Discrete Time Markov Chains. Applications in Bioinformatics

Javier Campos Universidad de Zaragoza <u>http://webdiis.unizar.es/~jcampos/</u>

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

- Markov processes: special class of stochastic processes that satisfy the Markov Property (MP):
 - Given the state of the process at time *t*, its state at time *t* + *s* has probability distribution which is a function of *s* only.
 - i.e. the future behaviour after *t* is independent of the behaviour before *t*.
 - Often intuitively reasonable, yet sufficiently "special" to facilitate effective mathematical analysis.

• We consider Markov processes with discrete state (sample) space.

They are called **Markov chains**.

- If time parameter is discrete $\{t_0, t_1, t_2...\}$ they are called **Discrete Time Markov Chains** (DTMC).
- If time is continuous ($t \ge 0$, $t \in IR$), they are called **Continuous Time Markov Chains** (CTMC).

• Let $X = \{X_n\}$, with $n = 0, 1, ...; X_i \in IN, i \ge 0$ be a non-negative integer valued Markov chain with discrete time parameter n.

Markov Property states that:

$$P(X_{n+1} = j \mid X_0 = x_0, \dots, X_n = x_n) =$$
^(*)

=
$$P(X_{n+1} = j | X_n = x_n)$$
, for $j, n=0, 1...$

^(*) Remember "conditional probability", prob of A, given the occurrence of B: $P(A | B) = \frac{P(A \cap B)}{P(B)}$.

- Evolution of a DTMC is completely described by its 1-step transition probabilities $p_{ij}(n) = P(X_{n+1} = j | X_n = i)$ for $i, j, n \ge 0$
- If the conditional probability is invariant with respect to the time origin, the DTMC is said to be time-homogeneous

$$p_{ij}(n) = p_{ij}$$

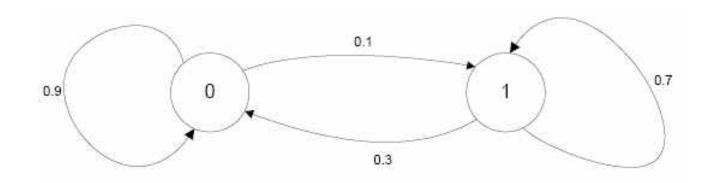
 $\sum_{j \in \Omega} p_{ij} = 1, \forall i \in \Omega$

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

- State transition diagram
 - Directed graph
 - number of nodes = number of states (if Ω finite)
 - An arc from *i* to *j* if and only if $p_{ij} > 0$

Telephone line example: line is either idle (state 0) or busy (state 1)

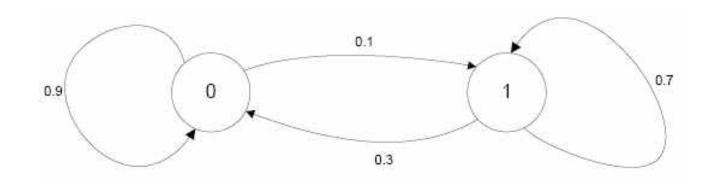


• Transition probability matrix

$$P = \begin{bmatrix} p_{00} & p_{01} & \cdots \\ p_{10} & p_{11} & \cdots \\ \vdots & & & \\ p_{i0} & p_{ii} & \cdots \\ \vdots & & \vdots & \end{bmatrix}$$
 in which all rows sum to 1

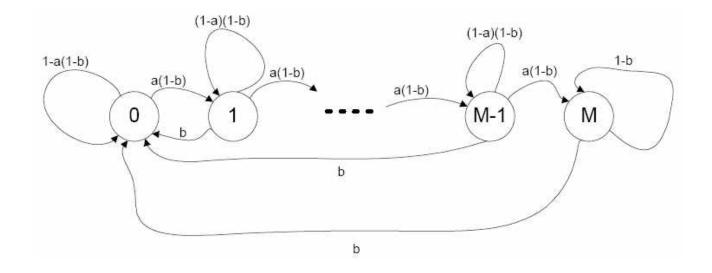
- dimension = number of states in Ω if finite, otherwise countably infinite
- conversely, any real matrix P s.t. $p_{ij} \ge 0$, $\sum_j p_{ij} = 1$ (called a stochastic matrix) defines a MC

