Computational Methods in Stochastics

Lecture I

Intro **Probability Review** Random variables **Common Distributions Random Number** Generators

Course "Statements"

With this course I try to give a reasonable understanding on the fundamentals of stochastic processes and of common computational methods used to simulate them.

- The formalism needed to state things precisely is tedious, sometimes even overwhelming. After this course you hopefully will be fluent enough with the concepts and notation to be able to read the literature, so that you can learn and understand new things.
- The field is vast! You will not run out of things to learn during your lifetime.
- When reading the slides, try to understand the messages conveyed by the formalism. These slides are also meant to be a reference for you; hence they are detailed.
- Try to develop a feeling for how formal things could be implemented as algorithms.

Course "Statements"

Most importantly, try to find the fun in implementing methods as algorithms. Optimally, python and stochastic simulation should start seeming like a playground for you.

For those into machine learning: understanding the concepts within this course will make understanding machine learning methods a whole lot easier.

The final exam will concentrate on understanding and I will write a page or two about what you should understand and at how detailed a level (of notation) before the time comes.

Sometimes understanding exactly what is expected in the assignments can be hard. **Please ask TAs in good time before the deadline.** TAs are not obliged to respond to zillions of Slack messages 1 hour before DL. – **And please be polite.**

Stochastic Modelling

The word "stochastic" derives from the Greek ($\sigma \tau \sigma \chi \alpha \zeta \epsilon \sigma \theta \alpha \iota$ to aim, to guess) and means "random" or "chance".

A deterministic model predicts a single outcome from a given set of circumstances.

A stochastic model predicts a set of possible outcomes weighted by their likelihoods or probabilities.

Stochastic modelling can be applied also to deterministic states whose outcome is not known (e.g. a hidden tossed coin).

Notation for stochastic processes is really compact: Computing a **deterministic equation** $x = -A \log(u)$ means computing x from a single value of u, whereas the **stochastic equation** $X = -A \log(U)$ requires computing from the distribution U.

Stochastic Modelling

Scientific modelling (SM) has three components: (i) *a natural phenomenon* under study, (ii) *a logical system* for deducing implications about the phenomenon, and (iii) *a connection* linking the elements of the natural system under study to the logical system used to model it.

Example: (i) The earth with airports, (ii) mathematics of spherical geometry, and (iii) viewing the airports in the physical system as points in the logical system.

Stochastic SM is based on **the law of large numbers**: The relative fraction of times in which an event occurs in a sequence of independent similar experiments approaches, in the limit of an infinite sequence, the probability of the occurrence of the event on any single trial. (Related to ergodicity.)

Stochastic Modelling

A stochastic process is a family of random variables X_t , where t is a parameter running over a suitable index set T. (Sometimes we write X(t) instead of X_t .)

In a common situation the index *t* corresponds to discrete units of time, and the index set is $T = \{0, 1, 2, ...\}$.

Stochastic processes for which $T = [0, \infty)$ are important. t often represents time. It may also represent e.g. distance from an arbitrary origin, and X_t may count the number of defects in the interval (0, t] along a thread, or the number of cars in the interval (0, t] along a highway.

Stochastic processes are distinguished by their **state space**, or the range of possible values for the random values X_t , by their index set T, and by the dependence relations among the random variables X_t .

Probability Review

- **For a clear review, see:** <u>Intro of the online book</u> Let *A* and *B* be **events.**
 - The event that at least one of *A* or *B* occurs: $A \cup B$ (union).
 - The event that both *A* and *B* occur: $A \cap B$, or *AB* (intersection).
- This notation extends to *finite* and *countable* sets of events $A_1, A_2, ...$
- At least one event occurs: $A_1 \cup A_2 \cup \dots = \bigcup_{i=1}^{\infty} A_i$. All events occur: $A_1 \cap A_2 \cap \dots = \bigcap_{i=1}^{\infty} A_i$.
- The probability of an event *A*: Pr{*A*}.
- The *certain* event is denoted by Ω : $Pr{\Omega} = 1$. The *impossible* event is denoted by \emptyset : $Pr{\emptyset} = 0$.

Probability Review

Disjoint events, $A \cap B = \emptyset$, cannot both occur.

The *addition law* for disjoint events: $\Pr{A \cup B} = \Pr{A} + \Pr{B}$; if events A_i and A_j are disjoint for $i \neq j$, then $\Pr{\bigcup_{i=1}^{\infty}} = \sum_{i=1}^{\infty} \Pr{A_i}$.

The law of total probability: Let $A_1, A_2, ...$ be disjoint events for which $\Omega = A_1 \cup A_2 \cup ...$ (exactly one event will occur). Then $\Pr{B} = \sum_{i=1}^{\infty} \Pr{B \cap A_i}$ for any event *B*.

Many important equations and principles are derived from the law of total probability.

Events are said to be *independent* if $Pr{A \cap B} = Pr{A} \times Pr{B}$, or $Pr{A_{i_1} \cap A_{i_2} \cap \cdots \cap A_{i_n}} = Pr{A_{i_1}}Pr{A_{i_2}} \cdots Pr{A_{i_n}}$ for every infinite set of distinct indices $i_1, i_2, ..., i_n$.

