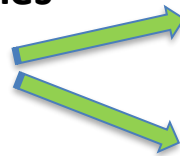


(B)



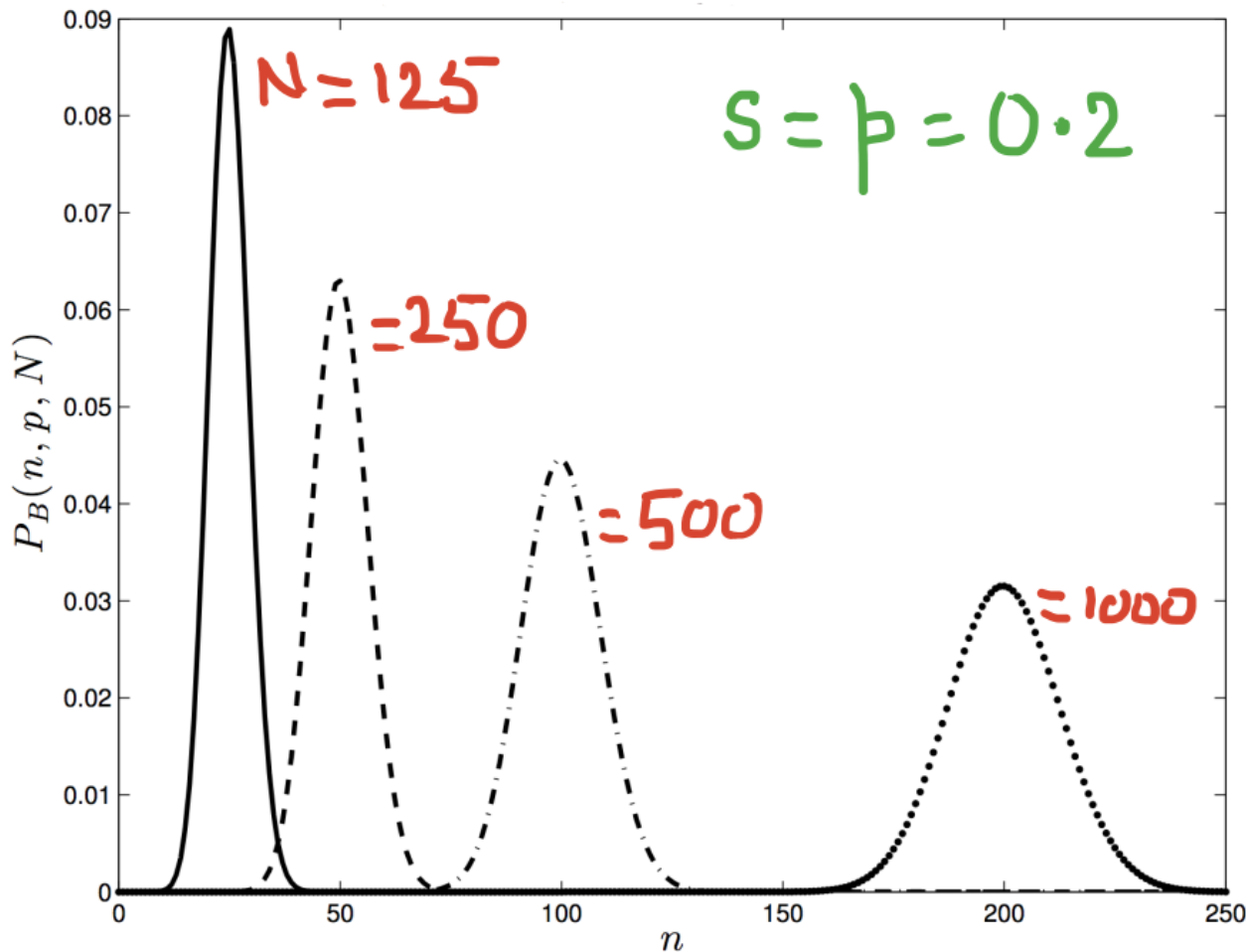
Success (or failure) probability distribution