Telephone line example



$$P = \begin{bmatrix} 0.9 & 0.1 \\ 0.3 & 0.7 \end{bmatrix}$$

Example 2: I/O buffer, capacity *M* records
 New record added in any unit of time with prob. *a* (if not full).
 Buffer emptied in any unit of time with prob. *b*.
 If both occur in same interval, insertion done first.
 Let X_n be the number of records in buffer at (discrete) time *n*.
 Then, assuming that insertions and emptying are independent of each other and of their own past histories, {X_n | n=0,1,...} is a MC with state space {0,1,...,M} and state diagram:

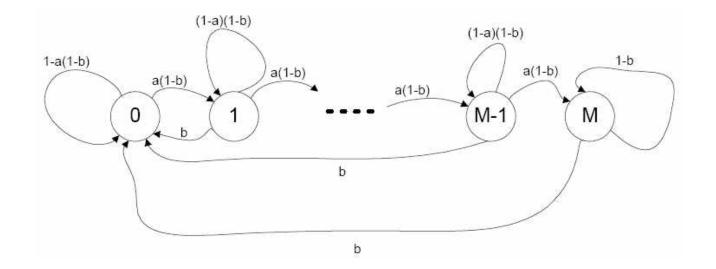


• The transition probability matrix follows immediately, e.g.:

$$p_{12} = a(1 - b) = p_{n,n+1}$$
, $0 \le n \le M - 1$

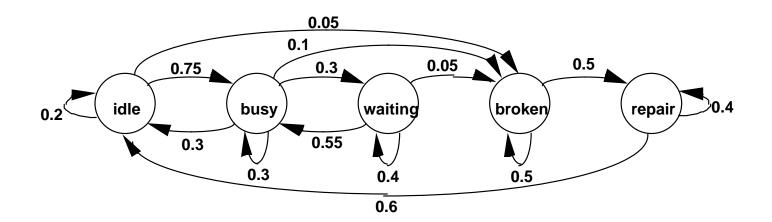
$$p_{MM} = 1 - b$$

etc.

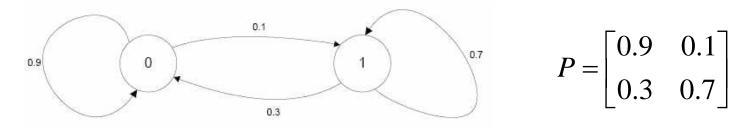


- Example 3:
 - A system that can be
 - Idle
 - Busy
 - Waiting for a resource
 - Broken
 - Repairing

	idle	busy	wait	broken repair	
	0.2	0.75	0.0	0.05	0.0
	0.3	0.3	0.3	0.1	0.0
<i>P</i> =	0.0	0.55	0.4	0.05	0.0
	0.0	0.0	0.0	0.5	0.5
	_0.6	0.0	0.0	0.0	0.4



• Time spent in a state:



- T_0 = random variable "time spent in state 0"

 $P(T_0=1) = (1-p_{00})$ $P(T_0=2) = p_{00} (1-p_{00})$ $P(T_0=3) = p_{00}^2 (1-p_{00})$... $P(T_0=n+1) = p_{00}^n (1-p_{00})$

 \rightarrow Geometrically distributed random variable Is the discrete analogue of exponential distribution \rightarrow memoryless

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

• Let the 2-step transition probability be

$$p_{ij}^{(2)} = P(X_{n+2} = j | X_n = i)$$

= $\sum_{k \in \Omega} P(X_{n+1} = k, X_{n+2} = j | X_n = i)$ by law of tot. prob.
= $\sum_{k \in \Omega} P(X_{n+2} = j | X_n = i, X_{n+1} = k) P(X_{n+1} = k | X_n = i)$
= $\sum_{k \in \Omega} P(X_{n+2} = j | X_{n+1} = k) P(X_{n+1} = k | X_n = i)$ by MP
= $\sum_{k \in \Omega} p_{ik} p_{kj}$
= $(P^2)_{ij}$

