Fick's laws (for diffusion)

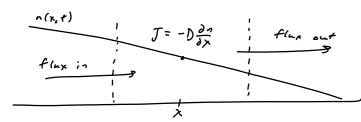
Fick's 1st law:

Diffusive flux goes from regions of high to low concentration with a magnitude proportional to the concentration gradient.

In 1-D,

flux
$$J_{\text{Diffusion}} = -D \frac{\partial n}{\partial x}$$

where D is the diffusive soe therent.



Mass conservation:

$$\frac{\partial n}{\partial t}$$
, $\frac{\partial}{\partial x}J = 0$ (or $\frac{\partial n}{\partial t} + \nabla \cdot J = 0$ in higher dimensions)

$$\Rightarrow \frac{\partial n}{\partial t} = \frac{\partial}{\partial x} \left(D \frac{\partial n}{\partial x} \right) \qquad \text{in} \qquad 1 - D.$$

Density-dependent diffusion

Some animals diffuse differently in response to population pressure. One extension of the classical diffusion model that is relevant to insect dispersal is when there is an increase in diffusion due to population pressure.

For example, the flux $J = -D(n)\nabla n$

where $\frac{dD}{dr} > 0$.

A typical Born is $D(n) = D_0 \left(\frac{n}{n_0}\right)^m$ where m > 0, $D_0 > 0$, $n_0 > 0$.

The dispersal equation is

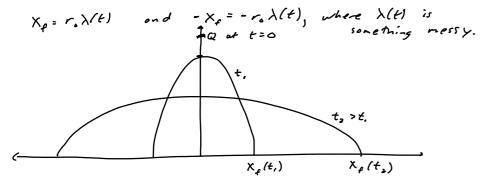
2- in 1-D,

$$\frac{\partial n}{\partial t} = D. \frac{\partial}{\partial x} \left(\left(\frac{n}{n_0} \right)^m \frac{\partial n}{\partial x} \right)$$

not constant with respect to X

with initial condition n(X,0) = Q & (X),
point mass of size Q at X=0 at time O.

The solution is fundamentally different from that of the classical cose when m=0, because D(0)=0. The solution has a wore front at



Chemotaxis

Many insects, animals, cells, etc are attracted up or down a chemical gradient of pheremones, cytokines, etc.

Motion up a chemical concentration is called chemotoxis.

Suppose the presence of a gradient in an attractant a(x,t) gives rise to the movement of individuals up the gradient.

Also, assume the flux of individuals will increase with the number of mandals n(x,t).

So, we may reasonably take the chemotochi flux to be

J= n X(a) Da

a gradient of chemoathroctont

where X(a) is a function of a called the chemotockic coefficient.

The "pure" chemotaxis equation is

$$\frac{\partial a}{\partial t} + \nabla \cdot J = 0$$

$$\Rightarrow \frac{\partial n}{\partial t} + \nabla \cdot (n \pi(a) \nabla a) = 0$$

Suppose we also have diffusion and a growth term f(n), then the basic reaction-diffusion-chemotoxis equation is

$$\frac{\partial n}{\partial t} = f(n) - \nabla \cdot (n \times (a) \nabla a) + \nabla \cdot (D \nabla n)$$
.

Assume attractort a(x,t) is governed by

The classical Keller-Segel model (1971) for a sline mould is

$$\frac{\partial n}{\partial t} = D \frac{\partial^2 n}{\partial x^2} - \sqrt{\partial x} \left(n \frac{\partial a}{\partial x} \right)$$

$$\frac{\partial a}{\partial t} = h_n - t a + D_0 \frac{\partial^2 a}{\partial x^2}$$

$$\begin{cases} \rho roduction & decay \end{cases}$$

A problem is that this model can give ise to solutions that blow up in finite time.

Chemotaxis modelling and analysis is a big are in mathematical bio.

People try to 'tix" the blow up problem by using

- e volume-filling method: there is a max density of cells/individuals n(x,t) at any point.
- one could saturate the effect of the gradient, so that extremely steep gradients don't althout better from very steep gradients.