Random Variables See Introduction to Probability, Statistics and Random Processes, Chapter 3

- A useful non-formal definition:
- A *random variable* is one that takes on its values by chance.
- Random variables are denoted by capital letters, e.g. X, Y, and Z.
- Real numbers are denoted by lowercase letters, e.g. *x*, *y*, and *z*.
- The expression $\{X \le x\}$ is the event that the random variable X assumes a value that is less than or equal to the real number x.
- The probability that this event occurs is $Pr{X \le x}$.

Allowing *x* to vary, this probability defines the *distribution function* (or cumulative distribution function, CDF) of the random variable *X* as

 $F(x) = \Pr\{X \le x\}, \qquad -\infty < x < +\infty.$

Subscripts are used, when several random variables appear in the same context, $F_X(\xi) = \Pr\{X \le \xi\}$ and $F_Y(\xi) = \Pr\{Y \le \xi\}$.

It's easy to see that, for example, $\Pr{X > a} = 1 - F(a)$, $\Pr{a < X \le b} = F(b) - F(a)$, and $\Pr{X = x} = F(x) - \lim_{\epsilon \downarrow 0} F(x - \epsilon) = F(x) - F(x-)$.

Discrete random variable X: There is a finite or denumerable set of distinct values $x_1, x_2, ...$ such that $a_i = \Pr\{X = x_i\} > 0$ for i = 1, 2, ... and $\sum_i a_i = 1$.

<u>The probability mass function (PMF)</u> for the random variable *X*: $p(x_i) = p_X(x_i) = a_i$ for i = 1, 2, ...

The relation between the probability mass and distribution function:

$$p(x_i) = F(x_i) - F(x_i)$$
 and $F(x) = \sum_{x_i \le x} p(x_i)$.

Probability Mass Function



Distribution function at value 3 would be F(x = 3) = p(0) + p(1) + p(2) + p(3). For discrete random variables *X*, *p* and *F* are discrete.

For a *continuous random variable X*: $Pr{X = x} = 0 \quad \forall x$.

If there is a nonnegative function $f(x) = f_X(x)$ defined for $-\infty < x < \infty$ such that

 $\Pr\{a < X \le b\} = \int_{a}^{b} f(x) \quad \text{for } -\infty < a < b < \infty,$ then $f(x) = f_X(x)$ is called <u>the *probability density function* (PDF)</u> for the random variable *X*.

Then <u>the *cumulative distribution* (CDF)</u> takes the form:

$$F(x) = F_X(x) = \Pr\{X \le x\} = \int_{-\infty}^x f(z) dz.$$

$$\Rightarrow f_X(x) = \frac{d}{dx} F_X(x).$$

Then the continuous distribution function $F(x) = \int_{-\infty}^{x} f(\xi) d\xi, \quad -\infty < x < \infty.$

If F(x) is differentiable in x, $f(x) = \frac{d}{dx}F(x) = F'(x)$, $-\infty < x < \infty$. $\Rightarrow \Pr\{x < X \le x + dx\} = F(x + dx) - F(x) = dF(x) = f(x)dx$.

More precisely, $\Pr\{x < X \le x + \Delta x + o(\Delta x)\}, \Delta x \downarrow 0$. $o(\Delta x)$ represents any term for which $\lim_{\Delta x \downarrow 0} o(\Delta x)/\Delta x = 0$.

The *m*th *moment* of a discrete random variable *X*:

$$E[X^m] = \sum_i x_i^m \Pr\{X = x_i\}$$

If the infinite sum diverges, the moment is said not to exist.

The *m*th *moment* of a continuous random variable *X*:

$$E[X^m] = \int_{-\infty}^{\infty} x^m f(x) dx$$

(the integral must converge absolutely).

The first moment, m = 1, is called <u>the *mean*</u> or *expected value* of *X*, denoted by m_X or μ_X .

<u>The *m*th *central moment* of *X* is defined as the *m*th moment of the random variable $X - \mu_X$.</u>

The second central moment is called the *variance* of *X* and written σ_X^2 or Var[*X*]. Var[*X*] = $E[(X - \mu)^2] = E[X^2] - \mu^2$.

 σ_X is called the *standard deviation* (stdev).

<u>The *median*</u> of a random variable *X* is any value *v* such that $\Pr{X \ge v} \ge \frac{1}{2}$ and $\Pr{X \le v} \ge \frac{1}{2}$. <u>The *mode*</u> of a random variable *X* is the value $x \in X$, where PMF or PDF has the maximum value: $\Pr{x = x_m} = \max{\Pr{x}}$.

The more asymmetric (= skewed) the distribution, the less descriptive is the mean value.



Skewness = 3 *(mean – median)/stdev is a *measure of distribution's asymmetry* (normal distribution is symmetrical).

Y = g(X) is also a random variable. The expectation of g(X): $E[g(X)] = \sum_{i=1}^{\infty} g(x_i) \Pr\{X = x_i\}.$

For continuous *X*: $E[g(X)] = \int g(x) f_X(x) dx$.

Generally, for both the discrete (in the limit of fine discretisation) and continuous cases:

 $E[g(X)] = \int g(x) dF_X(x),$

where F_X is the distribution function of the random variable *X*. (Lebesque-Stieltjes integral.)

