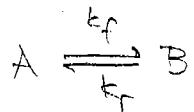


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BP 297

Stochastics & Fluctuations

The biochemical reactions considered so far are deterministic.



$$\frac{dA}{dt} = -k_f A + k_r B$$

$$\frac{dB}{dt} = k_f A - k_r B$$

For given initial conditions, at a given time  $t$ , the results are always exactly the same. But in real biological systems, we have 2 problems: fluctuations, and the small numbers problem.

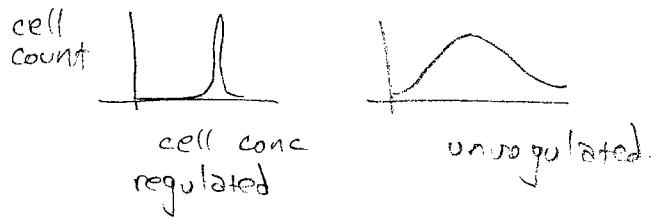
Small numbers problem: Compute the number of  $H^+$  atoms in an E coli: it's about 100. How many chromosomes in an E coli? 1.

Fluctuations matter. Examples

(1) In a race, it's not the average rate that matters. It's who gets there first. Also, activated kinetics is not about average molecules; it's about the highest-energy molecules.

② Biology works hard to beat the fluctuations

A Beckszi & L Serrano Nature 405 590 (2000)

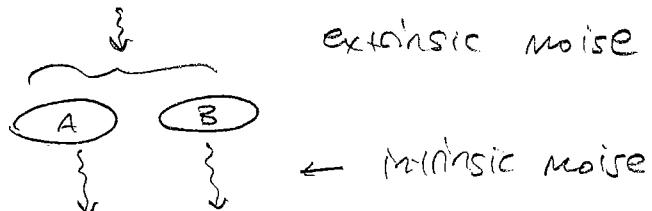


③ Stochastics determine biologically important variability

- Noise generates errors in DNA replica (Rao et al Nature 420 231 (2002)).
- Noise causes divergence of cell fates, creates cell individuality: lysis-lysogeny path, for example (J Hasty et al PNAS 97 2075 (2000)).
- Aggregation is very sensitive to flucs in prot conc: one person gets Alzheimer's; another doesn't.
- Noise causes divergence of cell fates, creates cell individuality: lysis-lysogeny path, for example (J Hasty et al PNAS 97 2075 (2000)).

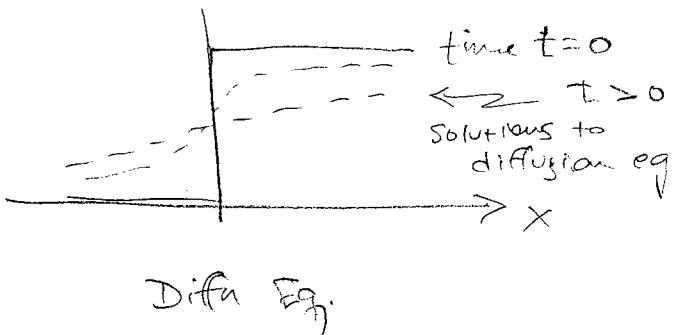
④ Recent work: extrinsic vs intrinsic noise:

- (a) Swain et al PNAS 99 12,795 (2002), (b) Elowitz et al Science 297 1183 (2002), (c) Erin O'Shea.

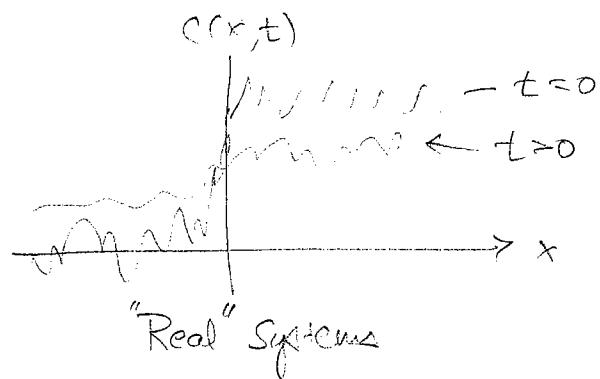


## First, Diffusion

The diffusion equation does not describe fluctuations  $c(x, t)$



Diff Eq.



