UQ, Semester 1, 2017, Companion to STAT2201/CIVL2530 Exam Formulae and Tables

To be provided to students with STAT2201 or CIVIL-2530 (Probability and Statistics) Exam.

Main exam date: Tuesday, 20 June.

Probability and Monte Carlo

- > An experiment that can result in different outcomes, even though it is repeated in the same manner every time, is called a **random experiment**.
- > The set of all possible outcomes of a random experiment is called the **sample space** of the experiment, and is denoted as Ω .
 - A sample space is **discrete** if it consists of a finite or countably infinite set of outcomes.
 - A sample space is **continuous** if it contains an interval (either finite or infinite) of real numbers, vectors or similar objects.
- \succ An event is a subset of the sample space of a random experiment.
 - The union of two events is the event that consists of all outcomes that are contained in either of the two events or both. We denote the union as $E_1 \cup E_2$.
 - The intersection of two events is the event that consists of all outcomes that are contained in both of the two events. We denote the intersection as $E_1 \cap E_2$.
 - The **complement** of an event in the sample space is the set of outcomes in the sample space that are not in the event. We denote the complement of the event E as \overline{E} . The notation E^c is also used. Note that $E \cup \overline{E} = \Omega$.
- > Two events, denoted E_1 and E_2 are **mutually exclusive** if: $E_1 \cap E_2 = \emptyset$ where \emptyset is called the **empty set** or **null event**.
- > A collection of events, E_1, E_2, \ldots, E_k is said to be **mutually exclusive** if for all pairs,

$$E_i \cap E_j = \emptyset$$

- > The definition of the complement of an event implies that: $(E^c)^c = E$.
- \succ The distributive law for set operations implies that

 $(A \cup B) \cap C = (A \cap C) \cup (B \cap C)$ and $(A \cap B) \cup C = (A \cup C) \cap (B \cup C).$

 \succ DeMorgan's laws imply that

$$(A \cup B)^c = A^c \cap B^c$$
 and $(A \cap B)^c = A^c \cup B^c$.

- ➤ Union and intersection are commutative operations: $A \cap B = B \cap A$ and $A \cup B = B \cup A$.
- Probability is used to quantify the likelihood, or chance, that an outcome of a random experiment will occur.
- > Whenever a sample space consists of a finite number N of possible outcomes, each equally likely, the probability of each outcome is 1/N.
- > For a discrete sample space, the **probability of an event** E, denoted as P(E), equals the sum of the probabilities of the outcomes in E.
- > If Ω is the sample space and E is any event in a random experiment,
 - (1) $P(\Omega) = 1.$
 - (2) $0 \le P(E) \le 1.$
 - (3) For two events E_1 and E_2 with $E_1 \cap E_2 = \emptyset$ (disjoint), $P(E_1 \cup E_2) = P(E_1) + P(E_2).$

- (4) $P(E^c) = 1 P(E)$.
- (5) $P(\emptyset) = 0.$

> The probability of event A or event B occurring is,

$$P(A \cup B) = P(A) + P(B) - P(A \cap B).$$

> If A and B are mutually exclusive events,

$$P(A \cup B) = P(A) + P(B).$$

► For a collection of **mutually exclusive events**,

$$P(E_1 \cup E_2 \cup \dots \cup E_k) = P(E_1) + P(E_2) + \dots + P(E_k).$$

- > The probability of an event B under the knowledge that the outcome will be in event A is denoted $P(B \mid A)$ and is called the **conditional probability** of B given A.
- > The conditional probability of an event B given an event A, denoted as $P(B \mid A)$, is

$$P(B \mid A) = \frac{P(A \cap B)}{P(A)} \quad \text{for} \quad P(A) > 0$$

- ➤ The multiplication rule for probabilities is: $P(A \cap B) = P(B \mid A)P(A) = P(A \mid B)P(B)$.
- > For an event B and a collection of mutual exclusive events, E_1, E_2, \ldots, E_k where their union is Ω . The **law of total probability** yields,

$$P(B) = P(B \cap E_1) + P(B \cap E_2) + \dots + P(B \cap E_k)$$

= $P(B \mid E_1)P(E_1) + P(B \mid E_2)P(E_2) + \dots + P(B \mid E_k)P(E_k).$

 \succ Two events A and B are **independent** if any one of the following equivalent statements is true:

- (1) $P(A \mid B) = P(A).$
- (2) $P(B \mid A) = P(B).$
- (3) $P(A \cap B) = P(A)P(B)$.

Observe that **independent** events and **mutually exclusive** events, are completely different concepts. Don't confuse these concepts.

 \succ For multiple events E_1, E_2, \ldots, E_n are independent if and only if for any subset of these events

$$P(E_{i_1} \cap E_{i_2} \cap \dots \cap E_{i_k}) = P(E_{i_1}) P(E_{i_2}) \dots P(E_{i_k})$$

➤ A pseudorandom sequence is a sequence of numbers U_1, U_2, \ldots with each number, U_k depending on the previous numbers $U_{k-1}, U_{k-2}, \ldots, U_1$ through a well defined functional relationship and similarly U_1 depending on the seed \tilde{U}_0 . Hence for any seed, \tilde{U}_0 , the resulting sequence U_1, U_2, \ldots is fully defined and repeatable. A pseudorandom sequence often lives within a discrete domain as $\{0, 1, \ldots, 2^{64} - 1\}$. It can then be normalised to floating point numbers with,

$$R_k = \frac{U_k}{2^{64} - 1}.$$

- \succ A good pseudorandom sequence has the following attributes among others:
 - 1. It is quick and easy to compute the next element in the sequence.
 - 2. The sequence of numbers R_1, R_2, \ldots resembles properties as an i.i.d. sequence of uniform(0,1) random variables (i.i.d. is defined in Unit 4).
- Computer simulation of random experiments is called Monte Carlo and is typically carried out by setting the seed to either a reproducible value or an arbitrary value such as system time.
- \succ Random experiments may be replicated on a computer using Monte Carlo simulation.

Distributions

- > A random variable X is a numerical (integer, real, complex, vector, etc.) summary of the outcome of the random experiment. The range or support of the random variable is the set of possible values that it may take. Random variables are usually denoted by capital letters.
- A discrete random variable is an integer/real-valued random variable with a finite (or countably infinite) range.
- A continuous random variable is a real-valued random variable with an interval (either finite or infinite) of real numbers for its range.
- > The **probability distribution** of a random variable X is a description of the probabilities associated with the possible values of X. There are several common alternative ways to describe the probability distribution, with some differences between discrete and continuous random variables.
- > While not the most popular in practice, a unified way to describe the distribution of any scalar valued random variable X (real or integer) is the **cumulative distribution function**,

$$F(x) = P(X \le x).$$

- \succ It holds that
 - (1) $0 \le F(x) \le 1$.
 - (2) $\lim_{x \to -\infty} F(x) = 0.$
 - (3) $\lim_{x\to\infty} F(x) = 1.$
 - (4) If $x \leq y$, then $F(x) \leq F(y)$. That is, $F(\cdot)$ is non-decreasing.
- > Distributions are often summarised by numbers such as the mean, μ , variance, σ^2 , or moments. These numbers, in general do not identify the distribution, but hint at the general location, spread and shape.
- > The standard deviation of X is $\sigma = \sqrt{\sigma^2}$ and is particularly useful when working with the Normal distribution.
- > Given a discrete random variable X with possible values x_1, x_2, \ldots, x_n , the **probability mass** function of X is,

$$p(x) = P(X = x).$$

Note: In [MonRun2014] and many other sources, the notation used is f(x) (as a pdf of a continuous random variable).

> A probability mass function, p(x) satisfies:

(1)
$$p(x_i) \ge 0.$$

(2) $\sum_{i=1}^{n} p(x_i) = 1$

> The cumulative distribution function of a discrete random variable X, denoted as F(x), is

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

> $P(X = x_i)$ can be determined from the *jump* at the value of x. More specifically

$$p(x_i) = P(X = x_i) = F(x_i) - \lim_{x \uparrow x_i} F(x_i).$$

 \succ The mean or expected value of a discrete random variable X, is

$$\mu = E(X) = \sum_{x} x \, p(x).$$

➤ The expected value of h(X) for some function $h(\cdot)$ is:

$$E[h(X)] = \sum_{x} h(x) p(x).$$

> The k'th moment of X is,

$$E(X^k) = \sum_x x^k \, p(x).$$

 \succ The variance of X, is

$$\sigma^2 = V(X) = E((X - \mu)^2) = \sum_x (x - \mu)^2 p(x) = \sum_x x^2 p(x) - \mu^2$$

> A random variable X has a **discrete uniform distribution** if each of the n values in its range, x_1, x_2, \ldots, x_n , has equal probability. I.e.

$$p(x_i) = 1/n.$$

Suppose that X is a discrete uniform random variable on the consecutive integers $a, a + 1, a + 2, \ldots, b$, for $a \le b$. The **mean** and **variance** of X are

$$E(X) = \frac{b+a}{2}$$
 and $V(X) = \frac{(b-a+1)^2 - 1}{12}$.

 \succ The setting of *n* independent and identical Bernoulli trials is as follows:

- (1) There are n trials.
- (1) The trials are independent.
- (2) Each trial results in only two possible outcomes, labelled as "success" and "failure".
- (3) The probability of a success in each trial denoted as p is the same for all trials.
- The random variable X that equals the number of trials that result in a success is a **binomial** random variable with parameters $0 \le p \le 1$ and n = 1, 2, ... The probability mass function of X is

$$p(x) = {\binom{n}{x}} p^x (1-p)^{n-x}, \qquad x = 0, 1, \dots, n$$

 \succ Useful to remember from algebra: the binomial expansion for constants a and b is

$$(a+b)^n = \sum_{k=0}^n \binom{n}{k} a^k b^{n-k}.$$

> If X is a binomial random variable with parameters p and n, then,

$$E(X) = n p$$
 and $V(X) = n p (1-p)$.

> Given a continuous random variable X, the **probability density function** (pdf) is a function, f(x) such that,

- (1) $f(x) \ge 0.$
- (2) f(x) = 0 for x not in the range.
- (3) $\int_{-\infty}^{\infty} f(x) \, dx = 1.$
- (4) For small Δx , $f(x) \Delta x \approx P(X \in [x, x + \Delta x))$.
- (5) $P(a \le X \le b) = \int_{a}^{b} f(x)dx = \text{area under } f(x) \text{ from } a \text{ to } b.$

> Given the pdf, f(x) we can get the cdf as follows:

$$F(x) = P(X \le x) = \int_{-\infty}^{x} f(u)du \quad \text{for} \quad -\infty < x < \infty.$$

> Given the cdf of a continuous random variable, F(x) we can get the pdf:

$$f(x) = \frac{d}{dx} F(x).$$

 \succ The mean or expected value of a continuous random variable X, is

$$\mu = E(X) = \int_{-\infty}^{\infty} x f(x) dx.$$

> The **expected value** of h(X) for some function $h(\cdot)$ is:

$$E\Big[h(X)\Big] = \int_{-\infty}^{\infty} h(x)f(x) \, dx.$$

 \succ The k'th **moment** of X is,

$$E(X^k) = \int_{-\infty}^{\infty} x^k f(x) \, dx$$

≻ The **variance** of X, is

$$\sigma^{2} = V(X) = E((X - \mu)^{2}) = \int_{-\infty}^{\infty} (x - \mu)^{2} f(x) dx = \int_{-\infty}^{\infty} x^{2} f(x) dx - \mu^{2}.$$

> A continuous random variable X with probability density function

$$f(x) = \frac{1}{b-a}, \qquad a \le x \le b.$$

is a **continuous uniform random variable** or "uniform random variable" for short.

> If X is a continuous uniform random variable over $a \le x \le b$, the **mean** and **variance** are:

$$\mu = E(X) = \frac{a+b}{2}$$
 and $\sigma^2 = V(X) = \frac{(b-a)^2}{12}$.

 \succ A random variable X with probability density function

$$f(x) = \frac{1}{\sigma\sqrt{2\pi}} e^{\frac{-(x-\mu)^2}{2\sigma^2}}, \qquad -\infty < x < \infty,$$

is a **normal random variable** with parameters μ where $-\infty < \mu < \infty$, and $\sigma > 0$. For this distribution, the parameters map directly to the mean and variance,

$$E(X) = \mu$$
 and $V(X) = \sigma^2$.

The notation $N(\mu, \sigma^2)$ is used to denote the distribution. Note that some authors and software packages use σ for the second parameter and not σ^2 .

 \succ A normal random variable with a mean and variance of:

$$\mu = 0$$
 and $\sigma^2 = 1$

is called a **standard normal random variable** and is denoted as Z. The cumulative distribution function of a standard normal random variable is denoted as

$$\Phi(z) = F_Z(z) = P(Z \le z),$$

and is tabulated.

► It is very common to compute
$$P(a < X < b)$$
 for $X \sim N(\mu, \sigma^2)$. This is the typical way:

$$P(a < X < b) = P(a - \mu < X - \mu < b - \mu)$$

= $P\left(\frac{a - \mu}{\sigma} < \frac{X - \mu}{\sigma} < \frac{b - \mu}{\sigma}\right)$
= $P\left(\frac{a - \mu}{\sigma} < Z < \frac{b - \mu}{\sigma}\right)$
= $\Phi\left(\frac{b - \mu}{\sigma}\right) - \Phi\left(\frac{a - \mu}{\sigma}\right).$

We get:

$$F_X(b) - F_X(a) = F_Z\left(\frac{b-\mu}{\sigma}\right) - F_Z\left(\frac{a-\mu}{\sigma}\right)$$

> The exponential distribution with parameter $\lambda > 0$ is given by the survival function,

$$\overline{F}(x) = 1 - F(x) = P(X > x) = e^{-\lambda x}.$$

- > The random variable X that equals the distance between successive events from a Poisson process with mean number of events per unit interval $\lambda > 0$.
- \succ The probability density function of X is

$$f(x) = \lambda e^{-\lambda x}$$
 for $0 \le x < \infty$.

Note that sometimes a different parameterisation, $\theta = 1/\lambda$ is used (e.g. in the Julia Distributions package).

 \succ The mean and variance are:

$$\mu = E(X) = \frac{1}{\lambda} \quad \text{ and } \quad \sigma^2 = V(X) = \frac{1}{\lambda^2}.$$

> The exponential distribution is the only continuous distribution with range $[0, \infty)$ exhibiting the **lack of memory property**. For an exponential random variable X,

$$P(X > t + s \,|\, X > t) = P(X > s).$$

[➤] Monte Carlo simulation makes use of methods to transform a uniform random variable in a manner where it follows an arbitrary given distribution. One example of this is if $U \sim \text{Uniform}(0,1)$ then $X = -\frac{1}{\lambda} \log(U)$ is exponentially distributed with parameter λ .

Joint Probability Distributions

- A joint probability distribution of two random variables is also referred to as a **bivariate probability distribution**.
- > A joint probability mass function for discrete random variables X and Y, denoted as $p_{XY}(x, y)$, satisfies the following properties:
 - (1) $p_{XY}(x,y) \ge 0$ for all x, y.
 - (2) $p_{XY}(x, y) = 0$ for (x, y) not in the range.
 - (3) $\sum \sum p_{XY}(x,y) = 1$, where the summation is over all (x,y) in the range.
 - (4) $p_{XY}(x,y) = P(X = x, Y = y).$
- > A joint probability density function for continuous random variables X and Y, denoted as $f_{XY}(x, y)$, satisfies the following properties:
 - (1) $f_{XY}(x,y) \ge 0$ for all x, y.
 - (2) $f_{XY}(x,y) = 0$ for (x,y) not in the range.
 - (3) $\int_{-\infty}^{\infty} \int_{-\infty}^{\infty} f_{XY}(x,y) \ dx \ dy = 1.$

(4) For small
$$\Delta x$$
, Δy : $f_{XY}(x,y) \Delta x \Delta y \approx P((X,Y) \in [x, x + \Delta x) \times [y, y + \Delta y))$

(5) For any region R of two-dimensional space,

$$P((X,Y) \in R) = \iint_R f_{XY}(x,y) \ dx \ dy.$$

- > A joint probability density function can also be defined for n > 2 random variables (as can be a joint probability mass function). The following needs to hold:
 - (1) $f_{X_1X_2...X_n}(x_1, x_2, ..., x_n) \ge 0.$
 - (2) $\int_{-\infty}^{\infty} \int_{-\infty}^{\infty} \dots \int_{-\infty}^{\infty} f_{X_1 X_2 \dots X_n}(x_1, x_2, \dots, x_n) dx_1 dx_2 \dots dx_n = 1.$
- > Most of the concepts in this section, carry over from bivariate to general multivariate distributions (n > 2).
- > The marginal distributions of X and Y as well as the conditional distribution of X given a specific value Y = y and vice versa can be obtained from the joint distribution.
- > If the random variables X and Y are independent, then $f_{XY}(x, y) = f_X(x) f_Y(y)$ and similarly in the discrete case.
- \succ The expected value of a function of two random variables is:

$$E[h(X,Y)] = \iint h(x,y)f_{XY}(x,y) \ dx \ dy \qquad \text{for } X,Y \text{ continuous.}$$

> The covariance is a common measure of the relationship between two random variables (say X and Y). It is denoted as cov(X, Y) or σ_{XY} , and is given by:

$$\sigma_{XY} = E\Big[(X - \mu_X)(Y - \mu_Y)\Big] = E(XY) - \mu_X\mu_Y.$$

 \succ The covariance of a random variable with itself is its variance.

> The correlation between the random variables X and Y, denoted as ρ_{XY} , is

$$\rho_{XY} = \frac{\operatorname{cov}(X, Y)}{\sqrt{V(X)V(Y)}} = \frac{\sigma_{XY}}{\sigma_X \sigma_Y}.$$

- ► For any two random variables X and Y, $-1 \le \rho_{XY} \le 1$.
- > If X and Y are independent random variables, $\sigma_{XY} = 0$ and $\rho_{XY} = 0$. The opposite case does not always hold: In general $\rho_{XY} = 0$ does not imply independence. But for jointly Normal random variables it does. In any case, if $\rho_{XY} = 0$ then the random variables are called uncorrelated.
- > When considering several random variables, it is common to consider the (symmetric) Covariance Matrix, Σ with $\Sigma_{i,j} = \operatorname{cov}(X_i, X_j)$.
- \succ The probability density function of a bivariate normal distribution is

$$f_{XY}(x, y; \sigma_X, \sigma_Y, \mu_X, \mu_Y, \rho) = \frac{1}{2\pi\sigma_X\sigma_Y\sqrt{1-\rho^2}} \\ \times \exp\left\{\frac{-1}{2(1-\rho^2)} \left[\frac{(x-\mu_X)^2}{\sigma_X^2} - \frac{2\rho(x-\mu_X)(y-\mu_Y)}{\sigma_X\sigma_Y} + \frac{(y-\mu_Y)^2}{\sigma_Y^2}\right]\right\}$$

for $-\infty < x < \infty$ and $-\infty < y < \infty$, with parameters $\sigma_X > 0$, $\sigma_Y > 0$, $-\infty < \mu_X < \infty$, $-\infty < \mu_Y < \infty$, and $-1 < \rho < 1$.

> Given random variables X_1, X_2, \ldots, X_n and constants c_1, c_2, \ldots, c_n , the (scalar) linear combination (with possible affine term b),

$$Y = c_1 X_1 + c_2 X_2 + \dots + c_n X_n + b$$

is often a random variable of interest.

 \succ The mean of the linear combination is the linear combination of the means,

$$E(Y) = c_1 E(X_1) + c_2 E(X_2) + \dots + c_n E(X_n) + b$$

This holds even if the random variables are not independent.

 \succ The variance of the linear combination is as follows:

$$V(Y) = c_1^2 V(X_1) + c_2^2 V(X_2) + \dots + c_n^2 V(X_n) + 2 \sum_{i < j} \sum c_i c_j \operatorname{cov}(X_i, X_j).$$

> If X_1, X_2, \ldots, X_n are **independent** (or even if they are just uncorrelated).

$$V(Y) = c_1^2 V(X_1) + c_2^2 V(X_2) + \dots + c_n^2 V(X_n).$$

- > In case the random variables X_1, \ldots, X_n were jointly Normal then, $Y \sim \text{Normal}(E(Y), V(Y))$. That is, **linear combinations of Normal random variables remain Normally distributed**.
- > A collection of random variables, X_1, \ldots, X_n is said to be **i.i.d.**, or **independent and identically distributed** if they are mutually independent and identically distributed. This means that the (*n* - dimensional) joint probability density is a product of the individual densities.
- > In the context of statistics, a **random sample** is often modelled as an i.i.d. vector of random variables. X_1, \ldots, X_n .
- \succ An important linear combination associated with a random sample is the **sample mean**:

$$\overline{X} = \frac{\sum_{i=1}^{n} X_i}{n} = \frac{1}{n} X_1 + \frac{1}{n} X_2 + \dots + \frac{1}{n} X_n.$$

➤ If X_i has mean μ and variance σ^2 then sample mean (of an i.i.d. sample) has,

$$E(\overline{X}) = \mu, \qquad V(\overline{X}) = \frac{\sigma^2}{n}.$$

Descriptive Statistics

- Descriptive statistics deals with summarizing data using numbers, qualitative summaries, tables and graphs.
- \succ Here are some types of **data configurations**:
 - 1. Single sample: x_1, x_2, \ldots, x_n .
 - 2. Single sample over time (time series): $x_{t_1}, x_{t_2}, \ldots, x_{t_n}$ with $t_1 < t_2 < \ldots < t_n$.
 - 3. Two samples: x_1, \ldots, x_n and y_1, \ldots, y_m .
 - 4. Generalizations from two samples to k samples (each of potentially different sample size, n_1, \ldots, n_k).
 - 5. Observations in tuples: $(x_1, y_1), (x_2, y_2), \dots, (x_n, y_n)$.
 - 6. Generalizations from tuples to vector observations (each vector of length ℓ),

$$(x_1^1,\ldots,x_1^\ell),\ldots,(x_n^1,\ldots,x_n^\ell).$$

- Individual variables may be categorical or numerical. Categorical variables (taking values in one of several categories) may be ordinal meaning that they can be sorted (e.g. "low", "moderate", "high"), or not (e.g. "cat", "dog", "fish").
- > A statistic is a quantity computed from a sample (assume here a single sample x_1, \ldots, x_n). Here are very common and useful statistics:
 - 1. The sample mean: $\bar{x} = \frac{x_1 + \dots + x_n}{n} = \frac{\sum_{i=1}^n x_i}{n}.$ 2. The sample variance: $s^2 = \frac{\sum_{i=1}^n (x_i - \bar{x})^2}{n-1} = \frac{\sum_{i=1}^n x_i^2 - n \, \bar{x}^2}{n-1}.$
 - 3. The sample standard deviation: $s = \sqrt{s^2}$.
 - 4. Order statistics work as follows: Sort the sample to obtain the sequence of sorted observations, denoted $x_{(1)}, \ldots, x_{(n)}$ where, $x_{(1)} \leq x_{(2)} \leq \ldots \leq x_{(n)}$. Some common order statistics:
 - (a) The **minimum** $\min(x_1, ..., x_n) = x_{(1)}$.
 - (b) The **maximum** $\max(x_1, ..., x_n) = x_{(n)}$.
 - (c) The median

median =
$$\begin{cases} x_{(\frac{n+1}{2})} & \text{if } n \text{ is odd,} \\ \frac{1}{2} \left(x_{(\frac{n}{2})} + x_{(\frac{n}{2}+1)} \right) & \text{if } n \text{ is even.} \end{cases}$$

Note that the median is the 50'th percentile and the 2nd quartile (see below).

- (d) The q th quantile $(q \in [0, 1])$ or alternatively the p = 100q percentile (measured in percents instead of a decimal), is the observation such that p percent of the observations are less than it and (1-p) percent of the observations are greater than it. In cases (as is typical) that there is not such a precise observation, it is a linear interpolation between two neighbouring observations (as is done for the median when n is even). In terms of order statistics, the q th quantile is approximately (not taking linear interpolations into account) $x_{([q*n])}$. Here [z] denotes the nearest integer in $\{1, \ldots, n\}$ to z.
- (e) The first **quartile**, denoted Q1 is the 25th percentile. The second quartile (Q2) is the median. The **third quartile**, denoted Q3 is the 75th percentile. Thus half of the observations lie between Q1 and Q3. In other words, the quartiles break the sample into 4 quarters. The difference Q3 Q1 is the **interquartile range**.
- (f) The sample range is $x_{(n)} x_{(1)}$.

\succ Constructing a Histogram (Equal Bin Widths)

- (1) Label the bin (class interval) boundaries on a horizontal scale.
- (2) Mark and label the vertical scale with **frequencies** or **counts**.
- (3) Above each bin, draw a rectangle where height is equal to the frequency (or count).
- > A Kernel Density Estimate (KDE) is a way to construct a Smoothed Histogram. While construction is not as straightforward as steps (1)-(3) above, automated tools can be used.
- ➤ Both the histogram and the KDE are not unique in the way they summarize data. With these methods, different settings (e.g. number of bins in histograms or **bandwidth** in a KDE) may yield different representations of the same data set. Nevertheless, they are both very common, sensible and useful visualisations of data.
- ➤ The box plot is a graphical display that simultaneously describes several important features of a data set, such as centre, spread, departure from symmetry, and identification of unusual observations or outliers. It is often common to plot several box plots next to each other for comparison.
- > An anachronistic, but useful way for summarising small data-sets is the stem and leaf diagram.
- ➤ In a **cumulative frequency plot** the height of each bar is the total number of observations that are less than or equal to the upper limit of the bin.
- ➤ The Empirical Cumulative Distribution Function (ECDF) is,

$$\hat{F}(x) = \frac{1}{n} \sum_{i=1}^{n} \mathbf{1}\{x_i \le x\}$$

Here $\mathbf{1}\{\cdot\}$ is the **indicator function**. The ECDF is a function of the data, defined for all x.

- ➤ Given a candidate distribution with cdf F(x), a probability plot is a plot of the ECDF (or sometimes just it's jump points) with the y-axis stretched by the inverse of the cdf $F^{-1}(\cdot)$. The monotonic transformation of the y-axis is such that if the data comes from the candidate F(x), the points would appear to lie on a straight line. Names of variations of probability plots are the **P-P plot** and **Q-Q plot** (these plots are similar to the probability plot). A very common probability plot is the **Normal probability plot** where the candidate distribution is taken to be Normal(\overline{x}, s^2).
- ➤ The Normal probability plot can be useful in identifying distributions that are symmetric but that have tails that are "heavier" or "lighter" than the Normal.
- ➤ A time series plot is a graph in which the vertical axis denotes the observed value of the variable and the horizontal axis denotes time.
- A scatter diagram is constructed by plotting each pair of observations with one measurement in the pair on the vertical axis of the graph and the other measurement in the pair on the horizontal axis.
- > The sample correlation coefficient r_{xy} is an estimate for the correlation coefficient, ρ , presented in the previous unit:

$$r_{xy} = \frac{\sum_{i=1}^{n} (y_i - \bar{y})(x_i - \bar{x})}{\sqrt{\sum_{i=1}^{n} (y_i - \bar{y})^2 \sum_{i=1}^{n} (x_i - \bar{x})^2}}$$

Statistical Inference Ideas

- Statistical Inference is the process of forming judgements about the parameters of a population, typically on the basis of random sampling.
- > The random variables X_1, X_2, \ldots, X_n are an (i.i.d.) random sample of size n if
 - (a) the X_i 's are independent random variables and
 - (b) every X_i has the same probability distribution.
- ➤ A statistic is any function of the observations in a random sample, and the probability distribution of a statistic is called the sampling distribution.
- > Any function of the observation, or any statistic, is also a random variable. We call the probability distribution of a statistic a sampling distribution. A point estimate of some population parameter θ is a single numerical value $\hat{\theta}$ of a statistic $\hat{\Theta}$. The statistic $\hat{\Theta}$ is called the point estimator.
- > The most common statistic we consider is the sample mean, \overline{X} , with a given value denoted by \overline{x} . As an estimator, the sample mean is an estimator of the population mean, μ .
- > Central Limit Theorem (for sample means): If X_1, X_2, \ldots, X_n is a random sample of size *n* taken from a population with mean μ and finite variance σ^2 and if \overline{X} is the sample mean, the limiting form of the distribution of

$$Z = \frac{\overline{X} - \mu}{\sigma / \sqrt{n}}$$

as $n \to \infty$, is the standard normal distribution.

- > This implies that \overline{X} is approximately normally distributed with mean μ and standard deviation σ/\sqrt{n} .
- > The standard error of \overline{X} is given by σ/\sqrt{n} . In most practical situations σ is not known but rather estimated in this case, the estimated standard error, (denoted in typical computer output as "SE"), is s/\sqrt{n} where s is the point estimator,

$$s = \sqrt{\frac{\sum\limits_{i=1}^{n} x_i^2 - n \overline{x}^2}{n-1}}$$

> Central Limit Theorem (for sums): Manipulate the central limit theorem (for sample means and use $\sum_{i=1}^{n} X_i = n\overline{X}$. This yields,

$$Z = \frac{\sum_{i=1}^{n} X_i - n\,\mu}{\sqrt{n\sigma^2}}$$

which follows a standard normal distribution as $n \to \infty$.

- > This implies that $\sum_{i=1}^{n} X_i$ is approximately normally distributed with mean $n \mu$ and variance $n \sigma^2$.
- Knowing the sampling distribution (or the approximate sampling distribution) of a statistic is the key for the two main tools of statistical inference that we study:
 - (a) Confidence intervals a method for yielding error bounds on point estimates.
 - (b) **Hypothesis testing** a methodology for making conclusions about population parameters.

> The formulas for most of the statistical procedures use quantiles of the sampling distribution. When the distribution is N(0,1) (standard normal), the α quantile is denoted z_{α} and satisfies:

$$\alpha = \int_{-\infty}^{z_{\alpha}} \frac{1}{\sqrt{2\pi}} e^{\frac{-x^2}{2}} dx.$$

A common value to use for α is 0.05 and in procedures the expressions $z_{1-\alpha}$ or $z_{1-\alpha/2}$ appear. Note that in this case $z_{1-\alpha/2} = 1.96 \approx 2$.

> A confidence interval estimate for μ is an interval of the form $l \leq \mu \leq u$, where the end-points l and u are computed from the sample data. Because different samples will produce different values of l and u, these end points are values of random variables L and U, respectively. Suppose that

$$P(L \le \mu \le U) = 1 - \alpha.$$

The resulting **confidence interval** for μ is

 $l \leq \mu \leq u.$

The end-points or bounds l and u are called the **lower-** and **upper-confidence limits** (bounds), respectively, and $1 - \alpha$ is called the **confidence level**.

> If \bar{x} is the sample mean of a random sample of size *n* from a normal population with known variance σ^2 , a 100(1 - α)% confidence interval on μ is given by

$$\bar{x} - z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}} \le \mu \le \bar{x} + z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}}.$$

- > Note that it is roughly of the form, $\overline{x} 2 \operatorname{SE}(\overline{x}) \leq \mu \leq \overline{x} + 2 \operatorname{SE}(\overline{x})$.
- > Confidence interval formulas give insight into the **required sample size**: If \bar{x} is used as an estimate of μ , we can be $100(1 \alpha)\%$ confident that the error $|\bar{x} \mu|$ will not exceed a specified amount Δ when the sample size is not smaller than

$$n = \left(\frac{z_{1-\alpha/2} \ \sigma}{\Delta}\right)^2.$$

- > A statistical hypothesis is a statement about the parameters of one or more populations. The null hypothesis, denoted H_0 is the claim that is initially assumed to be true based on previous knowledge. The alternative hypothesis, denoted H_1 is a claim that contradicts the null hypothesis.
- > For some arbitrary value μ_0 , a two-sided alternative hypothesis would be expressed as follows:

$$H_0: \mu = \mu_0 \qquad H_1: \mu \neq \mu_0,$$

whereas a **one-sided alternative hypothesis** would be expressed as:

$$H_0: \mu = \mu_0$$
 $H_1: \mu < \mu_0$ or $H_0: \mu = \mu_0$ $H_1: \mu > \mu_0$.

- > The standard scientific research use of hypothesis is to "hope to reject" H_0 so as to have statistical evidence for the validity of H_1 .
- > An hypothesis test is based on a **decision rule** that is a function of the **test statistic**. For example: Reject H_0 if the test statistic is below a specified threshold, otherwise don't reject.

> Rejecting the null hypothesis H_0 when it is true is defined as a **type I error**. Failing to reject the null hypothesis H_0 when it is false is defined as a **type II error**.

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	H_0 Is True	H_0 Is False		
Fail to reject H_0 :	No error	Type II error		
Reject H_0 :	Type I error	No error		

 $\alpha = P(\text{type I error}) = P(\text{reject } H_0 \mid H_0 \text{ is true}).$

 $\beta = P(\text{type II error}) = P(\text{fail to reject } H_0 \mid H_0 \text{ is false}).$

- > The **power** of a statistical test is the probability of rejecting the null hypothesis H_0 when the alternative hypothesis is true.
- > A typical example of a simple hypothesis test has $H_0: \mu = \mu_0$ vs. $H_1: \mu = \mu_1$, where μ_0 and μ_1 are some specified values for the population mean. This test isn't typically practical but is useful for understanding the concepts at hand.
- > Assuming that $\mu_0 < \mu_1$ and setting a threshold, τ , reject H_0 if the $\overline{x} > \tau$, otherwise don't reject.
- > Explicit calculation of the relationships of τ , α , β , n, σ , μ_0 and μ_1 is possible in this case.
- > In most hypothesis tests used in practice (and in this course), a specified level of type I error, α is predetermined (e.g. $\alpha = 0.05$) and the type II error is not directly specified.
- > The probability of making a type II error β increases (power decreases) rapidly as the true value of μ approaches the hypothesized value.
- > The probability of making a type II error also depends on the sample size n increasing the sample size results in a decrease in the probability of a type II error.
- > The population (or natural) variability (e.g. described by σ) also affects the power.
- > The *P*-value is the smallest level of significance that would lead to rejection of the null hypothesis H_0 with the given data. That is, the *P*-value is based on the data. It is computed by considering the location of the test statistic under the sampling distribution based on H_0 . It can also be viewed as the probability of observing a set of data which is as consistent or more consistent with the alternative hypothesis than the observed data, when the null hypothesis is true.
- > It is customary to consider the test statistic (and the data) significant when the null hypothesis H_0 is rejected; therefore, we may think of the *P*-value as the smallest α at which the data are significant. In other words, the *P*-value is the **observed significance level**.
- > Clearly, the *P*-value provides a measure of the credibility of the null hypothesis. Computing the exact *P*-value for a statistical test is not always doable by hand.
- > It is typical to report the *P*-value in studies where H_0 was rejected (and new scientific claims were made). Typical ("convincing") values can be of the order 0.001.

≻ A General Procedure for Hypothesis Tests is

- (1) **Parameter of interest:** From the problem context, identify the parameter of interest.
- (2) Null hypothesis, H_0 : State the null hypothesis, H_0 .
- (3) Alternative hypothesis, H_1 : Specify an appropriate alternative hypothesis, H_1 .
- (4) Test statistic: Determine an appropriate test statistic.
- (5) **Reject** H_0 if: State the rejection criteria for the null hypothesis.
- (6) **Computations:** Compute any necessary sample quantities, substitute these into the equation for the test statistic, and compute the value.
- (7) **Draw conclusions:** Decide whether or not H_0 should be rejected and report that in the problem context.

Single Sample Inference

- > The setup is a sample x_1, \ldots, x_n (collected values) modelled by an i.i.d. sequence of random variables, X_1, \ldots, X_n .
- > The parameter at question in this unit is the population mean, $\mu = E[X_i]$. A point estimate is \overline{x} (described by the random variable \overline{X}).
- > We devise hypothesis tests and confidence intervals for μ , distinguishing between the (unrealistic but simpler) case where the population variance, σ^2 , is known, and the more realistic case where it is not known and estimated by the sample variance, s^2 .
- > For very small samples, the results we present are valid only if the population is normally distributed. But for non-small samples (e.g. n > 20, although there isn't a clear rule), the central limit theorem provides a good approximation and the results are approximately correct.

≻ Testing Hypotheses on the Mean, Variance Known (Z-Tests)

Model: $X_i \stackrel{i.i.d.}{\sim} N(\mu, \sigma^2)$ with μ unknown but σ^2 known.

Null hypothesis: $H_0: \mu = \mu_0.$

Test statistic: $z = \frac{\overline{x} - \mu_0}{\sigma/\sqrt{n}}, \qquad Z = \frac{\overline{X} - \mu_0}{\sigma/\sqrt{n}}.$

Alternative Hypotheses	P-value	Rejection Criterion for Fixed-Level Tests
$H_1: \mu \neq \mu_0$	$P = 2 \big[1 - \Phi \big(z \big) \big]$	$z > z_{1-\alpha/2}$ or $z < z_{\alpha/2}$
$H_1: \mu > \mu_0$	$P = 1 - \Phi(z)$	$z > z_{1-\alpha}$
$H_1: \mu < \mu_0$	$P = \Phi(z)$	$z < z_{\alpha}$

> Note: For $H_1: \mu \neq \mu_0$, a procedure identical to the preceding fixed significance level test is:

Reject H_0 : $\mu = \mu_0$	if either	$\bar{x} < a \text{ or } \bar{x} > b$
	where	
$a = \mu_0 - z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}}$	and	$b = \mu_0 + z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}}$

Compare these results with the confidence interval formula (presented in previous unit):

$$\bar{x} - z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}} \le \mu \le \bar{x} + z_{1-\alpha/2} \frac{\sigma}{\sqrt{n}}.$$

- > In this case, if H_0 is not true and H_1 holds with a specific value of $\mu = \mu_1$, then it is possible to compute the probability of type II error, β .
- > In the (very realistic) case where σ^2 is not known, but rather estimated by S^2 , we would like to replace the test statistic, Z, above with,

$$T = \frac{\overline{X} - \mu_0}{S/\sqrt{n}},$$

but in general, T no longer follows a Normal distribution.

> Under $H_0: \mu = \mu_0$, and for moderate or large samples (e.g. n > 100) this statistic is approximately Normally distributed just like above. In this case, the procedures above work well.

- > But for smaller samples, the distribution of T is no longer Normally distributed. Nevertheless, it follows a well known and very famous distribution of classical statistics: The Student-t Distribution.
- > The probability density function of a Student-t Distribution with a parameter v, referred to as degrees of freedom, is,

$$f(x;v) = \frac{\Gamma[(v+1)/2]}{\sqrt{\pi v} \Gamma(v/2)} \cdot \frac{1}{\left[(x^2/v) + 1 \right]^{(v+1)/2}} \qquad -\infty < x < \infty$$

where $\Gamma(\cdot)$ is the Gamma-function. It is a symmetric distribution about 0 and as $v \to \infty$ it approaches a standard Normal distribution.

- > The following mathematical result makes the *t*-distribution useful: Let X_1, X_2, \ldots, X_n be an i.i.d. sample from a Normal distribution with mean μ and variance σ^2 . The random variable, *T* has a *t*-distribution with n-1 degrees of freedom.
- > Now, knowing the distribution of T (and noticing it depends on the sample size, n), allows us to construct hypothesis tests and confidence intervals when σ^2 is not known, analogous to the (Z-tests and confidence intervals) presented above.
- > If \bar{x} and s are the mean and standard deviation of a random sample from a normal distribution with unknown variance σ^2 , a 100(1 - α)% confidence interval on μ is given by

$$\bar{x} - t_{1-\alpha/2,n-1} \frac{s}{\sqrt{n}} \le \mu \le \bar{x} + t_{1-\alpha/2,n-1} \frac{s}{\sqrt{n}},$$

where $t_{1-\alpha/2,n-1}$ is the $1-\alpha/2$ quantile of the t distribution with n-1 degrees of freedom.

> A related concept is a $100(1 - \alpha)\%$ prediction interval (PI) on a single future observation from a normal distribution is given by

$$\bar{x} - t_{1-\alpha/2,n-1}s\sqrt{1+\frac{1}{n}} \le X_{n+1} \le \bar{x} + t_{1-\alpha/2,n-1}s\sqrt{1+\frac{1}{n}}.$$

This is the range where we expect the n + 1 observation to be, after observing n observations and computing \overline{x} and s.

> Testing Hypotheses on the Mean, Variance Unknown (T-Tests)

Model: $X_i \stackrel{i.i.d.}{\sim} N(\mu, \sigma^2)$ with both μ and σ^2 unknown.

Null hypothesis: $H_0: \mu = \mu_0.$

Test statistic:

$$t = \frac{\overline{x} - \mu_0}{s/\sqrt{n}}, \qquad T = \frac{\overline{X} - \mu_0}{S/\sqrt{n}}.$$

Alternative Hypotheses	P-value	Rejection Criterion for Fixed-Level Tests
$H_1: \mu \neq \mu_0$	$P = 2\left[1 - F_{n-1}(t)\right]$	$t > t_{1-\alpha/2, n-1}$ or $t < t_{\alpha/2, n-1}$
$H_1: \mu > \mu_0$	$P = 1 - F_{n-1}(t)$	$t > t_{1-\alpha,n-1}$
$H_1: \mu < \mu_0$	$P = F_{n-1}(t)$	$t < t_{\alpha,n-1}$

Note that here, $F_{n-1}(\cdot)$ denotes the cdf of the t-distribution with n-1 degrees of freedom. As opposed to $\Phi(\cdot)$, it is not tabulated in standard tables and like $\Phi(\cdot)$ it cannot be explicitly evaluated. So to calculate *P*-values, we use software.

Two Sample Inference

- The setup is a sample x_1, \ldots, x_{n_1} modelled by an i.i.d. sequence of random variables, X_1, \ldots, X_{n_1} and another sample y_1, \ldots, y_{n_2} modelled by an i.i.d. sequence of random variables, Y_1, \ldots, Y_{n_1} . Observations, x_i and y_i (for same *i*) are not paired. In fact, it is possible that $n_1 \neq n_2$ (unequal sample sizes).
- ➤ The model assumed is, X_i ^{i.i.d.} N(µ₁, σ₁²), Y_i ^{i.i.d.} N(µ₂, σ₂²). Variations are: (i) equal variances: σ₁² = σ₂² := σ². (ii) unequal variances: σ₂² ≠ σ₂².
- > We could carry single sample inference for each population separately. Specifically, for $\mu_1 = E[X_i]$ and $\mu_2 = E[Y_i]$. However we focus on,

$$\Delta_{\mu} := \mu_1 - \mu_2 = E[X_i] - E[Y_i].$$

For this difference in means we can carry out inference jointly.

- > It is very common to ask if Δ_{μ} (=, <, >) 0, i.e. if μ_1 (=, <, >) μ_2 . But we can also replace the "0" with other values, e.g. $\mu_1 \mu_2 = \Delta_0$ for some Δ_0 .
- > A point estimator for Δ_{μ} is $\overline{X} \overline{Y}$ (difference in sample means). The estimate from the data is denoted by $\overline{x} \overline{y}$ (the difference in the individual sample means), with,

$$\overline{x} = \frac{1}{n_1} \sum_{i=1}^{n_1} x_i, \qquad \overline{y} = \frac{1}{n_2} \sum_{i=1}^{n_2} y_i.$$

➤ In the case (ii) of **unequal variances**: Point estimates for σ_1^2 and σ_2^2 are the individual sample variances,

$$s_1^2 = \frac{1}{n_1 - 1} \sum_{i=1}^{n_1} (x_i - \overline{x})^2, \qquad s_2^2 = \frac{1}{n_2 - 1} \sum_{i=1}^{n_2} (y_i - \overline{y})^2.$$

≻ In case (i) of equal variances, both S_1^2 and S_2^2 estimate σ^2 . In this case, a more reliable estimate can be obtained via the **pooled variance estimator**

$$S_p^2 = \frac{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}{n_1 + n_2 - 2}$$

> In case (i), under H_0 :

$$T = \frac{\overline{X} - \overline{Y} - \Delta_0}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}} \sim t(n_1 + n_2 - 2).$$

That is, the T test statistic follows a t-distribution with $n_1 + n_2 - 2$ degrees of freedom.

> In case (ii), under H_0 , there is only the approximate distribution,

$$T = \frac{\overline{X} - \overline{Y} - \Delta_0}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}} \quad \sim^{\text{approx}} \quad t(v).$$

where the degrees of freedom are

$$v = \frac{\left(\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}\right)^2}{\frac{\left(s_1^2/n_1\right)^2}{n_1 - 1} + \frac{\left(s_2^2/n_2\right)^2}{n_2 - 1}}.$$

If v is not an integer, may round down to the nearest integer (for using a table).

\succ Case (i):

Testing Hypotheses on Differences of Mean, Variance Unknown and Assumed Equal (two sample T-Tests with equal variance)

Model:	$X_i \stackrel{i.i.d.}{\sim} N(\mu_1, \sigma^2), \qquad Y_i \stackrel{i.i.d.}{\sim} N(\mu_2, \sigma^2).$	
Null hypothesis:	$H_0: \mu_1 - \mu_2 = \Delta_0.$	
Test statistic:	$t = \frac{\bar{x} - \bar{y} - \Delta_0}{s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}, \qquad T = \frac{\bar{X} - \bar{Y} - \Delta_0}{S_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}}.$	
Alternative Hypotheses	P-value	Rejection Criterion for Fixed-Level Tests
Hypotheses	$P-\text{value}$ $P = 2\left[1 - F_{n_1+n_2-2}(t)\right]$	0
Hypotheses $H_1: \mu_1 - \mu_2 \neq \Delta_0$		for Fixed-Level Tests

 \succ Case (ii):

Testing Hypotheses on Differences of Mean, Variance Unknown and NOT Equal (two sample T-Tests with unequal variance)

Model:	$X_i \stackrel{i.i.d.}{\sim} N(\mu_1, \sigma_1^2), \qquad Y_i \stackrel{i.i.d.}{\sim} N(\mu_2, \sigma_2^2).$	
Null hypothesis:	$H_0: \mu_1 - \mu_2 = \Delta_0.$	
Test statistic:	$t = \frac{\bar{x} - \bar{y} - \Delta_0}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}}, \qquad T = \frac{\bar{X} - \bar{Y} - \Delta_0}{\sqrt{\frac{S_1^2}{n_1} + \frac{S_2^2}{n_2}}}.$	
$\begin{array}{c} {\rm Alternative} \\ {\rm Hypotheses} \end{array}$	P-value	Rejection Criterion for Fixed-Level Tests
Hypotheses	$P-\text{value}$ $P = 2\left[1 - F_v(t)\right]$	U U
Hypotheses	$P = 2\left[1 - F_v(t)\right]$	for Fixed-Level Tests

 \succ Case (i) (Equal variances) - confidence interval:

$$\bar{x} - \bar{y} - t_{1-\alpha/2, n_1+n_2-2} \ s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}} \ \leq \ \mu_1 - \mu_2 \ \leq \ \bar{x} - \bar{y} + t_{1-\alpha/2, n_1+n_2-2} \ s_p \sqrt{\frac{1}{n_1} + \frac{1}{n_2}}$$

➤ Case (ii) (NOT Equal variances) - confidence interval:

$$\bar{x} - \bar{y} - t_{\alpha/2,v} \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}} \leq \mu_1 - \mu_2 \leq \bar{x} - \bar{y} + t_{\alpha/2,v} \sqrt{\frac{s_1^2}{n_1} + \frac{s_2^2}{n_2}}$$

Linear Regression

> The collection of statistical tools that are used to model and explore relationships between variables that are related in a nondeterministic manner is called **regression analysis**. Of key importance is the conditional expectation,

$$E(Y \mid x) = \mu_{Y \mid x} = \beta_0 + \beta_1 x \quad \text{with} \quad Y = \beta_0 + \beta_1 x + \epsilon,$$

where x is not random and ϵ is a Normal random variable with $E(\epsilon) = 0$ and $V(\epsilon) = \sigma^2$.

> Simple Linear Regression is the case where both x and y are scalars, in which case the data is,

$$(x_1, y_1), \ldots, (x_n, y_n)$$

Then given estimates of β_0 and β_1 denoted by $\hat{\beta}_0$ and $\hat{\beta}_1$ we have

$$y_i = \hat{\beta}_0 + \hat{\beta}_1 x_i + e_i \qquad i = 1, 2, \dots, n_i$$

where e_i , are the **residuals** and we can also define the **predicted observation**,

$$\hat{y}_i = \hat{\beta}_0 + \hat{\beta}_1 x_i$$

Ideally it would hold that $y_i = \hat{y}_i$ ($e_i = 0$) and thus **total mean squared error**

$$L := SS_E = \sum_{i=1}^n e_i^2 = \sum_{i=1}^n (y_i - \hat{y}_i)^2 = \sum_{i=1}^n (y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i)^2,$$

would be zero. But in practice, unless $\sigma^2 = 0$ (and all points lie on the same line), we have that L > 0.

> The standard (classic) way of determining the statistics $(\hat{\beta}_0, \hat{\beta}_1)$ is by minimisation of L. The solution, called the **least squares estimators** must satisfy

$$\frac{\partial L}{\partial \beta_0}\Big|_{\hat{\beta}_0\hat{\beta}_1} = -2\sum_{i=1}^n (y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i) = 0$$
$$\frac{\partial L}{\partial \beta_1}\Big|_{\hat{\beta}_0\hat{\beta}_1} = -2\sum_{i=1}^n (y_i - \hat{\beta}_0 - \hat{\beta}_1 x_i)x_i = 0$$

Simplifying these two equations yields

$$n\hat{\beta}_{0} + \hat{\beta}_{1}\sum_{i=1}^{n} x_{i} = \sum_{i=1}^{n} y_{i}$$
$$\hat{\beta}_{0}\sum_{i=1}^{n} x_{i} + \hat{\beta}_{1}\sum_{i=1}^{n} x_{i}^{2} = \sum_{i=1}^{n} y_{i}x_{i}$$

These are called the **least squares normal equations**. The solution to the normal equations results in the **least squares estimators** $\hat{\beta}_0$ and $\hat{\beta}_1$. Using the sample means, \bar{x} and \bar{y} the estimators are,

$$\hat{\beta}_{0} = \bar{y} - \hat{\beta}_{1}\bar{x}, \qquad \qquad \hat{\beta}_{1} = \frac{\sum_{i=1}^{n} y_{i}x_{i} - \frac{\left(\sum_{i=1}^{n} y_{i}\right)\left(\sum_{i=1}^{n} x_{i}\right)}{n}}{\sum_{i=1}^{n} x_{i}^{2} - \frac{\left(\sum_{i=1}^{n} x_{i}\right)^{2}}{n}}.$$

 \succ The following quantities are also of common use:

$$S_{xx} = \sum_{i=1}^{n} (x_i - \bar{x})^2 = \sum_{i=1}^{n} x_i^2 - \frac{\left(\sum_{i=1}^{n} x_i\right)^2}{n}$$
$$S_{xy} = \sum_{i=1}^{n} (y_i - \bar{y})(x_i - \bar{x}) = \sum_{i=1}^{n} x_i y_i - \frac{\left(\sum_{i=1}^{n} x^i\right)\left(\sum_{i=1}^{n} y^i\right)}{n}$$
$$\hat{\beta}_i = \frac{S_{xy}}{n}$$

Hence,

$$\hat{\beta}_1 = \frac{S_{xy}}{S_{xx}}$$

Further,

$$SS_T = \sum_{i=1}^n (y_i - \bar{y})^2, \qquad SS_R = \sum_{i=1}^n (\hat{y}_i - \bar{y})^2, \qquad SS_E = \sum_{i=1}^n (y_i - \hat{y}_i)^2.$$

➤ The Analysis of Variance Identity is

$$\sum_{i=1}^{n} \left(y_i - \bar{y} \right)^2 = \sum_{i=1}^{n} \left(\hat{y}_i - \bar{y} \right)^2 + \sum_{i=1}^{n} \left(y_i - \hat{y}_i \right)^2$$

or,

$$SS_T = SS_R + SS_E$$

Also, $SS_R = \hat{\beta}_1 S_{xy}$.

> An Estimator of the Variance, σ^2 is

$$\hat{\sigma}^2 := MS_E = \frac{SS_E}{n-2}$$

 \succ A widely used measure for a regression model is the following ratio of sum of squares, which is often used to judge the adequacy of a regression model:

$$R^2 = \frac{SS_R}{SS_T} = 1 - \frac{SS_E}{SS_T}.$$

$$E\left(\hat{\beta}_{0}\right) = \beta_{0}, \qquad V\left(\hat{\beta}_{0}\right) = \sigma^{2}\left[\frac{1}{n} + \frac{\bar{x}^{2}}{S_{XX}}\right]$$
$$E\left(\hat{\beta}_{1}\right) = \beta_{1}, \qquad V\left(\hat{\beta}_{1}\right) = \frac{\sigma^{2}}{S_{XX}}.$$

In simple linear regression, the estimated standard error of the slope and the estimated standard error of the intercept are

$$se(\hat{\beta}_1) = \sqrt{\frac{\hat{\sigma}^2}{S_{XX}}}$$
 and $se(\hat{\beta}_0) = \sqrt{\hat{\sigma}^2 \left[\frac{1}{n} + \frac{\bar{x}^2}{S_{XX}}\right]}$

 \succ The Test Statistic for the Slope is

$$T = \frac{\hat{\beta}_1 - \beta_{1,0}}{\sqrt{\hat{\sigma}^2 / S_{XX}}}$$

$$H_0: \beta_1 = \beta_{1,0} \qquad \qquad H_1: \beta_1 \neq \beta_{1,0}$$

Under H_0 the test statistic T follows a **t** - **distribution** with "n-2 degree of freedom".

> An alternative is to use the F statistic as is common in **ANOVA** (Analysis of Variance) – not covered fully in the course.

$$F = \frac{SS_R/1}{SS_E/(n-2)} = \frac{MS_R}{MS_E}$$

Under H_0 the test statistic F follows an **F** - **distribution** with "1 degree of freedom in the numerator and n-2 degrees of freedom in the denominator".

Analysis of Variance Table for Testing Significance of Regression

Source of Variation		Degrees of Freedom	Mean Square	F ₀
Regression	$SS_R = \hat{\beta}_1 S_{xy}$ $SS_E = SS_T - \hat{\beta}_1 S_{xy}$	1	MS_R	MS_R/MS_E
Error	$SS_E = SS_T - \hat{\beta}_1 S_{xy}$	n-2	MS_E	
Total	SS_T	n-1		

- > There are also confidence intervals for β_0 and β_1 as well as prediction intervals for observations. We don't cover these formulas.
- > To check the regression model assumptions we plot the residuals e_i and check for (i) