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Constitutive Expression for N for molecular diffusion

1. Empirical - Fick


2. Theory

- analogy to current & heat
- thermodynamics
- statistical mechanics

Example - classical thermodynamics

- reduce free energy of a chemical system by reducing chemical potential;
- this can be accomplished by net movement of molecular species down a gradient of chemical potential $\nabla \mu_i$

\underline{N} - flux



$$\frac{\text{molecular species}}{\text{area} \cdot \text{time}}$$

c_i - conc. (molecules)

$$\frac{\text{vol}}$$

chemical potential of molecular species, i

$$\underline{N} = \alpha_i c_i \boxed{\nabla \mu_i}$$

↑ mobility of species

driving force

mobility is theoretically derived from movement of a particle in a solvent

← e.g. Stokes-Einstein theory.

Deen 1.5 (size, viscosity, structure)

$$\mu_i = RT \ln(\gamma_i y_i) = \frac{c_i}{c_T}$$

↑ gas constant.

abs. temp.

activity coeff. of species i

← total solution

↑ mole fraction of species in solution

γ_i parameterizes nonideal molecular interactions (% of high concentrations) under dilute conditions ($y_i \ll 1$)

$\gamma_i \rightarrow 1$ (ideal behavior)

issue: many biological systems are non-ideal

$$\therefore \nabla \mu_i = RT \nabla (\gamma_i \ln y_i) \Rightarrow \text{for } \gamma_i = 1$$

$$\underline{N} = -[\alpha_i RT] \nabla c_i$$

diffusion coefficient for species i

$$\therefore \nabla \mu_i = RT \nabla (\ln[\gamma_i y_i])$$

\Rightarrow for $\gamma_i = 1$:

$$\underline{N} = -\underbrace{[\alpha_i RT]}_{D_i} \nabla c_i$$

If $\gamma_i \neq 1$, then D_i is a function of γ_i , which maybe a function of c_i

$$\Rightarrow \underline{N} = -\nabla (D_i c_i)$$

So, if $\gamma_i \neq 1$, D_i is not constant, then

$$\underline{N} = \boxed{-D_i \nabla c_i} - \underbrace{c_i \nabla D_i}_{\text{potential}}$$

Fick's Law

esp. in cellular situations
extra contribution in non-ideal solutions

Back to mass cons. eq:

$$\frac{\partial c}{\partial t} = -\nabla \cdot (-D \nabla c) + R$$

Assume D is constant (no nonidealities)

$$\frac{\partial c}{\partial t} = D \nabla^2 c + R$$

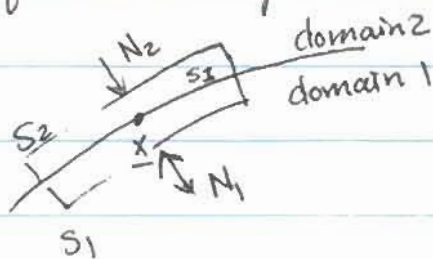
Can we solve this eqn for $c(x, t)$?

Need 1 Initial condition - specify $c(x, t_0) = c(x)$

2 Boundary conditions - @ 2 x locations specify

IC/BC con. laws &/or constitutive expressions

Conservation eqns - around points @ a boundary or interface



But here, as $V \rightarrow 0$

$$\frac{S}{V} \rightarrow \infty$$

so, $\int_V [] dV$ can be neglected.

Then cons. eqn for species is $0 = + \int_{S_1} - \underline{n}_I \cdot \underline{N}_1 dS' - \int_{S_2} - \underline{n}_I \cdot \underline{N}_2 dS' + \int_{S_I} R_s ds$

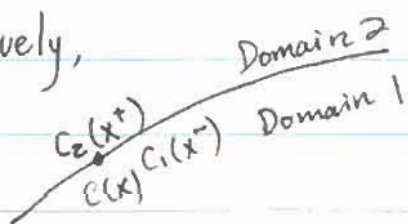
\nwarrow \underline{n}_I is unit normal vector on S_I

\nwarrow net generation @ interfacial surface

Take limit $S \rightarrow 0$ $0 = \underline{n}_I \cdot (\underline{N}_1 - \underline{N}_2) + R_s$

BC = "flux matching"

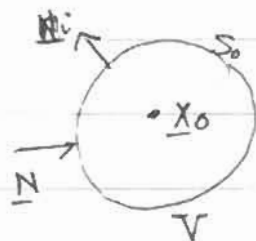
Alternatively,



Assume both domains are in equilibrium w/ one another.

$$\frac{C_2(x^+)}{C_1(x^-)} = K \text{ equilibrium partition coefficient}$$

can have "BC" @ an interior point



no generation and symmetric properties.