Why? Because Fick's Law  $J = -D \frac{\partial c}{\partial x}$

is about averages,  $\langle J \rangle = -D \frac{\partial \langle c \rangle}{\partial x}$ , not about distributions. Here's where it comes from

$$(1) \quad \langle J \rangle = \langle c \rangle \langle v \rangle$$

$$(2) \quad \text{force } \langle f \rangle = \overline{\gamma} \langle v \rangle$$

$$(3) \quad \text{force } \langle f \rangle = - \frac{\partial \mu}{\partial x} = -kT \frac{\partial \ln c(x)}{\partial x}$$

Combine:

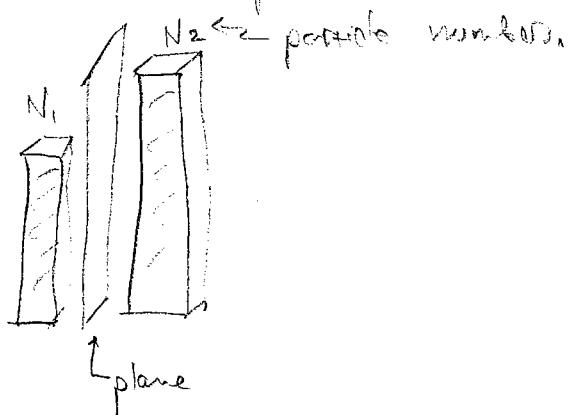
$$\langle J \rangle = - \frac{c k T}{\overline{\gamma}} \frac{\partial \ln c}{\partial x} = - \frac{k T}{\overline{\gamma}} \frac{\partial c}{\partial x} = -D \frac{\partial c}{\partial x}$$

$$\text{where } D = \frac{k T}{\overline{\gamma}}.$$

1

Let's use a 2-layer model instead to capture fluctuations

The basic question: What is the physical driving force for flow of particles?



Thermo says the number of config in this system is  $(N_1 + N_2)!$

$$\Omega = \frac{(N_1 + N_2)!}{N_1! N_2!}$$

$$= -N_1 \ln\left(\frac{N_1}{N}\right) - N_2 \ln\left(\frac{N_2}{N}\right)$$

The entropy  $S = \ln \Omega$  is maximal when  $N_1^* = N_2^*$ , so that's the state of equilibrium.

Our question is how it gets there. A particle can jump from either column to the other in time  $\Delta t$  with probability  $p$ . It stays, with probability  $q = 1-p$ .

Unlike thermo, where we define states, for dynamics we define a process that takes place between time  $t$  and  $t + \Delta t$ , in which  $m_1$  particles jump from column 1 and  $m_2$  particles jump from column 2. Particles are independent and jumps are too.

(5)

## Principle of Maximum Caliber

We write a dynamical position function

$$W(n, N_1, n_2, N_2) = \left[ P^{m_1} g^{N_1 - m_1} \frac{N_1!}{m_1! (N_1 - m_1)!} \right] \left[ P^{m_2} g^{N_2 - m_2} \frac{N_2!}{m_2! (N_2 - m_2)!} \right]$$

that applies between times  $t$  and  $t + \Delta t$  when  $N_1 = N_1(t)$  and  $N_2 = N_2(t)$ .

In the same spirit that the Second Law says to maximize  $S$  to predict equilibrium, now too for dynamics, we maximize  $W$  over all possible processes (ie over  $m_1$  and  $m_2$ ), to predict the flux. Since they are independent, we maximize  $W_1$  and  $W_2$  separately.

$$\frac{\partial \ln W_1}{\partial m_1} \Bigg|_{N_1, m_1 = m_1^*} = 0$$

$$\ln W = m_1 \ln P + (N_1 - m_1) \ln g + N_1 \ln N_1 - m_1 \ln m_1 - (N_1 - m_1) \ln (N_1 - m_1)$$

We call  $C = \ln W$  the caliber.