Given a pair (*X*, *Y*) of random variables, their *joint distribution function* is given by

$$\widetilde{F}_{XY}(x,y) = F(x,y) = \Pr\{X \le x \text{ and } Y \le y\}.$$

A joint distribution is said to possess a (joint) probability density if there exists a function f_{XY} of two real variables for which

$$F_{XY}(x,y) = \int_{-\infty}^{x} \int_{-\infty}^{y} f_{XY}(\xi,\eta) d\eta d\xi \quad \forall x,y.$$

The marginal distribution functions of X and Y are $F_X(x) = \lim_{y \to \infty} F_{XY}(x, y)$ and $F_Y(y) = \lim_{x \to \infty} F_{XY}(x, y)$, respectively. $\swarrow y$ "integrated out"

The marginal density functions are $f_X(x) = \int_{-\infty}^{\infty} f(x, y) dy$ and $f_Y(y) = \int_{-\infty}^{\infty} f(x, y) dx$.

If *X* and *Y* are jointly distributed, then E[X + Y] = E[X] + E[Y].

The random variables *X* and *Y* are said to be *independent* if $F(x, y) = F_X(x) \times F_Y(y) \quad \forall x, y.$

Then the joint density function $f(x, y) = f_X(x)f_Y(y) \forall x, y$.

Given that the jointly distributed *X* and *Y* have means μ_X and μ_X and finite variances, the *covariance* of *X* and *Y* is $Cov[X,Y] = \sigma_{XY} = E[(X - \mu_X)(Y - \mu_Y)] = E[XY] - \mu_X\mu_Y$

X and *Y* are said to be *uncorrelated* if $\sigma_{XY} = 0$.

Independent random variables having finite variances are uncorrelated, but the converse is not true; there are uncorrelated random variables that are not independent.

Correlation coefficient:
$$\rho = \frac{\sigma_{XY}}{\sigma_X \sigma_Y}$$
.

The *joint distribution function* of any finite collection $X_1, X_2, ..., X_n$:

$$F(x_1, \dots, x_n) = F_{X_1, \dots, X_n}(x_1, \dots, x_n) = \Pr\{X_1 \le x_1, \dots, X_n \le x_n\}$$

Independence of $X_1, X_2, ..., X_n$:

$$F(x_1, \dots, x_n) = F_{X_1}(x_1) \dots F_{X_n}(x_n) \ \forall x_1, \dots, x_n$$

A joint distribution function is said to have a *probability density function* $f(\xi_1, ..., \xi_n)$ if

$$F(x_1,\ldots,x_n) = \int_{-\infty}^{x_1} \cdots \int_{-\infty}^{x_n} f(\xi_1,\ldots,\xi_n) d\xi_1,\ldots,d\xi_n \ \forall x_1,\ldots,x_n.$$

For jointly distributed $X_1, X_2, ..., X_n$ and arbitrary functions $h_1, ..., h_m$ of *n* variables each, the *expectation* is:

$$E\left[\sum_{j=1}^{m} h_{j}(X_{1}, \dots, X_{n})\right] = \sum_{j=1}^{m} E\left[h_{j}(X_{1}, \dots, X_{n})\right].$$

If *X* and *Y* are independent random variables having distribution functions F_X and F_Y , respectively, then the distribution function of their sum Z = X + Y is the *convolution* of F_X and F_Y :

$$F_Z(z) = \int_{-\infty}^{\infty} F_X(z-\xi) dF_Y(\xi) = \int_{-\infty}^{\infty} F_Y(z-\eta) dF_X(\eta).$$

Respectively, for probability density functions:

$$f_Z(z) = \int_{-\infty}^{\infty} f_X(z-\xi) f_Y(\xi) d\xi = \int_{-\infty}^{\infty} f_Y(z-\eta) f_X(\eta) d\eta.$$

(for nonnegative random variables, replace the lower limit $-\infty$ by 0.)

The variance: $\sigma_Z^2 = \sigma_X^2 + \sigma_Y^2$

For all events *A* and *B* such that $Pr{B} > 0$, **the conditional probability** of *A* given *B* is written

$$\Pr\{A|B\} = \frac{\Pr\{A \cap B\}}{\Pr\{B\}} \quad \text{if } \Pr\{B\} > 0.$$

This is the **Bayes Theorem**.

If
$$\Omega = B_1 \cup B_2 \cup \cdots$$
 and $B_i \cap B_j = \emptyset$ for $i \neq j$, then

$$\Pr\{A\} = \sum_{i=1}^{\infty} \Pr\{A \cap B_i\} = \sum_{i=1}^{\infty} \Pr\{A|B_i\} \Pr\{B_i\}.$$

This is the law of total probability.

Bernoulli Distribution

Random variable *X* has two possible values 0 and 1.

The probability mass function (PMF) p(1) = p and p(0) = 1 - p where 0 .The mean: <math>E[X] = p. The variance: Var[X] = p(1 - p).

Bernoulli random variables occur frequently as indicators of events. The *indicator of an event A* is the random variable

 $\mathbf{1}(A) = \mathbf{1}_A = \begin{cases} 1 & \text{if } A \text{ occurs} \\ 0 & \text{if } A \text{ does not occur} \end{cases}$

 $\mathbf{1}_A$ is a Bernoulli random variable with parameter $p = E[\mathbf{1}_A] = \Pr{A}$.

