

# Part II: Mathematical Biology - Revision Notes

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## 1. Deterministic Systems

### 1.1. Single population models

#### 1.1.3. Populations with age structure

##### Notations

$n(a, t)$ : Number of individuals with age  $a$  at time  $t$ .

$N = \int_0^\infty n(a, t) da$ : Total population at time  $t$ .

$b(a)$ : Birth rate from individuals with age  $a$ .

$\mu(a)$ : Death rate from individuals with age  $a$ .

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$$n(a + \delta t, t + \delta t) = n(a, t) - \mu(a) \cdot \delta t \cdot n(a, t) + O(\delta t^2)$$

Also, by Taylor expansion,

$$n(a + \delta t, t + \delta t) = n(a, t) + \delta t \frac{\partial n}{\partial a} + \delta t \frac{\partial n}{\partial t} + O(\delta t^2)$$

Comparing and dividing by  $\delta t$ :

$$\frac{\partial n}{\partial a} + \frac{\partial n}{\partial t} = -\mu(a)n(a, t)$$

Boundary condition: Number of newborn babies satisfies

$$n(0, t) = \int_0^\infty b(a)n(a, t) da$$

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##### Normal node solutions

Set  $n(a, t) = r(a)e^{\gamma t} \implies r'(a) = -(\mu(a) + \gamma) r(a)$

$$r(a) = r(0)e^{-\gamma a} e^{\int_0^a \mu(s) ds}$$

$$n(a, t) = r(0)e^{\gamma(t-a)} e^{\int_0^a \mu(s) ds}$$

By boundary condition, we have

$$1 = \int_0^\infty b(a)e^{-\gamma a} e^{-\int_0^a \mu(s) ds} da := \phi(\gamma)$$

So if we can find a  $\gamma$  s.t.  $\phi(\gamma) = 1$ , then we have a valid normal node solution.

$\phi(\gamma)$  strictly decreasing  $\implies \phi(\gamma) = 1$  has a unique solution.

$$\phi(0) = \int_0^\infty b(a)e^{-\int_0^a \mu(s) ds} da$$

$\phi(0) > 1 \implies \gamma > 0 \implies$  growth

$\phi(0) < 1 \implies \gamma < 0 \implies$  decay

Biological interpretation of  $\phi(0)$ : mean number of offspring from one individual.

## 1.2. Discrete time

## 1.3. Multi-species models

### 1.3.1. Competition models

$$\dot{N}_1 = r_1 N_1 \left(1 - \frac{N_1}{K_1} - b_{12} \frac{N_2}{K_2}\right)$$

$$\dot{N}_2 = r_2 N_2 \left(1 - \frac{N_2}{K_2} - b_{21} \frac{N_1}{K_1}\right)$$

**Physical interpretation of each term:**

Each species along has a simple logistic dynamics with linear growth ( $r_1 N_1$  and  $r_2 N_2$ );

Negative quadratic terms mean each species has some stable equilibrium sizes ( $K_1$  and  $K_2$ ; the carrying capacity);

Terms with  $b_{12}$  and  $b_{21}$ : interactions between species. If the negative interaction terms are not too big, then two species will coexist at some stable equilibrium value; if too big, one species will win out.

**Rescaling:**  $M_1 = \frac{N_1}{K_1}$ ,  $M_2 = \frac{N_2}{K_2}$ ,  $\tau = r_1 t$ .

$$\frac{dM_1}{d\tau} = M_1(1 - M_1 - b_{12}M_2)$$

$$\frac{dM_2}{d\tau} = \rho M_2(1 - M_2 - b_{21}M_1)$$

where  $\rho = \frac{r_2}{r_1}$ .