(6)

So

$$\frac{d\ln N_1}{dm_1} = \ln P - \ln g - \ln m_1 - 1 + \ln(N_1 - m_1) + 1 = 0$$

$$\Rightarrow \ln \left( \frac{m^*}{N-m^*} \right) = \ln \left( \frac{P}{1-P} \right) = \ln \alpha$$

$$\Rightarrow m^* = \alpha N - \alpha m^*$$

$$\Rightarrow \frac{m^*}{N} = \frac{\alpha}{1+\alpha} = P$$

$$\Rightarrow \boxed{m^* = PN}$$

So, the flux is

$$\langle J \rangle = m_1^* - m_2^* = P(N_1(t) - N_2(t))$$

which is equivalent to Fick's law

$$\langle J \rangle = -D \frac{\Delta C}{\Delta x} = -D \Delta C \text{ if } \Delta x = 1 \text{ unit.}$$

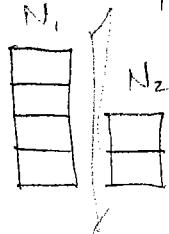
\* The principle of Maximum Caliber recovers for us Fick's Law for the mean flux at time  $t$ .

Now, let's look at the fluctuations

\* Note that this derivation of Fick's Law does not require an equilibrium assumption  $f = -\frac{d\mu}{dx}$ , where  $\mu = kT\ln c$ , or averaging  $\langle J \rangle = \langle c \rangle v$ .

## Now, the Fluctuations

Simple example  $N_1 = 4; N_2 = 2$



For simplicity  $p = q = \frac{1}{2}$  : Multiplicity table

$m_1$	0	1	2	
0	1	2	1	
1	4	8	4	
2	6	12	6	
3	4	8	4	
4	1	2	1	

Maxwell's Demon. (backflux).

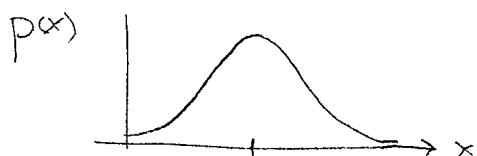
$m_1^* = 2 \quad m_2^* = 1 \Rightarrow \langle J \rangle = 1$

superflux

You can use  $\mathcal{W}$  to compute the relative populations of all the possible fluxes.

## Some Properties of the Fluctuations

Note that a binomial distribution resembles a gaussian function, for large enough  $x$



(8)

$$p(x) \propto e^{-\frac{(x - \langle x \rangle)^2}{2\sigma^2}} \approx e^{-\frac{(m_1 - N_1 p)^2}{2N_1 p g}}$$

where the mean is  $\langle x \rangle = N_1 p = m_1^*$

and standard dev is  $\sigma_1 = \sqrt{N_1 p g}$

We are interested in the distribution of flux,  $P(J)$ , not the distributions of  $m_1$  &  $m_2$  individually.

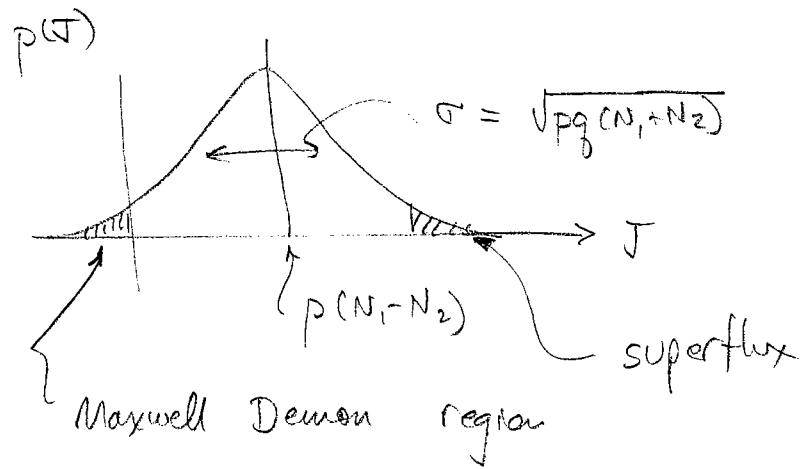
$$J = m_1 - m_2. \quad (1)$$

A law of gaussian functions: If you have a gaussian in  $m_1$  with mean  $m_1^*$  and STD  $\sigma_1$  and a gaussian in  $m_2$  with mean  $m_2^*$  and "  $\sigma_2$ , then the distribution of the difference, Eq (1) (or sum) is also a gaussian, with mean  $(m_1^* - m_2^*)$  and STD  $\sigma^2 = \sigma_1^2 + \sigma_2^2$ , so

$$P(J) \propto e^{-\frac{(J - \text{PAN})^2}{2pgN}}$$

where  $\Delta N = N_1 - N_2$  and  $N = N_1 + N_2$

(9)



Some properties of  $p(J)$ .