Binomial Distribution

Independent events $A_1, A_2, ..., A_n$ all having the same probability $p = \Pr{A_i}$ of occurrence. Let *Y* count the total number of events among $A_1, A_2, ..., A_n$ that occur. Then *Y* has a binomial distribution with parameters *n* and *p*. The probability mass function:

$$p_Y(k) = \Pr\{Y = k\} = \frac{n!}{k! (n-k)!} p^k (1-p)^{n-k} \text{ for } k = 0, 1 \dots, n.$$

Using indicator function, for which $\mathbf{1}(A) = 1$ or $\mathbf{1}(A) = 0$, when event A takes place or not, simplifies maths: Write Y as a sum of indicators $Y = \mathbf{1}(A_1) + \cdots + \mathbf{1}(A_n)$ to determine the moments.

$$E[Y] = E[\mathbf{1}(A_1)] + \dots + E[\mathbf{1}(A_n)] = np$$

Var[Y] = Var[\mathbf{1}(A_1)] + \dots + Var[\mathbf{1}(A_n)] = np(1-p)

Notation: $X \sim Bin(n, p)$ means X is a binomial random quantity based on *n* independent trials, each occurring with probability *p*.



Binomial is the discrete version of the normal/Gaussian.

The Poisson Distribution

The probability mass function of the Poisson distribution with parameter $\lambda > 0$

$$p(k) = \frac{\lambda^k e^{-\lambda}}{k!} \qquad \text{for } k = 0, 1, \dots$$

In calculations related to the Poisson distribution the series expansion for the exponential comes in handy

$$e^{\lambda} = 1 + \lambda + \frac{\lambda^2}{2!} + \frac{\lambda^3}{3!} + \cdots$$
$$\sum_{k=0}^{\infty} kp(k) = \lambda e^{-\lambda} \sum_{k=1}^{\infty} \frac{\lambda^{k-1}}{(k-1)!} = \lambda, \sum_{k=0}^{\infty} k(k-1)p(k) = \lambda^2$$

In terms of a Poisson distributed random variable $X \sim Po(\lambda)$:

Mean : $E[X] = \lambda$

Variance:
$$Var[X] = E[X^2] - {E[X]}^2 =$$

= $E[X(X - 1)] + E[X] - {E[X]}^2 = \lambda$

The simplest form of the *law of rare events:* The binomial distribution with parameter *n* and *p* converges to the Poisson distribution with parameter λ if $n \rightarrow \infty$ and $p \rightarrow 0$ so that $\lambda = np$ remains constant. (See 'An Introduction to Stochastic Modeling' for a proof.)

An important property: The sum of Poisson random quantities is also a Poisson random quantity.



The Multinomial Distribution

joint distribution of *r* variables taking nonnegative values0, ..., *n*. The joint probability mass function

$$\Pr\{X_1 = k_1, \dots, X_n = k_n\} = \begin{cases} \frac{n!}{k_1! \cdots k_r!} p_1^{k_1} \cdots p_r^{k_r} & \text{if } \sum_{i=1}^r k_i = n \\ 0 & \text{otherwise} \end{cases}$$

Here, $p_i > 0$ for i = 1, ..., r and $\sum_{i=1}^{r} p_i = 1$.

Mean: $E[X_i] = np_i$ Variance: $Var[X_i] = np_i(1 - p_i)$ Covariance: $Cov[X_iX_j] = -np_ip_j$

Multinomial is the generalisation of the binomial distribution.

Continuous Distributions

The Normal/Gaussian Distribution $N(\mu, \sigma^2)$

The probability density function $\phi(x;\mu,\sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}}, \quad -\infty < x < \infty.$

Mean: $E[X] = \mu$

Variance: $Var[X] = \sigma^2$

Standard normal distribution when $\mu = 0$ and $\sigma = 1$; N(0,1).

Continuous Distributions

The Normal Distribution



Continuous Distributions See the online book

The central limit theorem: For partial sums $S_n = \xi_1 + \dots + \xi_n$ of independent and identically distributed (i.i.d.) summands ξ_1, ξ_2, \dots having finite means $\mu = E[\xi_k]$ and finite variances $\sigma^2 = \text{Var}[\xi_k]$,

$$\lim_{n \to \infty} \Pr\left\{\frac{S_n - n\mu}{\sigma\sqrt{n}} \le x\right\} = \Phi(x) \ \forall x.$$

Here,
$$\Phi(x) = \int_{-\infty}^{x} \phi(\xi) d\xi = \int_{-\infty}^{x} \frac{1}{\sqrt{2\pi}} e^{-\xi^{2}/2} d\xi$$

(the standard normal distribution function)

So, the normal distribution results for numerous small additive, independent ξ , no matter how they are distributed.

Continuous Distributions

Equivalently, the *central limit theorem* for the sample mean $\overline{X}_n = \frac{1}{n}S_n$: $\lim_{n \to \infty} \Pr\left\{\frac{\overline{X}_n - \mu}{\sigma/\sqrt{n}} \le x\right\} = \Phi(x) \ \forall x.$

For the layman: Make independent observations of any random process \rightarrow The observed values are normally distributed (with the original mean, of course).

This <u>wiki page</u> is good demystification of the CLT.