Only **2 possible outcomes**
of an event with
 N attempts:



Success, probability $s (= p)$

Failure, probability $f = (1 - s)$



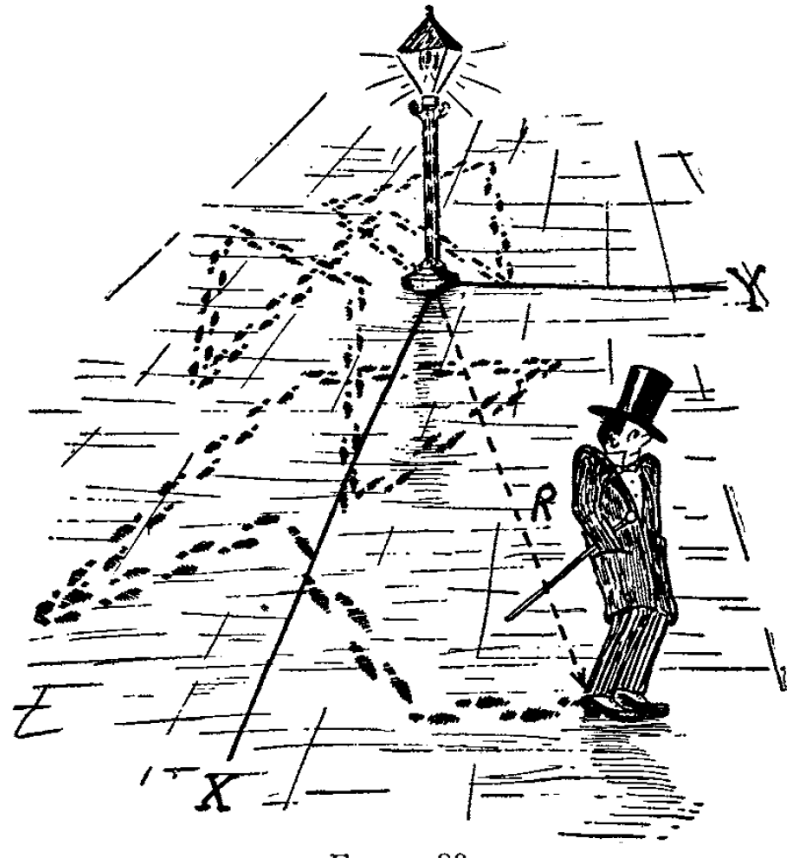
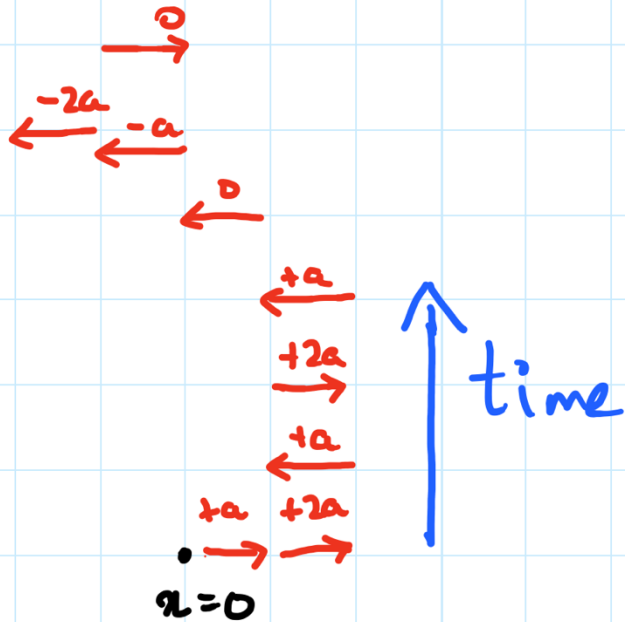
n = No. of success

At large N ,
Binomial disbn.
approaches the
Gaussian Disbn.

