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# APPENDIX A

## APPENDIX

# MODELING OF POPULATION WITH STAGE-STRUCTURE

Recent interest in a mathematical description of the age-structure of growing cell populations has centered on the use of an equation proposed by von-Foerster. This equation is a first order partial differential equation for a function n(t,a) defined so that n(t,a) is the number of organisms at time t whose ages lie in the interval (a, a + da). Note that most details in this part are collected and adapted from [2] and [18].

The age a that appears as an independent variable in n(t, a) is always interpreted as the chronological age of the organism. In consequence, the properties of two organisms belonging to a single age-group and subjected to identical environments are identical. Thus, the important feature of biological variability is omitted from the description of the age-structure of a single cohort.

The derivative of this kind of models is from the von-Foerster or the McKendrick equation. Let n(t,a) be the density of an age-structure population at time t of age a. Consider the partial differential equation

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

for t > a. Integrate along a characteristic curve given by  $t = a + \xi$ , and define

$$n_{\varepsilon}(a) = n(a + \xi, a).$$

$$\frac{dn_{\xi}}{da} = \left[\frac{\partial n}{\partial t} + \frac{\partial n}{\partial a}\right]_{t=a+\xi}$$
$$= -\mu(a)n(a+\xi,a)$$
$$= -\mu(a)n_{\xi}(a).$$

Hence,

$$n_{\xi}(a) = n_{\xi}(0)e^{-\int_{0}^{a}\mu(s)ds}$$

or

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$$n(a + \xi, a) = n(\xi, 0)e^{-\int_0^a \mu(s)ds}.$$

Putting  $a = \tau$  and  $\xi = t - \tau$  gives

$$n(t,\tau) = n(t-\tau,0)e^{-\int_0^u \mu(s)ds},$$

provided  $t > \tau$ .

Suppose that the total number of adults is N(t), where

$$N(t) = \int_{\tau}^{\infty} n(t,a) da.$$

Then

$$\frac{dN}{dt} = \int_{\tau}^{\infty} \frac{\partial n}{\partial t} da$$
$$= \int_{\tau}^{\infty} \left( -\frac{\partial n}{\partial a} - \mu(a)n(t,a) \right) da.$$

We can see that

$$\frac{dN}{dt} = n(t,\tau) - n(t,\infty) - \mu_M N(t), \quad \text{if } \mu(a) = \mu_M \text{ for all } a \ge \tau \,,$$

$$\frac{dN}{dt} = n(t - \tau, 0)e^{-\int_0^\tau \mu(s)ds} - \mu_M N(t), \text{ if } n(t, \infty) = 0.$$

But n(t,0) is the birth rate. If this is a function of the total number of adults, then n(t,0) = B(N(t)). So that

$$\frac{dN}{dt} = e^{-\int_0^t \mu(s)ds} B(N(t-\tau)) - \mu_M N(t).$$

We can see that term of age-structure  $e^{-\int_0^\tau \mu(s)ds}$  is absorbed. In my thesis, we assume that  $\mu(s)$  is a constant. The equation become

$$\frac{dN}{dt} = e^{-\delta\tau} B(N(t-\tau)) - \mu_M N(t),$$

where  $\mu(s) = \delta$ . Then, the above equation is similar to the selected model for analysis in Chapter IV:

$$N'(t) = -\gamma N(t) + \beta e^{-\delta \tau} f(N(t-\tau))N(t-\tau),$$

where  $\mu_M = \gamma$ . Note that the equation above is the model which we selected to study for the case of age-structure population models.

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