Continuous Distributions

The Lognormal Distribution

Here the natural logarithm of a nonnegative random variable *V* is normally distributed.

$$f_{V}(\nu) = \frac{1}{\sqrt{2\pi}\sigma\nu} \exp\left\{-\frac{1}{2}\left(\frac{\ln\nu - \mu}{\sigma}\right)^{2}\right\}, \quad \nu \ge 0.$$

Mean:
$$E[V] = \exp\left\{\mu + \frac{1}{2}\sigma^{2}\right\}$$

Variance: $Var[V] = \exp\left\{2\left(\mu + \frac{1}{2}\sigma^{2}\right)\right\} \left[\exp\{\sigma^{2}\} - 1\right].$

Lognormal distributions arise from **multiplicative stochastic processes**. Here, the random process is described by a product (instead of a sum) of independent random variables: for a large number of variables the distribution of ln *V* is normal (central limit theorem), so *V* is lognormally distributed.


(Skewness, see p. 17.)

The lognormal distribution is an example of a fat-tailed or skewed distribution. It is sometimes erroneously interpreted as a logarithmic distribution $p(x) \propto x^{-\alpha}$, x > 0 and the constant $\alpha > 0$. Moreover, many natural processes result in lognormal distribution although it may approximately look like a normal distribution.

Logarithmic central-limit theorem

Just like an **additive process** for independent random variables gives normal distribution in the limit of infinite number of samples, so will a **multiplicative process** for such variables give log-normal distribution in this limit.

In a multiplicative process, taking a logarithm of the variables Z, we see that variables $Y = \log Z$ would result from an additive process and will be distributed normally in the limit of infinite number of samples due to the central-limit theorem, and variables Z from the multiplicative process are distributed log-normally.

Logarithmic central-limit theorem

The outcome of such a multiplicative stochastic process is **log-normal distribution; PDF**:

$$f(x) = \frac{1}{\sqrt{2\pi}x\sigma} \exp\left(-\frac{(\ln x - \mu)^2}{2\sigma^2}\right)$$

A commonly used model for a binary distribution leading to normal distribution is the so-called Galton board, where grains are sifted through equilateral triangles and end in different slots at the bottom. Making the triangles' right sides longer in a certain way log-normal distribution results at the bottom for a sufficiently large number of grains and triangle layers. Multiplication follows this asymmetry in this triangle board (see the reference).



See Limpert, E., Stahel, W. A., and Abbt, M., "Log-normal Distributions across the Sciences: Keys and Clues," BioScience, Vol. 51, No. 5, May, 2001.<u>https://stat.ethz.ch/~stahel/logn</u> <u>ormal/bioscience.pdf</u> (Referred to in numpy reference for numpy.random.lognormal)

The Exponential Distribution

The random variable *T* has an exponential distribution with parameter $\lambda > 0$ ($T \sim Exp(\lambda)$), if the probability density function is

$$f_T(t) = \begin{cases} \lambda e^{-\lambda t} & \text{for } t \ge 0, \\ 0 & \text{for } t < 0. \end{cases}$$

The distribution function

$$F_T(t) = \begin{cases} 1 - e^{-\lambda t} & \text{for } t \ge 0, \\ 0 & \text{for } t < 0. \end{cases}$$

Mean: $E[T] = \frac{1}{\lambda}$ Variance: $Var[T] = \frac{1}{\lambda^2}$

There is an alternative definition for the parameter λ , so be sure to specify which one you are using in assignment problems.



Figure 6. Exponential pdf

To see if a distribution is really exponential, plot it in the **semilogarithmic coordinates** (y-axis logarithmic); you should have a straight line. (For logarithmic distributions, plot both with coordinates logarithmic scales/binning.)

The exponential distribution is **encountered in memoryless processes**, which is why it is relevant for (continuous) Markov chains. (Markov property: the next state is determined only by the present state.)

T is a lifetime. The unit has survived up to time *t*. What is the conditional distribution of the remaining life T - t?

$$\Pr\{T - t > x | T > t\} = \frac{\Pr\{T > t + x, T > t\}}{\Pr\{T > t\}}$$
(Bayes)
$$= \frac{\Pr\{T > t + x\}}{\Pr\{T > t\}}$$
(x > 0)
$$= \frac{e^{-\lambda(t+x)}}{e^{-\lambda t}} = e^{-\lambda x} \leftarrow \text{Independent of the past - no memory.}$$

The Uniform Distribution

The probability density function for a random variable *U* distributed uniformly over the interval [a, b], where a < b:

$$f_X(u) = \begin{cases} \frac{1}{b-a} & \text{for } a \le u \le b, \\ 0 & \text{otherwise.} \end{cases}$$

The distribution function

$$F_X(x) = \begin{cases} 0 & \text{for } u \leq a, \\ \frac{x-a}{b-a} & \text{for } a < x \leq b, \\ 1 & \text{for } x > b. \end{cases}$$

Notation: $X \sim U(a, b)$

Mean: $E[X] = \frac{1}{2}(a+b)$ Variance: $Var[X] = \frac{(b-a)^2}{12}$

The *standard* uniform distribution on the unit interval [0, 1] has a = 0 and b = 1. A random variable having this distribution is usually denoted by $U \sim U(0,1)$. E(U) = 1/2, Var(U) = 1/12.

Standard (pseudo)random number generators implement a random variable uniformly distributed over the interval (0, 1].