• Similarly, the *n*-step transition probability

$$p_{ij}^{(n)} = P(X_n = j | X_0 = i)$$

= $\sum_{k \in \Omega} P(X_n = j | X_{n-1} = k, X_0 = i) P(X_{n-1} = k | X_0 = i)$
= $\sum_{k \in \Omega} p_{ik}^{(n-1)} p_{kj}$

In matrix form:

$$P^{(n)} = P^{(n-1)} \times P$$

If n=2:

$$P^{(2)} = P^{(1)} \times P = P^2$$

And in general: $P^{(n)} = P^n$ i.e. $p_{ij}^{(n)} = (P^n)_{ij}$

- A more general version of previous equations
 - Chapman-Kolmogorov equations

$$p_{ij}^{(n+m)} = \sum_{k \in \Omega} p_{ik}^{(n)} p_{kj}^{(m)}$$

Because

$$p_{ij}^{(n+m)} = \sum_{k \in \Omega} P(X_{n+m} = j \mid X_n = k, X_0 = i) P(X_n = k \mid X_0 = i)$$

Thus

$$P^{(n+m)} = P^{(n)} \times P^{(m)} = P^n \times P^m$$

- Computation of transient distribution
 - Probabilistic behaviour of the Markov chain over any finite period time, given the initial state

$$P(X_n = j | X_0 = i) = p_{ij}^{(n)} = (P^n)_{ij}$$

E.g., in the example of the I/O buffer with capacity of M records, the average number of records in the buffer at time 50 is

$$E(X_{50} \mid X_0 = 0) = \sum_{j=1}^{\min(M, 50)} j q_{0j}^{(50)}$$

- Computation of transient distribution
 - n*th*-step distribution:

$$\pi_{j}(n) = P(X_{n} = j) = \sum_{i \in \Omega} P(X_{0} = i) P(X_{n} = j | X_{0} = i)$$
$$= \sum_{i \in \Omega} \pi_{i}(0) p_{ij}^{(n)} = \sum_{i \in \Omega} \pi_{i}(0) (P^{n})_{ij}$$

– in matrix form:

$$\pi(n)^{\mathsf{T}} = \pi(0)^{\mathsf{T}} P^n$$

- Problem: computationally expensive!

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

• State *j* is **accessible** from state *i* (writen written $i \rightarrow j$) if

 $p_{ij}^{(n)} > 0$, for some n

- A state *i* is said to communicate with state *j* (writen written *i* ↔ *j*) if *i* is accessible from *j* and *j* is accessible from *i*
- A set of states C such that each pair of states in C communicates is a **communicating class**
- A communicating class is closed if the probability of leaving the class is zero (no state out of C is accesible from states in C)
- A Markov chain is **irreducible** if the state space is a communicating class
- State *i* is an **absorbing** state if there is no state reachable from

- Periodicity:
 - A state *i* has **period** *k* if any return to state *i* must occur in some multiple of *k* time steps.

$$k = \gcd\{n: P(X_n = i | X_0 = i) > 0\}$$

- If k = 1, then the state is aperiodic; otherwise (k>1), the state is periodic with period k.
- It can be shown that every state in a communicating class must have the same period.
- An irreducible Markov chain is aperiodic if its states are aperiodic.

- Recurrence
 - A state *i* is transient if, given that we start in state *i*, there is a non-zero probability that we will never return back to *i*.
 - Formally, next return time to state *i* ("hitting time"):

 $T_i = \min\{n : X_n = i \mid X_0 = i\}$

- State *i* is transient if $P(T_i < \infty) < 1$ (i.e. $P(T_i = \infty) > 0$)
- If a state *i* is not transient (it has finite hitting time with probability 1), then it is said to be **recurrent**.
- Let M_i be the expected (average) return time, $M_i = E[T_i]$
 - Then, state *i* is **positive recurrent** if *M_i* is finite; otherwise, state *i* is **null recurrent**.
- It can be shown that a state is recurrent iff $\sum_{n=0}^{\infty} p_{ii}^{(n)} = \infty$

- In a finite DTMC:
 - All states belonging to a closed class are positive recurrent.
 - All states not belonging to a closed class are transient.
 - There are not null recurrent states.
- In an irreducible DTMC:
 - Either all states are transient or recurrent (in case of finite DTMC, all are positive recurrent).
- Ergodicity:
 - A state *i* is said to be **ergodic** if it is aperiodic and positive recurrent (finite average return time).
 - If all states in a DTMC are ergodic, the chain is said to be ergodic.