Proof:
not required;
request by email if
interested

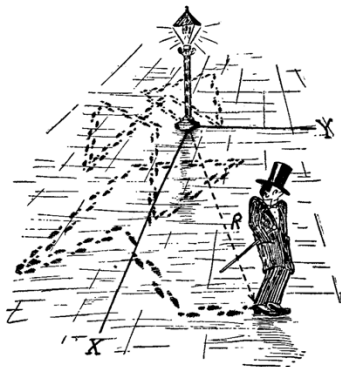
Random Walk

1-Dimensional Random Walk

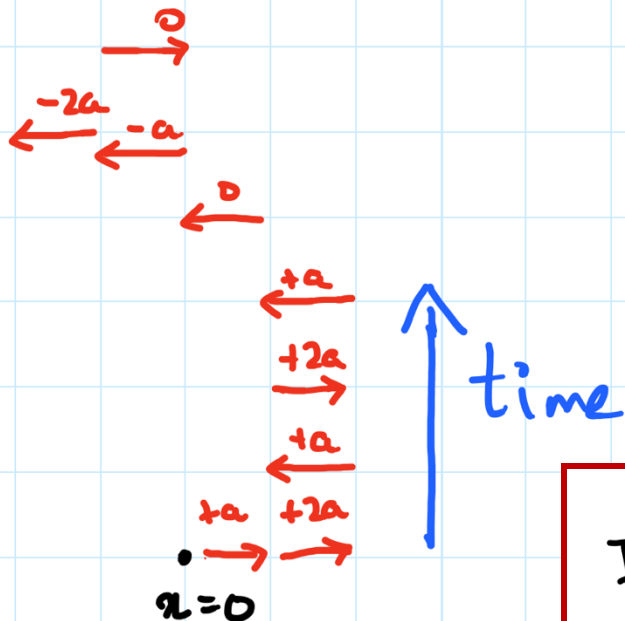


Biol. Physics, Nelson, Ch 2

Random Walk



1-Dimensional Random Walk



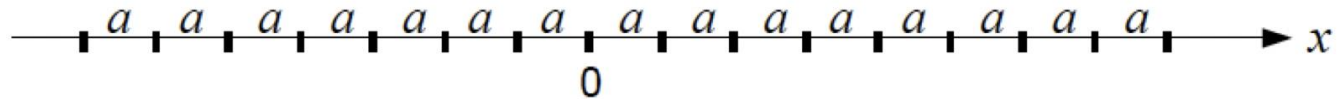
Mean Displacement in "N" steps,
and "n" forward steps is,

$$\begin{aligned}\langle x \rangle &= \sum_{n=0}^N P(n, p, N) [na - (N-n)a] \\ &= a \sum_{n=0}^N \frac{N!}{n!(N-n)!} p^n (1-p)^{N-n} (2n-N) \\ &\equiv Na(2p-1)\end{aligned}$$

skipping
steps

If probability of front and back step are equal,
 $\langle x \rangle = 0$

Mean displacement in unbiased walk: a simpler argument



a ---Length of each step

$x_0=0$ ---start point

x_n ---position after the n -th step

$k_n a$ ---displacement of the n -th step with $P(k_n=1)=P(k_n=-1)=1/2$

$k_n = +1$ (right move)
 $= -1$ (left move)

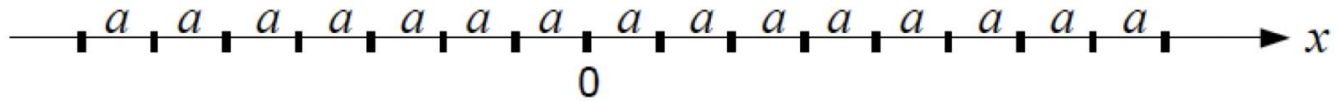
$$x_n = x_{n-1} + k_n a$$

$$\begin{aligned} \langle x_n \rangle &= \langle x_{n-1} + k_n a \rangle \\ &= \langle x_{n-1} \rangle + a \langle k_n \rangle \\ &= \langle x_{n-2} \rangle + a \langle k_{n-1} \rangle \\ &= \dots \langle x_1 \rangle + a \langle k_1 \rangle \end{aligned}$$