The Gamma Distribution

The random variable *X* has a gamma distribution with parameters α , $\beta > 0$, written X ~ $Ga(\alpha, \beta)$, if it has PDF

$$f(x) = \begin{cases} \frac{\beta^{\alpha}}{\Gamma(\alpha)} x^{\alpha-1} e^{-\beta x}, & x > 0\\ 0, & x \le 0, \end{cases}$$

where $\Gamma(\alpha) = \int_0^\infty y^{\alpha-1} e^{-y} dy$ is the *gamma function*. $Ga(1, \lambda) = \text{Exp}(\lambda)$, so the gamma distribution is a generalisation of the exponential distribution.

The mean and variance:

$$E[X] = \frac{\alpha}{\beta}, \qquad \operatorname{Var}[X] = \frac{\alpha}{\beta^2}.$$

A useful property that can be utilised in sampling variates from the gamma distribution is that if $Y = X_1 + X_2, X_1 \sim$ $Ga(\alpha_1, \beta)$, and $X_2 \sim Ga(\alpha_2, \beta)$ are independent and $Y \sim$ $Ga(\alpha_1 + \alpha_2, \beta)$, then $Y \sim Ga(\alpha_1 + \alpha_2, \beta)$.

Frequently CDF-based quantities are used to characterise a distribution:

Median
$$m$$
: $P(X \le m) = \frac{1}{2}$, or equivalently, $F_X(m) = 0.5$.
Lower quartile l : $F_X(l) = 0.25$
Upper quartile u : $F_X(u) = 0.75$

The cumulative distribution (CDF):

$$F(x) = F_X(x) = \Pr\{X \le x\} = \int_{-\infty}^{x} f(z) dz.$$

Quantifying noise

In stochastic modelling or when analysing data that is "noisy" or stochastic, one needs a measure of how noisy some random variable *X* is.

One can use the *variance* $\sigma^2 = Var[X]$, or, for having the noise in the same units as *X*, the *standard deviation* $SD[X] = \sigma$.

A consistent way is to give the noise magnitude relative to the random quantity *X*:

1. The *coefficient of variation* is defined as

$$CV[X] = \frac{SD[X]}{|E[X]|} = \frac{\sigma}{|\mu|}$$

2. *Signal-to-noise ratio* SNR = $\frac{1}{CV[X]}$ is more commonly used in engineering.

3. The *dispersion index*, or *variance-to-mean ratio*:

$$\mathrm{VMR}[X] = \frac{\mathrm{Var}[X]}{\mathrm{E}[X]} = \frac{\sigma^2}{\mu}.$$

Determining distributions

Whether you simulate a stochastic process or try to make sense of measured data, you will be plotting PDFs or PMFs. In other words, you generate histograms of the data. (For clarity you may plot the midpoints of the bins – the histogram bars.)

You will want to **change coordinates** in which the histograms are plotted. Linearly scaled coordinates will do nicely for looking at PDFs of normally distributed data, but in order to determine if the pertinent distribution is e.g **exponential or log-normal**, you need to plot the PDF in **semilogarithmic scale** (*x*- or *y*-axis logarithmic).

Determining distributions

For a first view of skewed distributions you can use a **semilog-scale where y-axis is logarithmic**.

For exponential distribution this view is perfect: a straight line.

Log-normal will not look Gaussian in this view, however.

To see the normally distributed log *x* of **log-normal** distribution you need to both use **logarithmic binning and** plot the PMF with l**ogarithmic** *x***-axis and linear** *y***-axis. Now the PMF looks like a normal distribution.**

To see if the PDF is **logarithmic** (scale invariant, power law) you need to use **logarithmic scale** (both the x- and y-axis logarithmic).

Binning distributions

Binning the data means assorting it to intervals along the abscissa (the horizontal axis).

Standard histogram uses **linear binning.** The intervals are of same size.

With some distributions **logarithmic binning** should be used. Here the interval boundaries are located as $\exp(i/r)$, where $i \in \mathbb{Z}$. This way, for skewed distributions sufficient amount of data will be binned in the intervals for large x to see what the distribution looks like.

Binning in more detail in the next lecture.

Random Number Generators

RNGs

Most computational methods use random numbers, for example Monte Carlo, molecular dynamics, stochastic optimization and cryptography. And, of course, all machine learning rests upon massive amounts of random numbers.

Deterministic algorithms. (Pseudorandom numbers.) Pseudorandom number generators (RNG's): Deterministic algorithms that mimic randomness → Generated numbers are only "pseudo-random" but approximate real random numbers reasonably well.

In what follows, the term "random number" means pseudorandom number.

RNGs

Most RNG algorithms produce pseudorandom uniformly distributed numbers $x_i \in (0,1], i = 1, 2, ..., n$. Roughly, $X \sim U(0,1)$

These uniformly distributed random numbers can be used further to produce different distributions of random numbers, in other words to *simulate* different distributions.

RNGs

A good RNG has the following *properties*:

The numbers have a *correct distribution*:

- in simulations, the sequence of random numbers must be uncorrelated
- in numerical integration, it is important that the distribution is flat (uniform)
- The sequence must have a *long period*:
- all RNGs repeat the same sequence of numbers eventually, but the sequence must be sufficiently long
- The sequences should be *reproducible*:
- for testing of simulation parameters

- for stopping a simulation and continuing later Starting from the same seed number gives you the same sequence \rightarrow store the seed, when testing.

The RNG must be *fast*. Simulations need loads of them.