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

- Transient behaviour: computationally expensive
- Easier and maybe more interesting to determine the limit or steady-state distribution

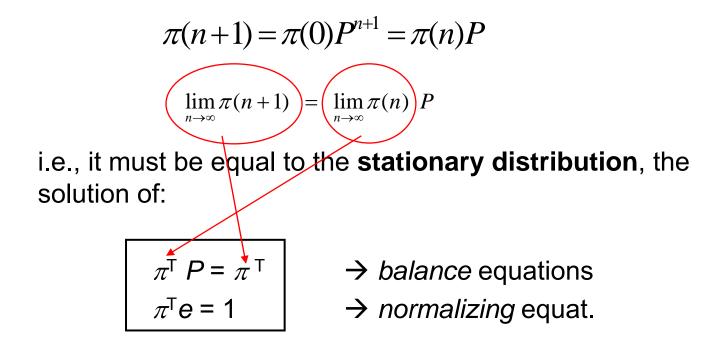
$$\pi_j = \lim_{n \to \infty} \pi_j(n)$$

In vector form

$$\pi = \lim_{n \to \infty} \pi(n)$$

- Does it exist?
- Is it unique?
- Is it independent of the initial state?

• If limit distribution exists... we know how to compute it!



where $e = (1, 1, ..., 1)^T$, and the initial distribution does not affect the limit distribution

- Other interpretation:
 - The solution of balance equations can be seen as the proportion of time that the process enters in each state in the long run
 - Let N_j(n) be the number of visits of the process to the state j until instant n
 - The occupation distribution can be defined as

$$\pi_j = \lim_{n \to \infty} \frac{E[N_j(n)]}{n+1}$$

- Of course, its inverse is the mean interval between visits, or mean return time (1/π_i)
- If the occupation distribution exists, it verifies

$$\pi^{\mathsf{T}} P = \pi^{\mathsf{T}}; \pi^{\mathsf{T}} e = 1$$

- But,
 - Does limit distribution exist?
 - Is it unique?
 - Is it independent of the initial state?

We know some cases where the answer is no

We know some cases where the answer is yes

- If a unique limit distribution exists, all rows of P^n must be equal in the limit, in this way the distribution of X_n does not depend on the initial distribution
- Example

$$P = \begin{bmatrix} 0 & 1 & 0 \\ 0.1 & 0 & 0.9 \\ 0 & 1 & 0 \end{bmatrix} \xrightarrow{1}_{0.9} \xrightarrow{0.1}_{1} P^{2n} = \begin{bmatrix} 0.1 & 0 & 0.9 \\ 0 & 1 & 0 \\ 0.1 & 0 & 0.9 \end{bmatrix} P^{2n-1} = \begin{bmatrix} 0 & 1 & 0 \\ 0.1 & 0 & 0.9 \\ 0 & 1 & 0 \end{bmatrix}$$

If *a* is the initial distribution, then the distribution of X_n , $n \ge 1$ is:

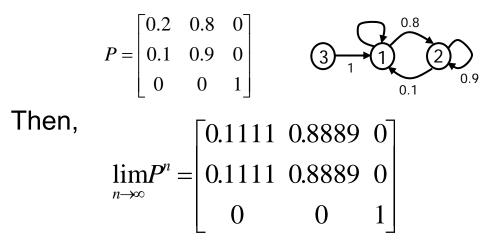
 $(0.1(a_1+a_3), a_2, 0.9(a_1+a_3))$, if *n* is odd $(0.1a_2, a_1+a_3, 0.9a_2)$, if *n* is even

Thus, the DTMC has not limit distribution.

If balance and normalization equations are solved, we get a unique solution $\pi = (0.05, 0.5, 0.45)$.

This means: if π is assumed as initial distribution, then π is also the distribution for X_n , for all n.

• Example: limit and stationary distributions may be non unique



Limit distribution exists, but it is not unique since it depends on the initial distribution: if *a* is the initial distribution

 $\pi = (0.111(a_1 + a_2), 0.8889(a_1 + a_2), a_3)$

is a limit distribution for X_n , and it is also a stationary distribution.

