

**SIMULATION AND MODELLING**

**CONTINUOUS TIME STOCHASTIC COMPARTMENTAL MODELS  
 OF DISCRETE POPULATIONS**

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PRELIMINARIES

Let  $[X(t); t > 0]$  denote continuous time,  $n$ -state semi-Markov process with stochastic transition matrix  $P = (p_{ij})$ , state residence time distribution function matrix  $W = [w_{ij}(z)]$ , and stochastic inter-val transition probability matrix  $F = [f_{ij}(t)]$  ( $i, j = 1, \dots, n$ ).  $X(t)$  is the state of the process at its most recent change of state and element  $f_{ij}(t)$  of  $F$  is the conditional probability that  $X(t) = j$  at time  $t$ , given that the initial state  $X(0+) = i$ . Elements of  $F$  are related to elements of  $P$  and  $W$  by a Markov renewal equation of the Volterra type whose solution can be expressed by conditioning on the number of changes of state of the process prior to time  $t$ :

$$f_{ij}(t) = \sum_{l=0}^{\infty} \Pr[X(t) = j / X(0+) = i, l \text{ changes of state in } (0, t)] \times$$

$$\times \Pr[l \text{ changes of state in } (0, t) / X(0+) = i] =$$

$$\delta_{ij} \cdot h_i(t) + p_{ij} \int_0^t w'_{ij}(z) \cdot h_j(t-z) dz +$$

$$+ \sum_{l=2}^{\infty} \left[ \sum_{q_1=1}^m p_{iq_1} \sum_{q_2=1}^{m_1} p_{q_1 q_2} \dots \sum_{q_{k-2}=1}^n p_{q_{k-2} q_{k-1}} \sum_{q_{k-1}=1}^n p_{q_{k-1} q_k} \right.$$

$$\left. - 1 \cdot p_{q_{k-1} q_1} \right] \times$$

$$\times \left[ \int_0^t (w_{iq_1} * w_{q_1 q_2} * \dots * w_{q_{k-2} q_{k-1}} * w_{q_{k-1} q_1})'(z) \cdot h_j(t-z) dz \right]$$

( $i, j = 1, \dots, n$ ) (eqn. 1)

where:

$$q_1 = j;$$

$$\int_0^t (w_{iq_1} * \dots * w_{q_{k-1} q_1})'(z) \cdot h_j(t-z) dz,$$

an  $l$ -fold convolution density convolved with  $h_j(t)$ , is multiplied by the probability  $(p_{iq_1} \dots p_{q_{k-1} q_1})$  that the  $l$ -step sequence of changes of state  $(i, q_1, \dots, q_{k-1}, q_1 = j)$  occurs:

$$h_i(t) = 1 - \sum_{k=1}^n p_{ik} \cdot w_{ik}(t);$$

state  $j$  is assumed to be reachable from state  $i$  so that there is at least one 1-step sequence with positive probability.

When  $P$  is upper or lower diagonal the infinite sum on the right hand side of eqn. (1) terminates for  $l > n+1$ .

Let  $C$  denote a discrete population in which the behavioral states of individuals are in one-to-one correspondence with the states of  $[X(t)]$ . Let  $S = (P, W, F)$  denote the system governing movement of individuals among behavioral states once they enter  $S$  from external sources. The conditional probability that an individual is in state  $j$  at time  $t > 0$ , given that it initially entered  $S$  at time  $z$  ( $0 < z < t$ ) in state  $i$  is  $f_{ij}(t-z)$ . Once inside  $S$  individuals are assumed to behave independently unless otherwise specified.

Subsets of states, aggregated into  $K$  non-overlapping and exhaustive subsets  $G_1, \dots, G_K$  are called compartments ( $K = 2, \dots, n-1$ ). The probability  $f_{iG_k}(t-z)$  that an individual entering state  $i$  at time  $z > 0$  is in compartment  $G_k$  at time  $t > z$  is:

$$f_{iG_k}(t-z) = \sum_{\text{state } j \text{ in } G_k} f_{ij}(t-z) \quad (\text{eqn. 2})$$

Let  $Y_{ij}(t)$  ( $i, j = 1, \dots, n$ ) be random variables denoting numbers of individuals in states  $1, \dots, n$  at time  $t > 0$  whose initial entry into  $S$  is through state  $i$ .