(Note: Red dashed arrows point from the $\langle k_n \rangle$ terms to a red '0' at the top right of the page.)

$$\langle x_n \rangle = \langle x_0 \rangle = 0$$

Mean squared displacement in 1-dimension



a ---Length of each step

$x_0=0$ ---start point

x_n ---position after the n -th step

$k_n a$ ---displacement of the n -th step with $P(k_n=1)=P(k_n=-1)=1/2$

$$\langle x_n^2 \rangle = \langle (x_{n-1} + k_n a)^2 \rangle$$

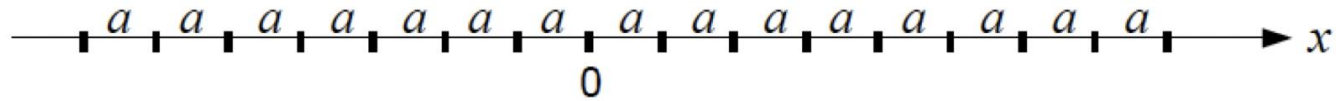
$$= \langle x_{n-1}^2 \rangle + \langle k_n^2 \rangle a^2 + 2a \langle k_n x_{n-1} \rangle$$

$$= \langle x_{n-1}^2 \rangle + a^2 = \langle x_{n-2}^2 \rangle + 2a^2 \quad \rightarrow 0$$

$$\langle k_n^2 \rangle = \frac{(+1)^2 + (-1)^2}{2} = 1$$

$$\begin{aligned} \langle k_n x_{n-1} \rangle &= x_{n-1} \times (+1) \times P(+1) \\ &\quad + x_{n-1} \times (-1) \times P(-1) \\ &= x_{n-1} \left[(+1) \times \left(\frac{1}{2}\right) + (-1) \times \left(\frac{1}{2}\right) \right] \\ &= 0 \end{aligned}$$

Mean squared displacement in 1-dimension



a ---Length of each step =

$x_0=0$ ---start point

x_n ---position after the n -th step

$k_n a$ ---displacement of the n -th step with $P(k_n=1)=P(k_n=-1)=1/2$

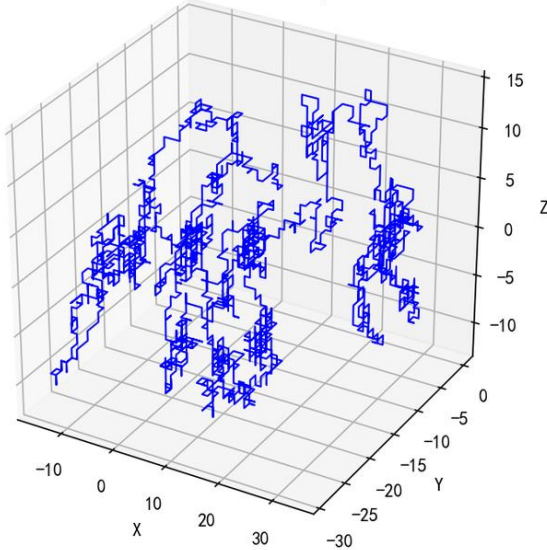
$$\begin{aligned}\langle x_n^2 \rangle &= \langle (x_{n-1} + k_n a)^2 \rangle \\ &= \langle x_{n-1}^2 \rangle + \langle k_n^2 \rangle a^2 + 2a \langle k_n x_{n-1} \rangle \\ &= \langle x_{n-1}^2 \rangle + \langle a^2 \rangle\end{aligned}$$

By iteration, for N steps,

$$\langle x_N^2 \rangle = N a^2$$

Mean squared displacement in 2- or 3- dimensions

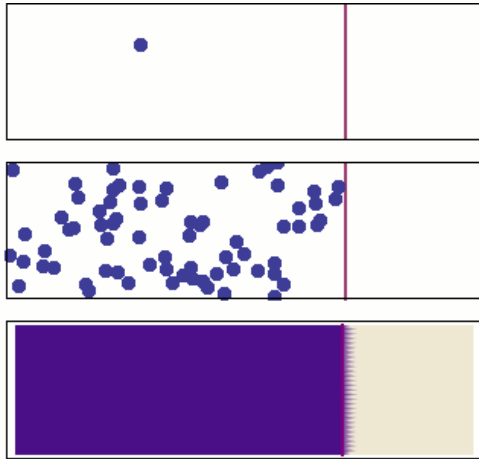
Random walk in 3D space



HW. Show that in ' d ' spatial dimensions, $\langle \mathbf{r}_N^2 \rangle = (d)Na^2$
Work out all steps.