### 1.3.2. The predator-prey models

$$\frac{dN}{dt} = N(a - bP)$$

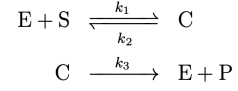
$$\frac{dP}{dt} = P(cN - d)$$

After rescaling,

$$\dot{u} = u(1 - v)$$

$$\dot{v} = -\alpha v(1 - u)$$

### 1.3.3. Chemical kinetics models



$$\dot{s} = -k_1 es + k_2 c$$

$$\dot{e} = -k_1 es + k_2 c + k_3 c$$

$$\dot{c} = +k_1 es - k_2 c - k_3 c$$

$$\dot{p} = +k_3 c$$

Initial conditions:  $s(0) = s_0$ ,  $e(0) = e_0$ ,  $c(0) = 0$ ,  $p(0) = 0$

Invariant quantities:  $\dot{e} + \dot{c} = 0$ ,  $\dot{s} + \dot{c} + \dot{p} = 0$ .  
 $\implies e + c = e_0$ ,  $s + c + p = s_0$ .

Hence, four equations reduce to two:

$$\dot{s} = -k_1(e_0 - c)s + k_2 c$$

$$\dot{c} = +k_1(e_0 - c)s - k_2 c - k_3 c$$

**Rescaling:**  $u = \frac{s}{s_0}$ ,  $v = \frac{c}{e_0}$  and suitable time rescaling gives:

$$u' = -u + (u + \mu - \lambda)v$$

$$\epsilon v' = u - (u + \mu)v$$

### 1.3.4. Epidemic models

$S$ : Susceptible,  $I$ : Infected,  $R$ : Recovered

$$\begin{aligned}\dot{S} &= -\beta IS \\ \dot{I} &= +\beta IS - \nu I \\ \dot{R} &= +\nu I\end{aligned}$$

$\beta$ =transmission rate,  $\nu$ =recovery rate

Total population size remain constant:  $N = S + I + R$ .  
Hence  
can determine  $R$  from  $R = N - S - I$ .

#### When can an epidemic happen?

$$\dot{I}(0) = [\beta S(0) - \nu]I(0)$$

**Assumptions:** Initially one or a few infected are introduced,  
i.e.  $I(0) > 0$ ; starting from nearly everyone susceptible, and  
only a very small number infected, i.e.  $S(0) \approx N$ .

Epidemic possible if and only if  $I(0) > 0$ , i.e.  $R_0 = \frac{\beta}{\nu}N > 1$ .

**Interpretation of  $R_0$  (reproduction ratio):**  $R$  has exponential distribution of time, with expected time to recovery  $\frac{1}{\nu}$ ,  
i.e. duration of an infection.  $\beta$  is the rate that one individual infects each susceptible. Hence,  $R_0$  is the mean number of infected from one infection.

#### Vaccination

Assume a proportion  $p$  of the population is vaccinated.  
Then  
 $S(0) = (1 - p)N$ ,

$$\frac{\beta}{\nu}S(0) = \frac{\beta}{\nu}(1 - p)N = (1 - p)R_0 > 1$$

for an epidemic to be possible.

#### Epidemic and final size

Consider  $\frac{dI}{dS} = \frac{\dot{I}}{\dot{S}} = -1 + \frac{N}{R_0} \frac{1}{S}$  (Note:  $\frac{dI}{dS} > -1$ )

Integrating wrt  $S$ :  $I = -S + \frac{N}{R_0} \log S + C$

Putting initial conditions:

$$I(t) - I(0) = -(S(t) - S(0)) + \frac{N}{R_0} \log \frac{S(t)}{S(0)} \quad (*)$$

where  $I(0) \ll N$ ,  $S(0) \approx N$ .

(Diagram: trajectories in  $(S, I)$  plane)

By differentiating  $(*)$ , we see that trajectories have a maximum number of infected as  $S = \frac{N}{R_0}$ .

**Define:**  $\sigma$  =proportion of the population that escape the epidemic and actually never get infected.

**Moral:**  $\sigma N$  is the left-most S-intercept in the diagram.

$$\begin{aligned}0 &= -[\sigma N - NB] + \frac{N}{R_0} \log\left(\frac{\sigma N}{N}\right) \\ \sigma - \frac{1}{R_0} \log \sigma &= 1\end{aligned}$$

(Diagram for  $\sigma$  as a function of  $R_0$ ).