Linear congruential generators (LCG)

LCGs are based on the **integer recursion** relation

 $x_{i+1} = (ax_i + b) \mod m$ AND to scale to (0,1): x_{i+1}/m .

Here, integers *a*, *b*, and *m* are constants.

LCG generates a sequence $x_1, x_2, ...$ of random integers that are distributed within the intervals

- [0, m 1] (if b > 0) or
- [1, m 1] (if b = 0). (You need to start with $x_0 \neq 0$ to get $x_i \neq 0$.)

Scaling: divide by *m* for the interval

- [0,1) (b > 0) or
- -(0,1)(b=0).

Classification:

- mixed (b > 0): LCG(a, b, m)

- multiplicative (b = 0): MLCG(a, m)

Parameter *m* determines the *period P* of the LCG (it is usually close to the largest integer of the computer).

A seed x_0 is needed as input.

(Zero values are often eliminated from RNGs.)

LCGs

Two standard LCGs:

GGL

MLCG(16807, 2³¹-1): $x_{i+1} = (16807x_i) \mod (2^{31} - 1)$

Available in some numerical software packages such as subroutine RAND in Matlab.

Simple and fast, but suffers from a short period of 2^{31} - 1 $\approx 2 \times 10^9$ steps.

Problems due to correlation.

LCGs

RAND

LCG(69069, 1, 2^{32}): $x_{i+1} = (69069x_i + 1) \mod (2^{32})$ Also has problems due to correlations.

Visual Test

LCGs have a serious drawback of correlations between consecutive numbers $x_{i+1}, x_{i+2}, ..., x_{i+d}$ in the sequence.

In *d*-dimensional space, the points given by these *d* numbers order on parallel **hyperplanes**. The average distance between these planes, **whose dimension is** *d* – 1, is constant. The smaller the number of these planes, the less uniform is the distribution.

Lagged Fibonacci Generators

These are generalisations of LCGs: The period of a LCG can be increased by the form

$$x_i = (a_1 x_{i-1} + a_2 x_{i-2} + \ldots + a_p x_{i-p}) \mod m$$

where p > 1 and $a_p \neq 0$.

An LF generator requires an initial set of elements $x_1, x_2, ..., x_r$ and then uses the integer recursion

$$x_i = (x_{i-r} \otimes x_{i-s}) \mod m$$

where *r* and *s* are two integer lags satisfying r > sand \otimes is one of the following binary operations: +, -, ×, or \oplus (exclusive-or). (To clarify: binary operations, x_i 's are integers.)

LFGs

The corresponding generators are termed $LF(r, s, \otimes)$.

The initialization requires a set of q random numbers that can be generated for example by using another RNG.

The properties of LF-generators are not very well known but a definite plus is **long period**. Some evidence suggests that the exclusive-or operation should not be used.

LFGs

RAN3

LF(55,24,-):
$$x_i = (x_{i-55} - x_{i-24}) \mod m$$

 $m = 2^{32}$

- also called a subtractive method
- period $2^{55} 1$
- initialisation requires 55 numbers
- does not suffer from similar correlations as LCGs

(Assignment 1.)

These can be viewed as the special case m = 2 of LF generators.

Feedback shift register algorithms are based on the theory of **primitive trinomials**

$$P(x; p, q) = x^{p} + x^{q} + 1 \qquad (x = 0, 1)$$

Given such a primitive trinomial and *p* initial binary digits, a sequence of bits $b = \{bi\}$ (i = 0, 1, 2, ...) can be generated using the following **recursion formula**:

$$b_i = b_{i-p} \oplus b_{i-p+q}$$

where p > q.

Using the recursion formula, random words W_i of size *i* can be formed by

$$W_i = b_i b_{i+d} b_{i+2d} \cdots b_{i+(l-1)d}$$

where *d* is a chosen *delay*.

The resulting binary vectors are treated as random numbers.

It can be shown that if *p* is a *Mersenne prime*, which means that $2^p - 1$ is also a prime, then the sequence of random numbers has a maximal possible period of $2^p - 1$.

If interested, see

https://en.wikipedia.org/wiki/Linear-feedback_shift_register

In generalized feedback shift register (GFSR) generators, *i*-bit words are formed by a recursion where two bit sequences are combined using the binary operation \oplus :

$$W_i = W_{i-p} \oplus W_{i-q}$$

The best choices for q and p are Mersenne primes, which satisfy the condition $p^2 + q^2 + 1 = prime$.

Examples of pairs that satisfy this condition:

p = 98	q = 27
p = 250	q = 103
p = 1279	q = 216, 418
p = 9689	q = 84, 471, 1836, 2444, 418

Generalized feedback shift register generators are denoted by $GFSR(p, q, \oplus)$.

R250

R250 for which p = 250 and q = 103 has been the most commonly used generator of this class.

The 32-bit integers (32-bit words) are generated by

$$x_i = x_{i-250} \oplus x_{i-103}$$

250 uncorrelated seeds (random integers) are needed to initialize R250.

The latest 250 random numbers must be stored in memory.

The period length is 2^{250} - 1.

R250 does not exhibit similar pair correlations as the LCG generators.

However, R250 has strong triple correlations:

$$\langle x_i \ x_{i-250} \ x_{i-103} \rangle \neq 0$$

In addition, R250 fails in some important physical applications such as random walks and simulations of the Ising model.