- ① The  $\frac{\text{std}}{\text{mean}}$  indicates the size of backflux.

$$= \frac{\sqrt{pqN}}{pfN} \sim \frac{1}{N^{1/2}} \quad \text{which} \rightarrow 0 \text{ as } N \uparrow.$$

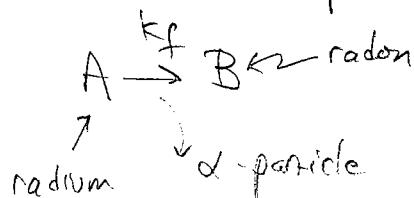
where  $f = \frac{N_1 - N_2}{N_1 + N_2}$ .  $\Rightarrow$  Small numbers problem: noise decreases as system size gets larger.

- ② Even if  $N$  gets large, if  $\Delta N \rightarrow 0$ , there's substantial backflux.

### Now, Chemistry & Fluctuations

$A \xrightleftharpoons[k_r]{k_f} B$  The rate coefficients  $k_f$  and  $k_r$  are for average rates. Sometimes an A converts faster, and sometimes slower, than this.

Best-known example. Radioactive decay



(10)

If you have  $N$  radium atoms, then in a time interval  $t$ , the probability is

$$P(m) = \frac{a^m e^{-a}}{m!} \quad m = 0, 1, 2, \dots$$

that  $m$  of them will decay in that time interval.  
This is a Poisson distribution  $\downarrow a = m^* = Np$ .

Now, we show that the Poisson distribution is just a manifestation of the model above when

- (1)  $p \ll q$
- (2)  $N \rightarrow \infty$  such that  $a = pqN$  is finite.

Again, start with

$$P(m, N) = P^m q^{N-m} \frac{N!}{m!(N-m)!} \quad (2)$$

We will use

$$\lim_{N \rightarrow \infty} \left(1 - \frac{a}{N}\right)^N = e^{-a}.$$

So, re-write Eq (2) as

$$\begin{aligned}
 P(m, n) &= p^m \frac{(1-p)^n}{(1-p)^m} \frac{n!}{(n-m)!} \frac{1}{m!} \\
 &= p^m \frac{\left(1 - \frac{a}{N}\right)^n}{(1-p)^m} N^m \frac{1}{m!} \\
 &= \frac{p^m e^{-a}}{m!}
 \end{aligned}$$

where  $(1-p)^m \approx 1$  since  $m$  is small and  $p \ll 1$ , and  $N(N-1)\dots(N-m+1) \approx N^m$ .

### Waiting Times

Single-molecule experiments often measure a quantity related to the Poisson dist; called the waiting time distribution. The WTD is the distribution of times you wait between events, rather than the number of events per unit time. For situations that give the Poisson dist for the latter, the WTD is exponential, ie

$$\text{WTD} \quad g(t) = \lambda e^{-\lambda t} \quad (3)$$

$$\text{PD if } p(m, t) = \frac{(\lambda t)^m e^{-\lambda t}}{m!}$$

We can derive  $g_t(t)$  from  $p_t(t)$ , or use MaxCal again in a simple way.

$$S = \sum g_t \ln g_t \quad \text{subject to}$$

$$\sum g_t = 1$$

$$\sum t g_t = \langle t \rangle = \bar{t}$$

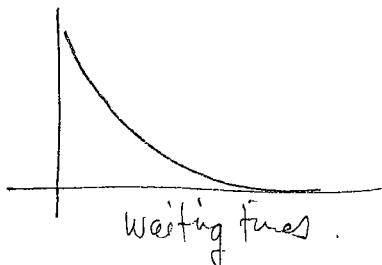
$$dC = - \sum (-1 - \ln g_t^* - \alpha - \lambda t) dg_t = 0$$

$$\Rightarrow g_t^* = \frac{e^{-\lambda t}}{\sum_t e^{-\lambda t}} = \frac{1}{Z} e^{-\lambda t}$$

$$\text{To normalize } g_t^* = \lambda e^{-\lambda t} \Rightarrow Z = \frac{1}{\lambda}$$