#### Extended epidemic model

In long term, consider natural births and deaths: let  $\mu$  be the constant birth/death rate per capita.

$$\dot{S} = -\beta IS - \mu S + \mu N$$

$$\dot{I} = \beta IS - \nu I - \mu I$$

$$\dot{R} = \nu I - \mu R$$

(normally ignore the equation for  $R$ ).

New reproduction ratio:  $R_0 = \frac{\beta}{\nu + \mu} N$ .

Consider  $\nu \gg \mu$ , i.e. the recovery rate is far higher than the natural death rate, i.e. the infectious period is much shorter than host lifetime.

$\exists$  a FP with disease present:  $S^* = \frac{N}{R_0}$ ,  $I^* = \frac{\mu}{\beta}(R_0 - 1)$ .

This is a stable focus with  $\omega \approx \sqrt{\mu\nu(R_0 - 1)}$  and period  $T \approx \frac{2\pi}{\sqrt{\mu\nu(R_0 - 1)}}$ .

$(\mu\nu)^{-1/2}$ : geometric mean of host lifetime and disease infectious period.

### 1.3.5. Excitable systems

## 2. Stochastic Systems

Write  $P(n, t)$  to represent the probability of the population size being  $n$  at time  $t$ . For simplicity, we write  $P(n, t) = p_n$ .

$$\phi(s, t) = \sum_{n=0}^{\infty} s^n p_n = \langle s^n \rangle$$

$$\phi(1, t) = 1$$

$\phi(0, t) = p_0$  = probability that the population has die out.

$$\phi(s, 0) = 1$$


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### 2.1. Discrete population sizes

**Single populations:** Constant probability rate  $\lambda$  of adding one individual to the population

**Master equation:**  $\dot{p}_n = \lambda(p_{n-1} - p_n)$

Try generating function:  $\phi(s, t) = \sum_{n=0}^{\infty} s^n p_n = \langle s^n \rangle$

$$\dot{\phi} = (s - 1)\lambda\phi$$

$$\phi = A(s)e^{(s-1)\lambda t}$$

$$\phi(s, 0) = 1 \Rightarrow \phi(s, t) = e^{(s-1)\lambda t}$$

$$\mu = \langle N \rangle = \frac{\partial \phi}{\partial s} \Big|_{s=1} = \lambda t$$

$$\langle N(N-1) \rangle = \frac{\partial^2 \phi}{\partial s^2} \Big|_{s=1} = (\lambda t)^2$$

$$var(N) = \lambda t$$


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**Import and death model:** Same import model as before with a per capita probability death rate  $\beta$  (so the total death rate is  $\beta n$ ).

**Master equation:**  $\dot{p}_n = \lambda(p_{n-1} - p_n) + \beta[(n+1)p_{n+1} - np_n]$

$$\phi_t = (s - 1)[\lambda\phi - \beta\phi_s]$$

Try  $\phi = e^{(s-1)f(t)} \Rightarrow \phi(s, t) = \exp[\frac{\lambda}{\beta}(s-1)(1 - e^{-\beta t})]$ .

$$\langle N \rangle = \frac{\lambda}{\beta}(1 - e^{-\beta t})$$

$$var(N) = \frac{\lambda}{\beta}(1 - e^{-\beta t})$$


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**Multiple population model**

## 2.2. Continuous population sizes

### 2.2.1. Fokker-Planck for a single variable

$W(n, r)$  represents the jump rate from  $n$  to  $n + r$ . Use  $x$  to represent continuous population size instead of  $n$ .

**Master equation:**

$$\frac{\partial}{\partial t} P(n, t) = \sum_r [W(n-r, r)P(n-r, t) - W(n, r)P(n, t)]$$

**Fokker-Planck Equation (FPE):**

$$\frac{\partial P}{\partial t} = -\frac{\partial}{\partial x}(AP) + \frac{1}{2}\frac{\partial^2}{\partial x^2}(BP)$$

*Derivation:* Do Taylor expansion to the master equation and define  $A(x) = \sum_r rW$  and  $B(x) = \sum_r r^2W$

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How expectation of a general function  $f(x)$  evolves in time:

$$\frac{d}{dt} \langle f(x) \rangle = \langle Af' \rangle + \frac{1}{2} \langle Bf'' \rangle$$

*Derivation:* By definition of expectation, then substitute Fokker-Planck equation, and lastly integration by parts with boundary conditions  $P$  and  $\frac{\partial P}{\partial x}$  tends to 0 as  $|x| \rightarrow \infty$ .