An efficient way of reducing correlations is to decimate the sequence by taking only every *k*th number (k = 3, 5, 7, ...).

It seems natural that shuffling a sequence or combining two separate sequences might help in reducing correlations.

The **combination sequence** z_i is defined by $z_i = x_i \otimes y_i$

where x_i and y_i are from some (good) generators and \otimes denotes a binary operation (+, -, ×, \oplus).

RANMAR

- the best known and tested combination generator

The first RNG is a lagged Fibonacci generator

$$x_{i} = \begin{cases} x_{i-97} - x_{i-33} & \text{if } x_{i-97} \ge x_{i-33} \\ x_{i-97} - x_{i-33} + 1 & \text{otherwise} \end{cases}$$

Only 24 most significant bits are used for single precision reals. **The second part** of the generator is a simple arithmetic sequence for the prime modulus $2^{24} - 3 = 16777213$.

The sequence is defined as

$$y_i = \begin{cases} y_i - c & \text{if } y_i \ge c & c = 7654321/16777216, \\ y_i - c + d & \text{otherwise} & d = 16777213/16777216 \end{cases}$$

The final random number z_i is produced by combining x_i and y_i :

$$z_{i} = \begin{cases} x_{i} - y_{i} & \text{if } x_{i} \ge y_{i} \\ x_{i} - y_{i} + 1 & \text{otherwise} \end{cases}$$

The total period of RANMAR is about 2¹⁴⁴.

The code is available in the lectures in MyCourses.

RANMAR is first initialized by the call (in C) crmarin(seed);

Here, seed is an integer seed.

The call

cranmar(crn,len);

fills the vector crn of length len with uniformly distributed random numbers.

RANMAR is a very fast generator.

RANMAR has also performed well in several tests, and should thus be suitable for most applications.

RNG Tests

No single test can prove that a RNG is suitable for all applications

It is always possible to construct a test where a given RNG fails (since the numbers are not truly random but generated by a deterministic algorithm).

Classification of test methods

1. Theoretical tests

- based on theoretical properties of algorithms
- exact but often very difficult to perform
- only *asymptotic*: important correlations between consecutive number sets are not measured
RNG Tests

2. Empirical tests

- based on testing algorithms and their implementations in practice
- can be tailored to measure particular correlations
- suitable for all algorithms
- often difficult to say how much testing is sufficient
- further division into *standard tests* (statistical tests) and *application specific tests* (physical quantities)

3. Visual tests

- can be used to locate global or local deviations from randomness
- e.g. pairs of random numbers can be used to plot points in a unit square Do visual tests when you can!

Mersenne Twister RNG

In large simulations currently the best and computationally heaviest RNG is the **Mersenne Twister**, see: https://en.wikipedia.org/wiki/Mersenne_Twister

Python uses the Mersenne Twister as the core generator. It produces 53-bit precision floats and has a period of 2**19937-1. The underlying implementation is in C.

In Python you invoke the Mersenne-Twister RNG included in module random (**import random**) by random.random(). See the link: https://github.com/james727/MTP

Example of a simple test

The *moment test* is a simple procedure to check that your RNG implementation works as it should.

The moments of the uniform distribution are known

$$\langle x^k \rangle = \frac{1}{k+1}$$

If we generate random numbers that should be uniformly distributed, the moments calculated from these numbers should be approximately equal to the analytical values *within statistical fluctuations*.

Example: the mean value of random numbers for 100 independent measurements over N = 100 and N = 1000 random numbers using RANMAR.



The 'measured' values fluctuate around the correct value 0.5. **Error goes as** $1/\sqrt{N}$ **for uncorrelated random numbers** (idealisation). **Note**: This is a way to detect correlations.

Central limit theorem

For any independently measured values $M_1, M_2, ..., Mm$ that come from the same (sufficiently short-ranged) distribution p(x), the average $\frac{1}{1} \sum_{m=1}^{m} \frac{m}{1}$

$$\langle M \rangle = \frac{1}{m} \sum_{i=1}^{m} M_i$$

will asymptotically follow a Gaussian distribution (normal distribution), whose mean is $\langle M \rangle$ (equal to the mean of the parent distribution p(x)) and standard deviation is $1/\sqrt{N}$ times the standard deviation of p(x).

We can use this result to analyse the errors in the calculated values of any of the moments.

Using Random Numbers Example

Denote the second moment $M = \langle x^2 \rangle$

The errors should follow the normal distribution and the width of this distribution should behave as $1/\sqrt{N}$

Let's take a set of *m* independent 'measurements' of the second moment, each consisting of an average obtained from *N* random numbers.

From each measurement we obtain a single value M_{α} .

The average of all *m* measurements is $\langle M \rangle = \frac{1}{m} \sum_{\alpha=1}^{m} M_{\alpha}$ and the variance is given by $\sigma^2 = \langle M^2 \rangle - \langle M \rangle^2$

Here we have the variance of $M = \langle x^2 \rangle$ obtained from m = 1000 measurements, each consisting of an average from *N* random numbers.



The variance behaves like $1/\sqrt{N}$, meaning that the second moment obeys the scaling of the central limit theorem. U Here, uniformly distributed random numbers from RANMAR.

Next time:

Sampling from different distributions Logarithmic binning