TMA4195 Mathematical Modelling Project

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Deriving the modelling equations

Diffusion equation

Flux J:

$$J = -D\nabla c$$

$$c_t = \kappa \Delta c$$
$$\frac{dc}{dt} = \kappa \nabla^2 c$$

Neumann BC:

$$\nabla c \cdot n = g(t, c)$$

The binding process

First we look at the reversible chemical reaction

$$R + N \rightleftharpoons RN$$

with reaction rate k_1^* to the right and k_2^* to the left, being respectively the probability for the reactions to occur in their direction. We get 3 ODE's from this chemical reaction:

$$\begin{split} \frac{d[\mathbf{R}]}{dt} &= -k_1^*[\mathbf{R}][\mathbf{N}] + k_2^*[\mathbf{R}\mathbf{N}], \\ \frac{d[\mathbf{N}]}{dt} &= -k_1^*[\mathbf{R}][\mathbf{N}] + k_2^*[\mathbf{R}\mathbf{N}], \\ \frac{d[\mathbf{R}\mathbf{N}]}{dt} &= k_1^*[\mathbf{R}][\mathbf{N}] - k_2^*[\mathbf{R}\mathbf{N}], \end{split}$$

where [R], [N] and [RN] are the concentrations of the receptors, neurotransmitters and the bound product of them. We may consider [N][R] the probability of a neurotransmitter meeting an unoccupied receptor, and \bar{k}_1^* the probability of the binding reaction happening. Likewise for \bar{k}_2^* . P^R is the probability of a receptor being unoccupied, $(1-P^R)$ the probability that the neurotransmitter is attached to the receptor, leeds to the following simplification of the above ODE's:

$$\frac{d[N]}{dt} = -\bar{k}_1^*[N]P^R + \bar{k}_2^*(1 - P^R),$$

$$\frac{dP^R}{dt} = -\bar{k}_1^*[N]P^R + \bar{k}_2^*(1 - P^R).$$

If we assume that the receptors are not uniformly distributed, we need to introduce a $\gamma(x)$ to describe the density of receptors. At the boundary:

$$\frac{d[N]}{dt} = -\bar{k}_1^*[N]\gamma P^R + \bar{k}_2^*\gamma (1 - P^R),$$

$$\frac{dP^R}{dt} = -\bar{k}_1^*[N]\gamma P^R + \bar{k}_2^*\gamma (1 - P^R),$$

which are Neumann boundary conditions (inserting c for [N])

$$\kappa \nabla c \cdot n = -\bar{k}_1^* c \gamma P^R + \bar{k}_2^* \gamma (1 - P^R),$$

$$\frac{dP^R}{dt} = -\bar{k}_1^* c P^R + \bar{k}_2^* (1 - P^R),$$

Glia cells

$$T + N \rightleftharpoons TN \rightarrow N_{inactive} + T$$

Define k_3, k_4, k_5 as the reaction rates of first rightward, first leftward, second rightward equation.

Similarly to the binding process, we get the following sets of equations:

$$\kappa \nabla c \cdot n = -\bar{k}_3 c \gamma^T P^T + \bar{k}_4 \gamma^T (1 - P^T),$$

$$\frac{dP^T}{dt} = -\bar{k}_3 c P^T + (1 - P^T)(\bar{k}_4 + \bar{k}_5),$$

Combining these equations, we get

$$\kappa \nabla c \cdot n = -c(\bar{k}_1^* \gamma^R P^R + \bar{k}_3 \gamma^T P^T) + \bar{k}_2^* \gamma^R (1 - P^R) + \bar{k}_4 \gamma^T (1 - P^T),
\frac{dP^R}{dt} = -c\bar{k}_1^* P^R + \bar{k}_2^* (1 - P^R),
\frac{dP^T}{dt} = -c\bar{k}_3 P^T + (\bar{k}_4 + \bar{k}_5)(1 - P^T),$$

1D solution using solver

Modelling the equation in one dimension is done by considering the points a and b, and the line between them. In this case, a is on one side of the synaptic cleft, and b is on the other side. Due to this, the boundary conditions for a and b differ. For a, we have

$$\kappa \nabla c = -k_3 c P^T + k_4 (1 - P^T),$$

and for b we have

$$\kappa \nabla c = -k_1 c P^R + k_2 (1 - P^R).$$

The next step is to combine these boundary conditions with the modelling equation to form a matrix equation.

The final equation was

$$\hat{M}\dot{X}(t) = -\kappa \hat{K}X(t) - k_3\hat{Q}^a(X(t))X(t) + k_4(1 - X_{N+1}(t))\hat{d}^a + (k_4 + k_5)(1 - X_{N+1}(t))\hat{e}^a - k_1\hat{Q}^b(X(t))X(t) + k_2(1 - X_{N-2}(t))\hat{d}^b,$$

and was found by Jorg Henrik Holstad 1 Here, X is a vector of length N + 2, consisting of the concentrations at the nodes, as well as the probabilities

$$P_a^T$$
 and P_b^R : $X = \begin{bmatrix} C \\ P_a^T \\ P_b^R \end{bmatrix}$.

A plot using N = 9 internal nodes is shown below:

Torg Henrik Holstad. (2011). Modellering av Diffusjon av Nevrotransmittere i den Ekstracellelaere Vaesken. Retrieved from https://www.duo.uio.no/bitstream/handle/10852/10871/MasteroppgaveHenrikHolstad.pdf

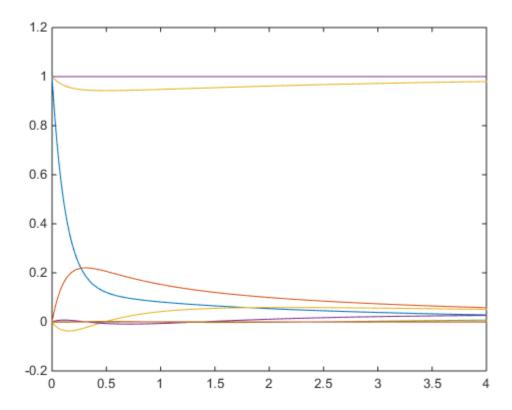


Figure 1: a = 0, b = 8, N = 9, $k_1 = k_2 = k_3 = k_4 = k_5 = 0.5$, $P_a^T(0) = P_b^R(0) = 1$