$$\begin{aligned}
 \langle r_N^2 \rangle &= \langle x_N^2 + y_N^2 + z_N^2 \rangle \\
 &= \langle (x_{N-1} + k_{xN}a)^2 \rangle + \dots + \dots \\
 &= \left(\langle x_{N-1}^2 \rangle + 2a \langle \cancel{x_{N-1}} \cancel{k_{xN}} \rangle + \langle k_{xN}^2 \rangle a^2 \right) \\
 &\quad + (\dots) + (\dots)
 \end{aligned}$$

Diffusion: average spatio-temporal pattern of random walks



In 2-dimensions:

$$\langle \mathbf{r}_N^2 \rangle = (d)Na^2$$

τ : time taken for step $(+a)$ or $(-a)$

T: total elapsed time

No. of steps, $N = \frac{T}{\tau}$

define **Diffusion Coefficient:** $D \equiv \frac{a^2}{2\tau}$

$$\begin{aligned} \langle r_N^2 \rangle &= 2Na^2 \\ &= 2\left(\frac{T}{\tau}\right)a^2 \\ &= (2\tau)(2D) \end{aligned}$$

Diffusion Relationship:

$$\langle r_N^2 \rangle = 4DT$$

HW. Generalize to ' d ' dimensions, ie. $\langle \mathbf{r}_N^2 \rangle = (2d)DT$

Diffusion: average spatio-temporal pattern of random walks

Diffusion is related to friction

EINSTEIN RELATIONSHIP

$$\zeta D = k_B T$$

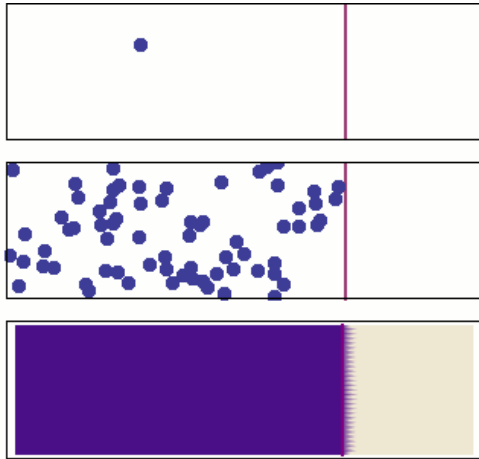
Viscous Friction Coefficient

At a given temperature, viscosity is inversely proportional to the diffusion coefficient.

STOKES-EINSTEIN RELATIONSHIP

$$6 \pi \overset{\text{viscosity}}{\eta} \overset{\text{radius}}{a} D = k_B T$$

Diffusion: movement of particles in unbiased random walks in any dimension



$$\langle r_N^2 \rangle = (d)Na^2$$

τ : time taken for step (+a) or (-a)

T: total elapsed time

No. of steps, $N = \frac{T}{\tau}$

define **Diffusion Coefficient:** $D \equiv \frac{a^2}{2\tau}$

Diffusion Relationship:

$$\langle r_N^2 \rangle = (2d)D(t)$$

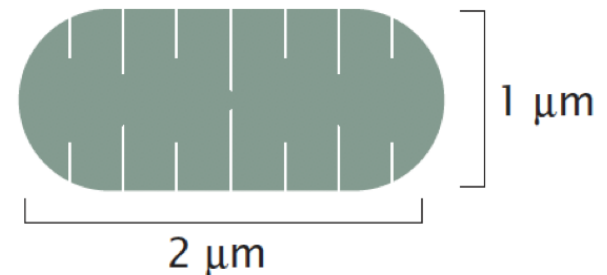
Stokes-Einstein (SE) Relationship:

$$6\pi\eta a D = k_B T$$

- Consider a ~spherical protein of radius 15 nm diffusing ("wandering aimlessly") in water.
- How long will it take to transverse *E. coli* length?
- At 298 K,

