## 1. Introduction and Examples

Models in Science and their relation to Mathematical Models
1.1 What is a stochastic process?
1.2 Topics to be discussed
1.3 Examples

- Waiting for a bus
- Play the winner rule
- Prevalence and Incidence Models
- Clinical Trials
- Early Detection of Disease
- Planning Early Detection Trials
- Cell Kinetics
- Models for Prevention Trials


## Mathematical Models

- A model can be used to vary the inputs and make predictions of the outputs.
- Mathematical models can be used to guide experiments.
- Nearly all mathematical models are simplifications of the phenomenon being observed. However a useful model captures the important features of the phenomenon.


## Models in Science

MODELS: Science has many different kinds of models.
Animal Models: To study human diseases.
Physical Models: To demonstrate the apparent truth of a principle.
Mathematical Models: To gain insight and understanding of a phenomenon which is being observed.

$$
\text { inputs } \Longrightarrow \text { MODEL } \Longrightarrow \text { Outputs }
$$

A Probability Model is a mathematical model which incorporates the probability aspects of the observed phenomenon. If the observed process has a time parameter (or something equivalent) we refer to the probability model as a stochastic process.

The model can be used to:

- Understand the phenomenon better
- Make predicitons of how outcome change with inputs


### 1.1 What is a Stochastic Process ?

A Stochastic Process is a collection of random variables $\{X(t), t \in T\}$ indexed by the quantity $t$ taking on values in the set $T$. Generally $t$ may be time.
$X(t)$ takes on values in a set $S$ called State Space.

## Two Major Cases (Time Domain)

Discrete Time: $T=\{0,1,2, \ldots\}$, then the process is written as $\left\{X_{n}, n \geq 0\right\}$
Continuous Time: $T=[0, \infty)$ we represent the process by $\{X(t), t \geq 0\}$.

## Two Major Cases (State Space)

Discrete or Countable: $S: 0,1,2, \ldots$
Continuous: $S:(-\infty, \infty)$
If $\{X(t), t \in T\}$ is an observed outcome of a stochastic process it is referred to as a sample path.
The set of all sample paths is the sample space.

## Classification of Processes

Time
Discrete Continuous

| State | Discrete | - | - |
| :--- | :--- | :--- | :--- |
| Space | Continuous | - | - |

## Goals of Analysis

- Find distribution of $X(t)$
- Find low order moments: mean, variance
- Find behavior at $t \rightarrow 0$ or $t \rightarrow \infty$
- Find special characteristics of process; e.g. First passage times (time to reach a state for first time).


### 1.2 Topics to be Discussed

- Review of Laplace Transforms
- Relations between Incidence and Prevalence of a disease
- Simple Time Dependent Models

Poisson Processes
Birth and Death Processes

- Markov Chains
- Renewal Processes
- Semi-Markov Process

Relation to Markov Chains and Processes

Topics to be Discussed (Cont.)

- Application to Clinical Trials
- Applications to Screening Problems
- Applications to Planning Disease Prevention Trials
- Miscellaneous Applications

Cell Kinetics
Family History Problems
1.3 Examples

## Waiting for a BUS

Problem: Why is it when one waits for a bus it always seems to take an unusually long time?

Hypothetical situation
Observed time between bus arrivals


5 Intervals in 60 minutes
MTBA $=$ Mean time between arrivals
$=\frac{\text { Total observed time }}{\text { No. of intervals }}=\frac{60}{5}=12$ mintes

Suppose a friend decides to catch a bus sometime during the observational period. How long does she have to wait for the bus to arrive (on the average)? What is her expected mean waiting time if MTBA $=12$ minutes?


MTBA $=12$ minutes



5 intervals

Equivilant to placing 5 points at random on clock; i.e.


Flip point 5 times

## Informal Analysis

Suppose the arrival point is placed on the clock initially


## Conclusion:

Placing a point on a clock is the same as spinning a pointer once. But we have 5 spins and one placed point. This is equivilant to 6 spins and results in 6 intervals. Hence average time between points is

$$
\frac{60 \text { minutes }}{6}=10 \text { minutes }
$$

Therefore average waiting time $=10$ minutes.

Reason for the unexpected result is that the arrival of friend at the bus stop is equivalent to choosing an inter-arrival interval. The probability of choosing an interval is proportional to the length of the interval.

## b. Play the Winner Rule (Two State Markov Chain)

Two treatments are to be evaluated. Outcome of individual observation is Success or Failure.

$$
\begin{aligned}
p_{i} & =P\{\text { Success on Treatment } i\} \quad q_{i}=1-p_{i}, \quad i=0,1 \\
X_{n} & = \begin{cases}1 & \text { if } \mathrm{n}^{t h} \text { patient is on } 1 \\
0 & \text { if } \mathrm{n}^{t h} \text { patient is on } 0\end{cases}
\end{aligned}
$$

PWR: If $(n-1)^{s t}$ patient is on $i$ and outcome is Success $\Rightarrow \operatorname{assign} n^{t h}$ patient to $i$. If outcome is Failure $\Rightarrow$ assign $n^{t h}$ patient to other treatment. Choose $n=1$ patient to be on treatment $i$ with prob. $=1 / 2$.

$$
\begin{aligned}
& P\left\{X_{1}=1\right\}=1 / 2 \\
& P\left\{X_{n}=1 \mid X_{n-1}=1\right\}=p_{1}, \quad P\left\{X_{n}=1 \mid X_{n-1}=0\right\}=q_{0} \\
S_{n}= & X_{1}+X_{2}+\ldots+X_{n}=\text { Number on treatment } 1 \text { for first } n \text { patients. }
\end{aligned}
$$

$\underline{\text { Sample Path: }} x_{1}, x_{2}, \ldots, x_{n}$
Joint Distribution: $P\left\{X_{1}=x_{1}, X_{2}=x_{2}, \ldots, X_{n}=x_{n}\right\}$

$$
P\left\{X_{1}, X_{2}, \ldots, X_{n}\right\}=P\left(x_{1}\right) P\left(x_{2} \mid x_{1}\right) P\left(x_{3} \mid x_{1}, x_{2}\right) \cdot \cdot P\left(x_{n} \mid x_{1}, \ldots x_{n-1}\right)
$$

But $P\left(x_{r} \mid x_{1}, x_{2}, \ldots, x_{r-1}\right)=P\left(x_{r} \mid x_{r-1}\right)$

$$
\therefore P\left(X_{1}, X_{2}, \ldots, X_{n}\right)=P\left(x_{1}\right) P\left(x_{2} \mid x_{1}\right)\left(x_{3} \mid x_{2}\right) \ldots P\left(x_{n} \mid x_{n-1}\right)
$$

where the probabilities can be expressed as

$$
P=\left[\begin{array}{ll}
P\left(X_{n}=0 \mid X_{n-1}=0\right) & P\left(X_{n}=1 \mid X_{n-1}=0\right) \\
P\left(X_{n}=0 \mid X_{n-1}=1\right) & P\left(X_{n}=1 \mid X_{n-1}=1\right)
\end{array}\right]=\left[\begin{array}{ll}
p_{0} & q_{0} \\
q_{1} & p_{1}
\end{array}\right]
$$

c. Prevalence and Incidence Models
$I(t): \quad$ Incidence of disease at time $t$
$P(t)$ : Prevalence of disease at time $t$
$t$ : may refer to age or chronological time
States: $\quad S_{0}$ : disease free
$S_{d}$ : disease
Sample $S_{d}$
Path $\quad S_{0}$

$I(t) d t=P\left\{S_{0} \rightarrow S_{d}\right.$ during $\left.t, t+d t\right\}$
What are relations between $I(t)$ and $P(t)$ ? Are there differences when $t$ is chronological time or age?

## d. Clinical Trials

Patients are entered in a clinical trial at time $t=0$. State of a patient at time $t=0$ is $S_{0}$.

As time goes on, possible other states are:
$S_{1}:$ response, $S_{2}$ : relapse, $S_{3}$ : death (absorbing state)



Note: $S_{3}$ is absorbing state.

- Survival Time is time from $S_{0} \rightarrow S_{3}$
- Does time in state depend on next state?
e. Early Detection of Disease

Suppose a person can be in one of three states; i.e.
$S_{0}$ : Disease Free
$S_{p}$ : Pre-clinical State (Individual has disease, but is unaware of it. Disease can be diagnosed by special exam)
$S_{c}: \quad$ Clinical Diagnosis

- Progressive Disease Model: $S_{0} \rightarrow S_{p} \rightarrow S_{c}$
- Goal of Early Detection Program is to find people in $S_{p}$


## Lead Time

Backward Recurrence Time
(Forward Recurrence Time)


Problems What is distribution of Lead Time? Note lead time is not observed.
What is distribution of backward recurrence time? (Time in $S_{p}$ before diagnosis). Note that the time when the transition $S_{0} \rightarrow S_{p}$ takes place is not observed.


At time of exam, those in $S_{p}$ are diagnosed (Assuming exam sensitivity is unity). Case finding is "throwing" a random vertical line on figure. Diagnosis is when vertical line intersects horizontal line.

Note larger horizontal lines have a greater chance of being intersected.
Longer horizontal line $\Longrightarrow$ Longer time in $S_{p}$
Longer Time in $S_{p} \Longrightarrow$ slower evolving disease.
Slower evolving disease $\Longrightarrow$ live longer.

## Problems

- What is distribution of time in $S_{p}$ for those diagnosed by special exam compared to those not diagnosed by special exam?
- What is probability of detection by exam if disease incidence depends on age?


## f. Planning Early Detection Trials

Eligible $\Longrightarrow\left|\begin{array}{c}\text { R } \\ \text { A } \\ \mathrm{N} \\ \mathrm{D} \\ \mathrm{O} \\ \mathrm{M} \\ \mathrm{I} \\ \mathrm{Z} \\ \mathrm{E}\end{array}\right|$ - Study Group ( $k$ special exams)

End-Point: Disease Mortality
Issues
How many exams ( $k$ )?
Spacing between exams (staggered, equidistant)
Optimal time for analysis?

## g. Cell Kinetics

Life cycle of cell consists of 4 states; i.e.

$$
\mathrm{G}_{1} \rightarrow \mathrm{~S} \rightarrow \mathrm{G}_{2} \rightarrow \text { Mitosis } \rightarrow \text { Daughter Cell }
$$

- Interest is in measuring total cell cycle time and time in S-state (DNA Synthesis State)
- Technique is to expose cells to a radio active label (Tridiated Thymadine, HTDR)
- Cells which are in $S$ state at time of labeling incorporate HTDR. Samples of cells are taken at subsequent times.
- Label Index $=$ Proportion of labeled cells $=L(t)$.


NON-DIVIDING CELLS


Population of asynchronous proliferating cells. Labeled cells are those in S-phase at time label is given to cell population.

A cell with long S-phase relative to cell cycle time has a higher probability of being labelled. Therefore population of labelled cells is not the same population as the initial cell population. The sample of cells selected by labelling is neither a representative or random sample of cells.

Problem: How to estimate cell cycle time?
h. Models for Prevention Trials (Cancer)

Define:
$S_{0}$ : Disease free state
$S_{c a}$ : Carcinogenic state
$S_{p}$ : Pre-clinical state
$S_{c}: \quad$ Clinical state
$S_{d}$ : Death
$S_{p} \rightarrow S_{c}$ and transitions $S_{0} \rightarrow S_{d}, S_{c} \rightarrow S_{d}$ are observable


To prevent disease, it is necessary to either reduce transition $S_{0} \rightarrow S_{c a}$ and/or $S_{c a} \rightarrow S_{c}$.
How does one model process?