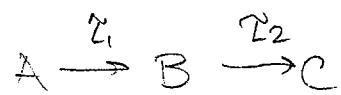
$$\text{so } \langle t \rangle = \bar{t} = \sum t g_t^* = \sum t e^{-\lambda t} = - \frac{\partial \ln Z}{\partial \lambda}$$

$$\text{so } \boxed{g_t^* = \lambda e^{-\lambda t} = \frac{1}{\lambda} e^{-t/\lambda}}$$



This is the waiting time used also in the Gillespie algorithm.

Two-Step Rxns.



$$g_A(t_1) = \frac{1}{\gamma_1} e^{-t/\gamma_1}$$

$$g_B(t_2) = \frac{1}{\gamma_2} e^{-t/\gamma_2}$$

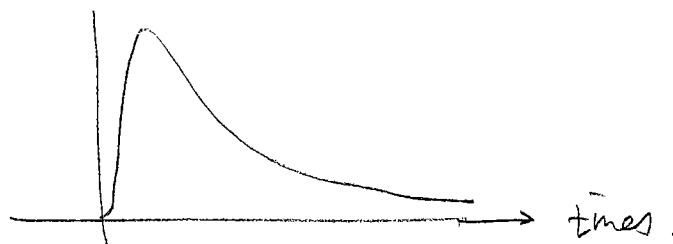
From A to C, we have

$$\begin{aligned} g(t) &= \int_0^t g_A(t_1) g_B(t-t_1) dt_1 \\ &= \int \left( \frac{1}{\gamma_1} e^{-t/\gamma_1} \right) \left( \frac{1}{\gamma_2} e^{-(t-t_1)/\gamma_2} \right) dt_1 \\ &= \left( \frac{1}{\gamma_1 \gamma_2} \right) e^{-t/\gamma_2} \underbrace{\int_0^t e^{-t_1 \left( \frac{1}{\gamma_1} - \frac{1}{\gamma_2} \right)} dt_1}_{C^{-t \left( \frac{1}{\gamma_1} - \frac{1}{\gamma_2} \right)} - 1} \\ &\quad - \left( \frac{1}{\gamma_1} - \frac{1}{\gamma_2} \right) \end{aligned}$$

$$= \frac{1/(\tau_1 \tau_2)}{\left(\frac{1}{\tau_1} - \frac{1}{\tau_2}\right)} (e^{-t/\tau_2} - e^{-t/\tau_1})$$

$$= \frac{1}{\tau_2 - \tau_1} (e^{-t/\tau_2} - e^{-t/\tau_1})$$

which looks like this



Langmuir Eqn / Gillespie / Stochsim

$$m \frac{dv}{dt} = -\xi v + f(t) \quad \text{Langmuir}$$

fluctuating force with props

$$\langle f(t) \rangle = 0$$

$$\langle f(t) f(t') \rangle = A \delta(t-t')$$

What you can do with this is to get various time correlation fns. For instance, multiply by  $v(0)$  and take the avg:

$$m \frac{d}{dt} \langle v(0) v(t) \rangle = -\xi \langle v(0) v(t) \rangle + \underbrace{\langle v(0) f(t) \rangle}_{=0} \\ \text{uncorrelated}$$

Solving gives

$$-\frac{\xi}{m} t$$

$$\langle v(0) v(t) \rangle = A e^{-\frac{\xi}{m} t}$$

Since  $\langle v(0)^2 \rangle = \frac{kT}{m}$ , it means  $A = \frac{kT}{m}$

For complex systems, this can be done by the Gillespie algorithm or STANHSIM:

- D. Gillespie ✓ Phys Chem 81 2390 (1977)
- M. Gibson & J. Bruck ✓ Phys Chem 104 1876 (2000)
- Rao et al Nature 420 231 (2002).