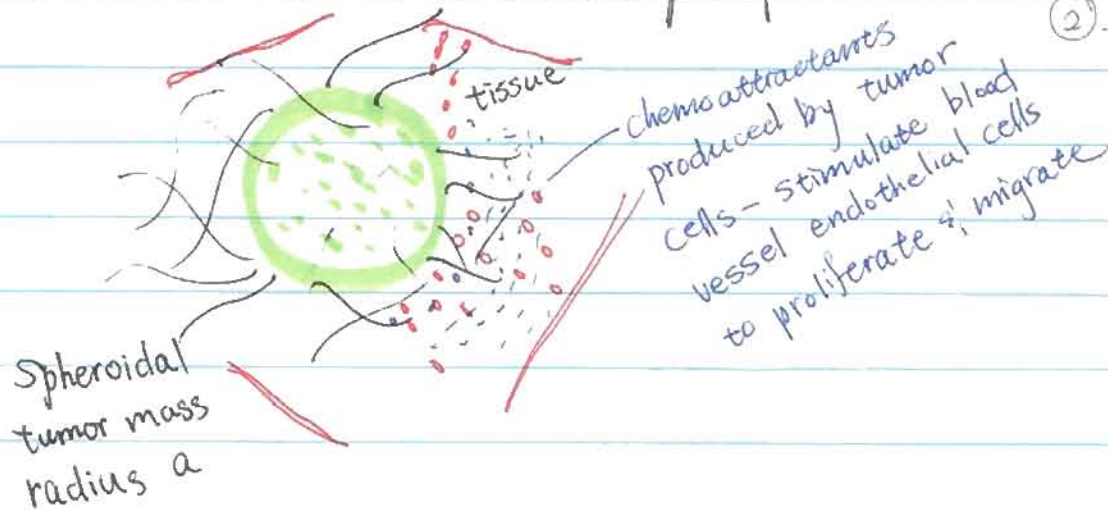
mass cons. eqn

$$\lim V \rightarrow 0 \Rightarrow 0 = - \underline{n}_I \cdot \underline{N} ; \Rightarrow \boxed{0 = \underline{N}} \text{ "no flux" symmetry condition @ an interior point.}$$

Let's formulate a model for an example problem

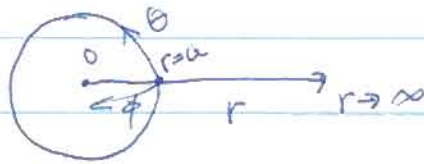
11. m-palmer @

(2)



Determine conc. profile of chemoattractant in tissue

$$\frac{\partial c}{\partial t} = D \nabla^2 c + R \quad \text{in 3-D radial coordinates}$$



Assume angular symmetry (no gradient w/ respect to θ, ϕ)

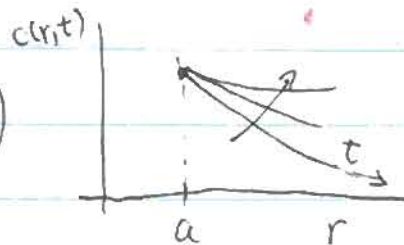
$$\nabla^2 c = \frac{1}{r^2} \frac{\partial}{\partial r} \left(r^2 \frac{\partial c}{\partial r} \right)$$

Assume for $r > a$, $R = 0$

-i.e., negligible degradation by tissue enzymes, negligible uptake by tissue cells, negligible loss to blood cells.

Then

$$\frac{\partial c}{\partial t} = \frac{D}{r^2} \frac{\partial}{\partial r} \left(r^2 \frac{\partial c}{\partial r} \right)$$



To start simply, think about time scales

- attractant profile establishment is much (10x? 100x?).

faster than time scale for tumor size change and endothelial cell response

⇒ assume attractant profile we are interested in is SS for any given α .

i.e. $\frac{\partial c(r)}{\partial t} \rightarrow 0$

$$\frac{\partial c}{\partial t} = 0 = \frac{D}{r^2} \frac{d}{dr} \left(r^2 \frac{dc}{dr} \right)$$

Solution by simple double integration $c(r) = \frac{B_1}{r} + B_2$

(B_1, B_2 are ~~not~~ undetermined coeff).

Determine B_1, B_2 .

BC: $r=a$ flux matching

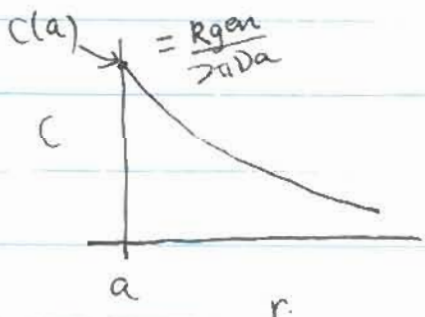
$$0 = R_{gen} + (2\pi a^2) D \frac{dc}{dr} \Big|_{r=a}$$

in tumor
↑
molecules/time
molecules
vol. dist. · time

$r \rightarrow \infty$ $c \rightarrow 0$?
 $\frac{dc}{dr} \rightarrow 0$?

can get B_1, B_2

$$c(r) = \frac{R_{gen}}{2\pi D r}$$



In vitro assay for endothelial cell response to tumor chemoattractant.



- steady state diffusion eqn, same assumptions

$$0 = \frac{D}{r} \frac{\partial}{\partial r} \left(r \frac{\partial c}{\partial r} \right)$$

solve by simple $\int \int$

$c(r) = B_1 \ln r + B_2$ now, in 2D cylindrical coordinates

B_1, B_2 undetermined constants $\nabla^2 c = \frac{1}{r} \frac{\partial}{\partial r} \left(r \frac{\partial c}{\partial r} \right)$

BC $r=a$ flux matching
 $r \rightarrow \infty, c \rightarrow 0$

$$0 = R_{gen} + (2\pi a) D \left. \frac{dc}{dr} \right|_{r=a}$$

no solution