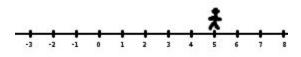
- Finite & irreducible DTMC \Rightarrow there exists a unique stationary distribution
- Finite & irreducible DTMC \Rightarrow there exists a unique occupation distribution, and it is equal to the stationary distribution
- Finite, irreducible & aperiodic DTMC \Rightarrow it has a unique limit distribution, and it is equal to the stationary distribution
- Positive recurrent & aperiodic DTMC \Rightarrow there exists limit distribution
 - If in addition DTMC is irreducible, the limit distribution is independent of the initial probability
- Irreducible, positive recurrent & periodic DTMC with period $d \Rightarrow \lim_{n \to \infty} p_{ij}^{(nd)} = d\pi_j$
- An irreducible & aperiodic DTMC is positive recurrent ⇔ there exists a unique solution of balance equation
- Irreducible, aperiodic & null recurrent DTMC $\Rightarrow \lim_{n \to \infty} p_{ij}^{(n)} = 0$

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

Examples

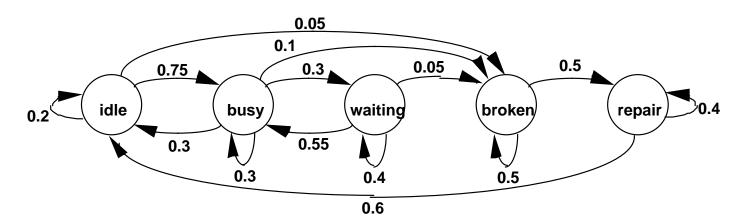
- Random walks
 - A random walk is a random process consisting of a sequence of discrete steps of fixed length.
 - One-dimensional nearest-neighbour random walk



- In this case, the reachable states are integer (or natural) numbers and at each step, the process jumps to the nearest-neighbour to the right with probability *p* or to the nearest-neighbour to the left with probability 1 − *p*.
- A random walk with state space equal to Z (integer numbers) can be transient or null recurrent depending on *p*.
 - If *p*=1/2 then it is null recurrent and in other cases it is transient.
 Look at the book by S. Ross (Example 4.3d).
- A random walk with reflectant barrier at 0 (i.e., state space equal to non negative integers) can be transient, null recurrent or positive recurrent. Look <u>here</u> (local copy).

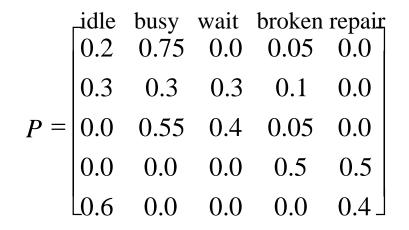
Examples

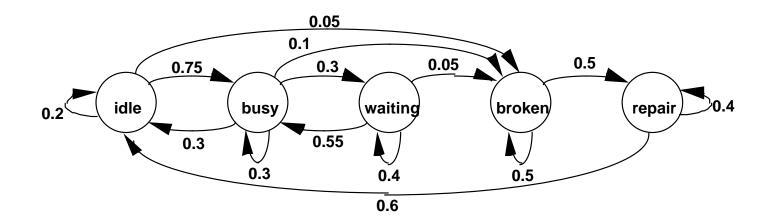
- A processor has certain tasks to perform
 - State transition diagram. Possible states:
 - idle (no task to do)
 - busy (working on a task)
 - waiting (stopped for some resource)
 - broken (no longer operational)
 - repair (fixing the failure)





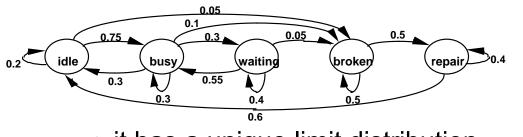
– Transition probability matrix representation





Examples

- Properties: finite state space, irreducible, aperiodic



 \Rightarrow it has a unique limit distribution, and it is equal to the stationary distribution

– Solution:

$$\pi = \lim_{n \to \infty} \pi(n) \qquad \qquad \pi^{\mathsf{T}} \mathsf{P} = \pi^{\mathsf{T}} \\ \pi^{\mathsf{T}} \mathsf{e} = \mathsf{1}$$