The number  $Y_{iG_k}(t)$  of individuals in compartment  $G_k$  at time  $t$  is:

$$Y_{iG_K}(t) = \text{state}_{\text{in } G_K} \sum_j Y_{ij}(t) \quad (\text{eqn. 3})$$

The mean and variance of  $Y_{iG_K}(t)$  is determined for different assumptions about processes of arrivals to S from external sources.

INDIVIDUAL POISSON ARRIVALS

Assume a Poisson stream of individual arrivals to S. Given that  $N_i$  arrivals occur in  $(0,t)$  to initial state i the joint p.f. of numbers in states  $1, \dots, n$  at time t is multinomial with parameters  $N_i, f_{i1}(t), \dots, f_{in}(t)$ . Multiplying the joint p.f. by the Poisson probability of  $N_i$  arrivals in  $(0,t)$  conditional on the arrival times of the first  $N_i$  arrivals being distributed as the order statistics of  $N_i$  independent samples from the d.f. on  $(0,t)$  having density  $\lambda_i(z) / \int_0^t \lambda_i(z) dz$  ( $0 < z < t$ ) the resulting joint p.f. is a product of n independent Poisson probabilities that  $Y_{i1}, \dots, Y_{in}$  individuals are in states  $1, \dots, n$  at time t:

$$P\{Y_{i1}(t)=y_{i1}, \dots, Y_{in}(t)=y_{in}, N_i \text{ arrivals}\} = \frac{m}{\prod_{j=1}^n y_{ij}!} \left( \lambda_i(z) \cdot f_{ij}(t-z) dz \right)^{Y_{ij}} \cdot e^{-\int_0^t \lambda_i(z) \cdot f_{ij}(t-z) dz} \quad (\text{eqn. 4})$$

$$(y_{i1} + \dots + y_{in} = N_i)$$

As shown by eqn. (4) the  $Y_{ij}(t)$ 's are mutually independent Poisson distributed r.v.'s Moreover:

i) the arrival stream to state j is Poisson distributed with intensity  $a_j(t)$  which satisfies the integral equation:

$$\int_0^t a_j(z) \cdot \sum_{k=1}^m p_{jk} \cdot [1 - w_{jk}(t-z)] dz = \int_0^t \lambda_i(z) \cdot f_{ij}(t-z) dz; \quad (j=1, \dots, n)$$

ii) the expectation of  $Y_{ij}(t)$  is:

$$E\{Y_{ij}(t)\} = \int_0^t \lambda_i(z) \cdot f_{ij}(t-z) dz \quad (\text{eqn. 5}) \quad (j=1, \dots, n)$$

Equations (5) when combined with equations (1) provide the basis for constructing families of regression models of inputs to the system S as well as inputs and outputs among states within S, from which parameters can be estimated. Maximum likelihood estimates of parameters can be obtained from eqn. (4).

The r.v.'s  $Y_{iG_k}(t)$  are independent and Poisson distributed with Poisson arrival intensities  $a_{iG_k}(t)$  and expectations:

$$E\{Y_{iG_k}(t)\} = \int_0^t \lambda_i(z) \cdot f_{iG_k}(t-z) dz = \int_0^t \lambda_i(z) \cdot dz \cdot \sum_{\text{state in } G_k} f_{ij}(t-z) \quad (\text{eqn. 6}) \quad (k=1, \dots, K)$$

BATCH POISSON ARRIVALS

Individuals arrive at initial state i in batches, at random (Poisson arrivals) where the mean and variance of the i.i. d. batch sizes are  $m_i$  and  $v_i$  respectively. The intensity of arrivals is  $\lambda_i(t)$ . The marginal d.f. of  $Y_{ij}(t)$  in this case is not Poisson unless  $v_i=0$  and  $m_i=1$ . The mean and variance of  $Y_{ij}(t)$  are:

$$E\{Y_{ij}(t)\} = m_i \cdot \int_0^t \lambda_i(z) \cdot f_{ij}(t-z) dz \quad (j=1, \dots, n) \quad (\text{eqn. 7})$$

and:

$$\text{Var}\{Y_{ij}(t)\} = m_i \cdot \int_0^t \lambda_i(z) \cdot f_{ij}(t-z) \cdot [1 - f_{ij}(t-z)] dz + (m_i^2 + v_i) \cdot \int_0^t \lambda_i(z) \cdot [f_{ij}(t-z)]^2 dz + v_i \cdot \left[ \int_0^t \lambda_i(z) \cdot f_{ij}(t-z) dz \right]^2 \quad (\text{eqn. 8}) \quad (j=1, \dots, n)$$

Equation (8) is demonstrated by first decomposing  $Y_{ij}(t)$  into the random sum of "clusters" of sizes  $1, 2, \dots, B$ :

$$Y_{ij}(t) = 1 \cdot D_{i1}(t) + 2 \cdot D_{i2}(t) + \dots + B \cdot D_{iB}(t) \quad (\text{eqn. 9})$$

where:

$D_{ik}(t)$  is a Poisson distributed r.v. with expectation:

$$E[D_{ik}(t)] = \int_0^t \lambda_i(z) \binom{B}{k} \cdot [f_{ij}(t-z)]^k \cdot [1-f_{ij}(t-z)]^{B-k} dz \quad (k=1,2,\dots,B)$$

For a batch of given size B arriving at initial state i at time z a cluster of k-out-of-B of the arriving individuals will be in state j (t-z) time units later with binomial probability:

$$\binom{B}{j} \cdot [f_{ij}(t-z)]^j \cdot [1-f_{ij}(t-z)]^{B-j}$$

Combining the relation:

$$\text{Var}[Y_{ij}(t)] = E[\text{Var}(Y_{ij}(t)/B)] + \text{Var}[E(Y_{ij}(t)/B)]$$

with eqn.(9), eqn (8) is obtained. The distribution of the number of clusters in state j at time t without regard to the cluster size for fixed size B of arriving batches is Poisson distributed with expectation:

$$\int_0^t \lambda_i(z) \cdot dz \cdot \sum_{r=0}^B \binom{B}{r} \cdot [f_{ij}(t-z)]^r \cdot [1-f_{ij}(t-z)]^{B-r}$$

Members of a given cluster have not necessarily been in residence in state j for the same length of time, however.

The mean and variance of the number of individuals in compartment  $G_k$  at time t are obtained by substituting  $f_{iG_k}(t-z)$

for  $f_{ij}(t-z)$  in equations 7 and 8.

ARBITRARY BUT FIXED INTERVALS BETWEEN ARRIVALS OF BATCHES

Batches of individuals, where batch sizes are i.i.d. random variables with mean  $m_1$  and variance  $v_1$  arrive at initial state i at arbitrary but fixed times  $t_1, t_2, \dots$ . For a batch arriving at time  $t_u$  and of conditional size  $B_u$  the joint p.f. of numbers in states  $1, 2, \dots, n$  at time  $t > t_u$  is multinomial

with parameters  $B_u, f_{i1}(t-t_u), \dots, f_{in}(t-t_u)$ .

The marginal p.f. of the number  $Y_{ij}(t)$  of individuals in state j at time t due only to the arriving batch at initial state i at time  $t_u$  of random size  $B_u$  has a compound form with mean and variance:

$$E[Y_{ij}(t)/t_u] = m_i \cdot f_{ij}(t-t_u) \quad \text{eqn. (10)}$$

and:

$$\text{Var}[Y_{ij}(t)/t_u] = m_i \cdot f_{ij}(t-t_u) \cdot [1-f_{ij}(t-t_u)] + v_i \cdot [f_{ij}(t-t_u)]^2 \quad \text{eqn. (11)}$$

If batch size is a fixed constant  $m_1$  then equation 11 is modified by setting  $v_i$  equal to zero.