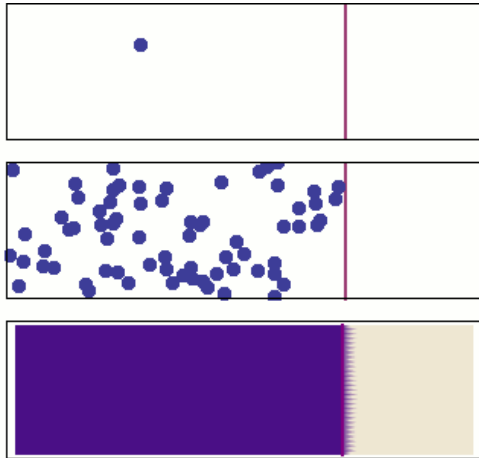
$$\eta_{H_2O} = 8.9 \times 10^{-4} \text{ Pa}\cdot\text{s}$$

$$= 0.890 \text{ centiPoise}$$



approx. *E. coli* dimensions

Diffusion: movement of particles in unbiased random walks in any dimension



$$\langle r_N^2 \rangle = (d)Na^2$$

τ : time taken for step $(+a)$ or $(-a)$

T: total elapsed time

No. of steps, $N = \frac{T}{\tau}$

define **Diffusion Coefficient:** $D \equiv \frac{a^2}{2\tau}$

Problem (Hw)

- Consider a ~spherical protein of radius **15 nm** ($= a$) diffusing ("wandering aimlessly") in water.
- How long will it take to transverse *E. coli* length?
- At 298 K, $\eta_{H_2O} = 8.9 \times 10^{-4} \text{ Pa}\cdot\text{s}$
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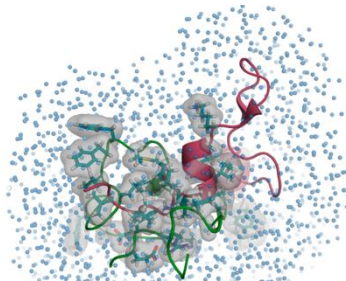
$\left\{ \begin{array}{l} \text{Pa}\cdot\text{s} : \text{SI} \\ \text{centiPoise} : \text{CGS} \end{array} \right.$

Diffusion Relationship:

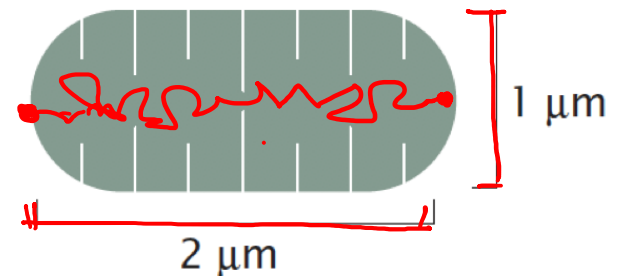
$$\langle r_N^2 \rangle = (2d)D(t)$$

Stokes-Einstein (SE) Relationship:

$$6\pi\eta a D = k_B T$$

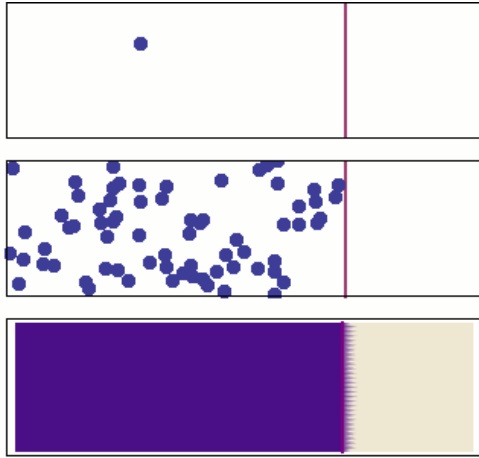


protein in water



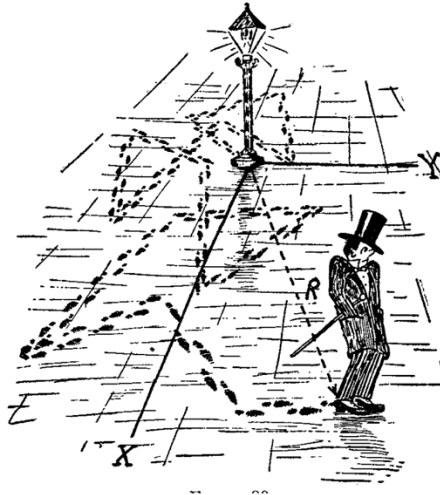
approx. *E. coli* dimensions

In General: Random Biological Processes



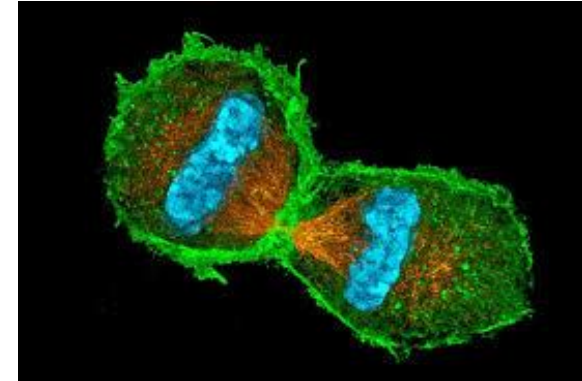
Diffusion

<https://commons.wikimedia.org/w/index.php?curid=8995324>



Random Walk

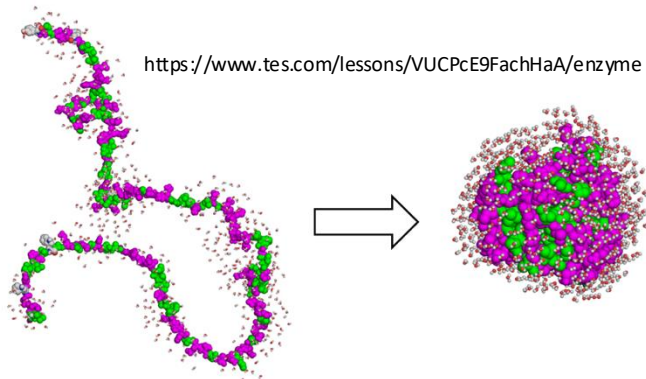
George Gamow, 1961



Cell division

<https://www.thoughtco.com/mitosis-and-cell-division-quiz-4078417>

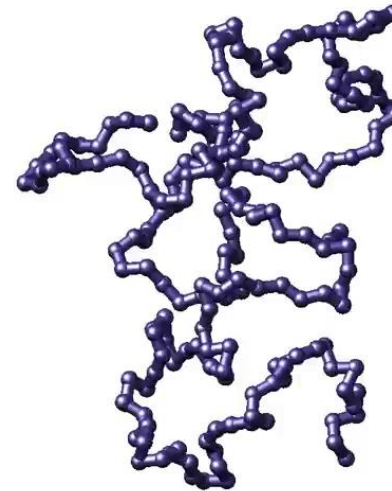
Polypeptide collapse



Unfolded

Folded

<https://www.tes.com/lessons/VUCPcE9FachHaA/enzyme>

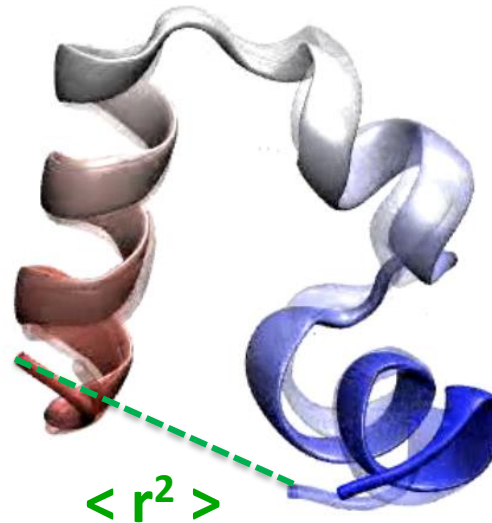


Polymer collapse

https://www.youtube.com/watch?v=ek9vd7_PDjk

Folding marker: **End-to-end distance**

Computationally folded structure superimposed on
experimentally determined crystal structure