Time evolution of mean:  $\langle x \rangle' = \langle A \rangle$  (time derivative)

Time evolution of variance:  $var(x)' = \langle B \rangle + 2cov(A, x)$

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### 2.2.2. Multivariate Fokker-Planck

**General master equation:**

$$\frac{\partial}{\partial t} P(\mathbf{x}, t) = \sum_{\mathbf{r}} [W(\mathbf{x} - \mathbf{r}, \mathbf{r})P(\mathbf{x} - \mathbf{r}, t) - W(\mathbf{x}, \mathbf{r})P(\mathbf{x}, t)]$$

**Multidimensional Fokker-Planck equation:**

$$\frac{\partial P}{\partial t} = -\frac{\partial}{\partial x_i}(A_i P) + \frac{1}{2}\frac{\partial^2}{\partial x_i \partial x_j}(B_{ij} P)$$

Time evolution of  $\mathbf{x}$ :  $\langle \mathbf{x} \rangle' = \langle \mathbf{A} \rangle$

Time evolution of covariance:

$$C'_{mn} = cov(A_m, x_n) + cov(A_n, x_m) + \langle B_{mn} \rangle$$

where  $C_{mn} = cov(x_m, x_n) = \langle x_m x_n \rangle - \langle x_m \rangle \langle x_n \rangle$

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**Behaviour at steady state**

**Lyapunov equation:**

$$\mathbf{aC} + \mathbf{Ca}^T + \mathbf{b} = 0$$

$$a_{ik}C_{kj} + a_{jk}C_{ki} + b_{ij} = 0$$


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### 3. Systems with spatial structure

$C(\mathbf{x}, t)$  is some quantity of interest.

$$\frac{d}{dt} \int_V C(\mathbf{x}, t) dV = \int_V F(\mathbf{x}, t) dV - \int_V \mathbf{J} \cdot \mathbf{n} dS$$

By divergence theorem, we get  $\frac{\partial C}{\partial t} = F - \nabla \cdot \mathbf{J}$

There are two types of flux:

1. Advection or active motion:  $\mathbf{J} = \mathbf{u}C$  where the stuff moves with velocity  $\mathbf{u}$ .
2. Diffusion:  $\mathbf{J} = -D\nabla C$

So transport equation becomes:

$$\frac{\partial C}{\partial t} + \nabla \cdot (\mathbf{u}C - D\nabla C) = F(\mathbf{x}, t)$$

#### 3.1. Diffusion and growth

##### 3.1.1. Linear diffusion in finite domain

**Basic setup:**  $D$  constant,  $F = 0$ ,  $\mathbf{u} = 0$ , one-dimension. So the transport equation is:  $\frac{\partial C}{\partial t} = D \frac{\partial^2 C}{\partial x^2}$ ; In finite linear domain of length  $L$ :  $x \in [0, L]$ ; Boundary condition:  $C(0, t) = C_0$ ,  $C(L, t) = C_1$ .

**Steady state solution:**

$$0 = D \frac{\partial^2 C}{\partial x^2}$$

Then  $C$  is just linear in  $x$ :

$$C(x, t) = C^*(x) = C_0 + (C_1 - C_0)x/L$$

**General solution:**

$$C(x, t) = C^*(x) + \hat{C}(x, t)$$

By linearity:

$$\frac{\partial \hat{C}}{\partial t} = D \frac{\partial^2 \hat{C}}{\partial x^2}$$

with boundary condition  $\hat{C}(0, t) = \hat{C}(L, t) = 0$ .

**Separation of variables:**

$$\hat{C}(x, t) = F(x)G(t)$$

$$\frac{G'(t)}{G(t)} = D \frac{F''(x)}{F(x)}$$

Boundary condition  $F(0) = F(L) = 0$  gives:

$$F(x) = \sin\left(\frac{n\pi x}{L}\right), \quad n = 1, 2, \dots$$

$$G(t) = e^{-\lambda_n t} \quad \text{with} \quad \lambda_n = D \frac{n^2 \pi^2}{L^2}$$

$$\hat{C}(x, t) = \sum_{n=1}^{\infty} a_n e^{-\lambda_n t} \sin\left(\frac{n\pi x}{L}\right)$$

$$C(x, t) = C_0 + (C_1 - C_0)x/L + \sum_{n=1}^{\infty} a_n e^{-\lambda_n t} \sin\left(\frac{n\pi x}{L}\right)$$

**Add linear growth to finite domain:**  $\frac{\partial C}{\partial t} = D \frac{\partial^2 C}{\partial x^2} + \lambda C$

Set  $C(x, t) = e^{\lambda t} \tilde{C}(x, t)$ , then

$$\frac{\partial \tilde{C}}{\partial t} = D \frac{\partial^2 \tilde{C}}{\partial x^2}$$

##### 3.1.2. Linear diffusion in an infinite domain

$$\frac{\partial C}{\partial t} = D \frac{\partial^2 C}{\partial x^2}$$

with boundary condition that  $C$  and  $\frac{\partial C}{\partial x}$  tends to 0 sufficiently fast as  $|x| \rightarrow \infty$ .

Total amount of stuff:  $M = \int_{-\infty}^{\infty} C(x, t) dx$ .

Check that  $M$  is constant:  $\frac{dM}{dt} = \frac{d}{dt} \int_{-\infty}^{\infty} C(x, t) dx = \int_{-\infty}^{\infty} \frac{\partial C}{\partial t} dx = \int_{-\infty}^{\infty} D \frac{\partial^2 C}{\partial x^2} dx = [D \frac{\partial C}{\partial x}]_{-\infty}^{\infty} = 0$

**Approach 1: Similarity solution**

Seek a solution of the form  $C(x, t) = \eta f(\xi)$  with  $\xi = \frac{x}{\sqrt{Dt}}$  and  $\eta = \frac{M}{\sqrt{Dt}}$

Then get  $f = Ae^{-\frac{1}{4}\xi^2}$ .

$$\int_{-\infty}^{\infty} f d\xi = 1 \implies A = (4\pi)^{-\frac{1}{2}}$$

**Approach 2: Sort powers of  $t$**

Set  $C(x, t) = t^\alpha G(\xi)$  with  $\xi = \frac{x}{t^\beta}$ .

Substitute this into the diffusion equation to get  $\beta = \frac{1}{2}$ .

$M = \int_{-\infty}^{\infty} C dx$  to get  $\alpha = -\frac{1}{2}$ .

**Add simple growth to linear diffusion in infinite domain**

$$\frac{\partial C}{\partial t} = D \frac{\partial^2 C}{\partial x^2} + \lambda C$$

Set  $C(x, t) = e^{\lambda t} \tilde{C}(x, t) \implies \frac{\partial \tilde{C}}{\partial t} = D \frac{\partial^2 \tilde{C}}{\partial x^2}$

##### 3.1.3. Non-linear diffusion

$$D = kC, \quad \frac{\partial C}{\partial t} = \frac{\partial}{\partial x} \left( D \frac{\partial C}{\partial x} \right) = k \frac{\partial}{\partial x} \left( C \frac{\partial C}{\partial x} \right)$$

$$\xi = \frac{x}{(Mkt)^{1/3}}, \quad \eta = \frac{M}{(Mkt)^{1/3}}$$

$$M = \int_{-\infty}^{\infty} C dx \implies \int_{-\infty}^{\infty} F d\xi = 1$$

Substitute new variable gets  $-\frac{\eta}{3t}(\xi F)' = \frac{\eta}{t}(FF')' \implies FF' + \frac{1}{3}\xi F = 0$  by noting that LHS goes to zero as  $x \rightarrow \infty$ .

$$F(\xi) = \begin{cases} A - \frac{1}{6}\xi^2 & \text{for } |\xi| < \sqrt{6A} \\ 0 & \text{otherwise} \end{cases}$$

Solve  $A$  by integration.

### 3.2. Travelling waves in reaction-diffusion systems