The mean and variance of  $Y_{iG_k}(t)$  are obtained by substituting  $f_{iG_k}(t-z)$  for  $f_{ij}(t-z)$  into equations 10 and 11.

The mean and variance of the marginal d.f. of the number of individuals in compartment  $G_k$  due to all arriving batches at times  $0 < t_1, \dots, t_u < t$  is, assuming independence of all movements of individuals entering upon S:

$$E[Y_{iG_k}(t)/t_1, \dots, t_u] = \sum_{r=1}^u E[Y_{iG_k}(t)/t_r] \quad (k=1, \dots, K) \quad \text{eqn. (12)}$$

and:

$$\text{Var}[Y_{iG_k}(t)/t_1, \dots, t_u] = \sum_{r=1}^u \text{Var}[Y_{iG_k}(t)/t_r] \quad (k=1, \dots, K) \quad \text{eqn. (13)}$$

As with equations 5, equations 6, 7, and 12 can be used as the basis of constructing regression estimates of parameters of the system S.

SUB AND SUPER SYSTEMS OF S

The system  $S=(P,W,F)$  may be decomposable into subsystems  $S_1, \dots, S_K$  identified

with compartments  $G_1, \dots, G_K$  or it may itself be a subsystem of a larger super-system of states in which  $S$  is identified with a compartment  $G_S$ . In either case it is important to maintain stochasticity of the state transition matrices and the interval transition matrices corresponding to each subcollection of states that are to be identified with a subsystem.

Let  $G_1, \dots, G_K$  denote a collection of compartments of  $S$  and arrange the transition matrix  $P$  into the form:

$$P = (P_{G_i G_j}) \quad (i, j = 1, 2, \dots, K)$$

where:

the submatrix  $P_{G_i G_j}$  has row and column dimension equal, respectively, to the number of states in compartments  $G_i$  and  $G_j$ .

Main diagonal submatrices contain state transition probabilities governing movements of individuals among states within compartment ( $G_i$  ( $i=1, \dots, K$ )). Either advance or return to states in  $G_j$  from states in  $G_i$  is restricted by the number and locations of positive entries in off-diagonal submatrices  $P_{G_i G_j}$ . If no positive entries occur in  $P_{G_i G_j}$  for all indices  $i$  and  $j$  then the system  $S$  consists of  $K$  independent subsystems. Each submatrix  $P_{G_i G_j}$  is stochastic as well as the submatrix  $F_{G_i G_i}$  of interval transition probabilities describing the time rate of movement of individuals among states of compartment  $G_i$ . Equations 1-13 are valid for each subsystem  $S_1, \dots, S_K$  in this case.

If compartments  $G_i$  and  $G_j$  are linked by positive entries in off-diagonal

submatrix  $P_{G_i G_j}$  then submatrix  $P_{G_i G_i}$  is not stochastic and movements of individuals within compartment  $G_i$  cannot be analyzed independent of other states of  $S$ . By joining  $G_i$  to one additional absorbing state accounting for movements of individuals out of  $G_i$  and assigning transition probabilities into the appended absorbing state equal to one minus the row sums  $P_{G_i G_i}$  for each row in the submatrix, the compartment  $G_i$  can be analyzed in either one of two ways:

- i) as a subsystem in which arrivals are from other states of  $S$  or
- ii) as a subsystem in which arrivals are assumed to occur without reference to prior movements in  $S$ .

If the system  $S$  is composed of states which are themselves a compartment of a supersystem, then  $S$  functions independently of other compartments or else  $S$  contains an absorbing state as described above so that movements of individuals within  $S$  can be analyzed independently of their movements within other compartments.