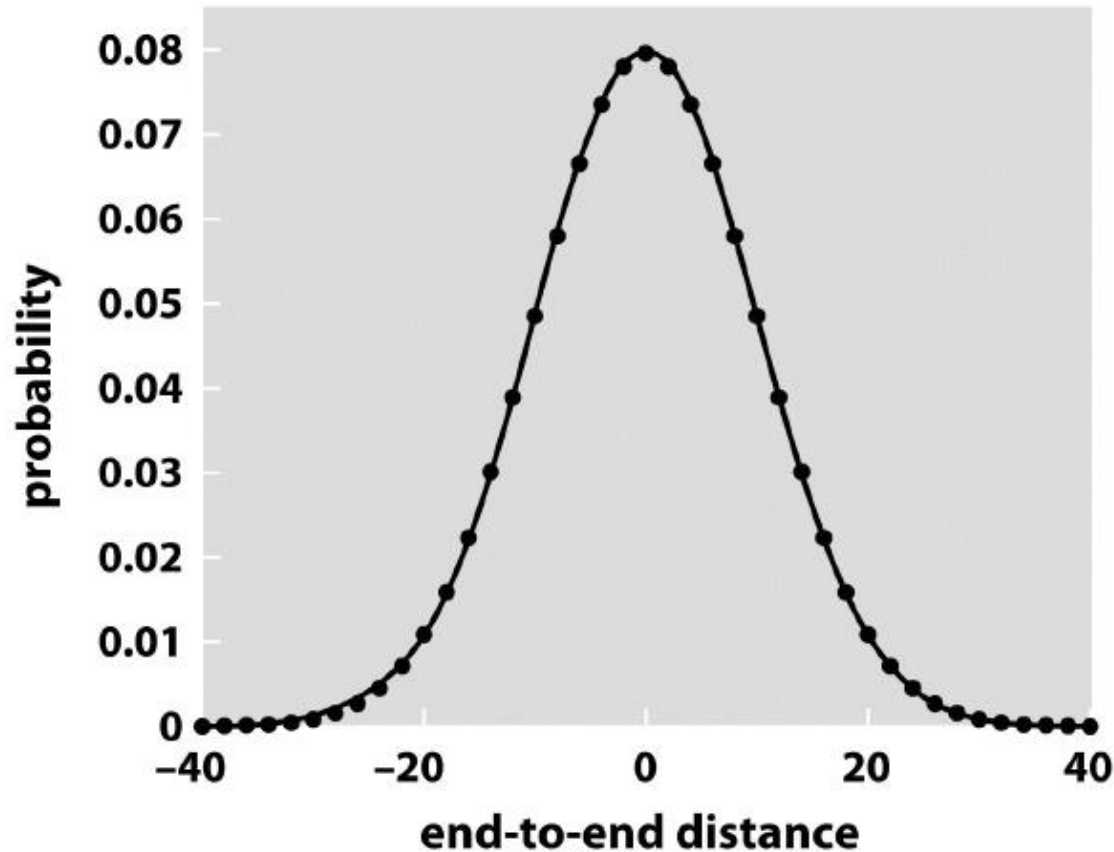


0:30.47

7584.0 ns

End-to-end distance distribution of protein samples with 'N' amino acids

$$P(R; N) = \frac{1}{\sqrt{2\pi N a^2}} e^{-R^2 / 2N a^2}$$



Parameter: $N=100$, $a=1/2$

Line: Gaussian distribution

Dot: binomial distribution

$$\langle R \rangle = 0, \quad \langle R^2 \rangle = N a^2$$