 $\pi = (0.2155, 0.3804, 0.1902, 0.1167, 0.0972)^{\mathsf{T}}$

Examples

$\pi = (0.2155, 0.3804, 0.1902, 0.1167, 0.0972)^{\mathsf{T}}$

- Compute performance indices:
 - Availability: P(idle + busy + wait) = 0.7861 (in other words, 78.61% of the time)
 - So, not available: 21.39% of the time
 - *Working time*: *P*(busy + wait) = 0.57

Outline

- Basic definitions
- Representations
- Multi-step transition probabilities
- Classification of states
- Steady-state behaviour
- Examples
- Applications in Bioinformatics

Simple model of virus mutation

- Suppose a virus can exist in *N* different strains and in each generation either stays the same, or with probability α mutates to another strain, which is chosen at random.
- What is the probability that the strain in the nth generation is the same as that in the Oth?

N-state chain, with $N \ge N$ transition matrix *P* given by

$$P_{ii} = 1 - \alpha$$

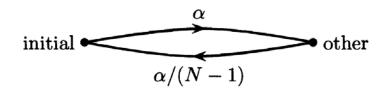
$$P_{ij} = \alpha / (N - 1), \text{ for } i \neq j.$$

Answer: computing $p_{11}^{(n)}$.

Much simpler: exploiting the symmetry present in the mutation rules...

Simple model of virus mutation

 At any time a transition is made from the initial state to another with probability α, and a transition from another state to the initial state with probability α /(N − 1) → two-state chain



$$P^{n+1} = P^n P \rightarrow p_{11}^{(n+1)} = p_{12}^{(n)} \beta + p_{11}^{(n)} (1 - \alpha), \text{ with } \beta = \alpha / (N - 1).$$

Since $p_{11}^{(n)} + p_{12}^{(n)} = 1$ then we get the recurrence $p_{11}^{(n+1)} = (1 - \alpha - \beta) p_{11}^{(n)} + \beta$, and $p_{11}^{(0)} = 1$

That has a unique solution

$$p_{11}^{(n)} = \begin{cases} \frac{\beta}{\alpha + \beta} + \frac{\alpha}{\alpha + \beta} (1 - \alpha - \beta)^n & \text{for } \alpha + \beta > 0\\ 1 & \text{for } \alpha + \beta = 0. \end{cases}$$

Then, in our case $p_{11}^{(n)} = \frac{1}{N} + \left(1 - \frac{1}{N}\right) \left(1 - \frac{\alpha N}{N-1}\right)^n$.

Life behaviour model exercise

- An octopus, called Paul, is trained to choose object A from a pair of objects A, B, by being given repeated trials in which it is shown both and is rewarded with food if it chooses A.
- Modelling its mind states...
 - state 1: it cannot remember which object is rewarded
 → it is equally likely to choose either
 - state 2: it remembers and chooses A but may forget again
 - state 3: it remembers and chooses A and never forgets
- Modelling its evolution in time
 - After each trial it may change its state of mind according to the transition matrix

state 1	1/2	1/2	0
state 2	1/2	1/12	5/12
state 3	0	0	1

Life behaviour model exercise

- Questions: It is in state 1 before the first trial,
 - What is the probablity that it is in state 1 just before the (*n*+1)*th* trial?
 - Answer.
 - What is the probability $P_{n+1}(A)$ that it chooses A on the (n+1)th trial?
 - <u>Answer</u>.
 - Someone suggests that the record of successive choices (a sequence of As and Bs) might arise from a two-state Markov chain with constant transition probabilities.

Discuss, with reference to the value of $P_{n+1}(A)$ that you have found, whether this is possible.

• <u>Answer</u>.

Other examples

- Population genetics, Norris' book:
 - Wright-Fisher model
 - Moran model
- Epidemic models, Linda Allen's book:
 - SIS Epidemic Model
 - Chain Binomial Epidemic Models

(you can read it at

http://webdiis.unizar.es/asignaturas/SPN/?page_id=104)

Many other applications...

- Birth-death processes used to model the evolution of populations
- Markov models appear also in chemical and biochemical kinetics
- Are the basis for Hidden Markov Models
- Et cetera...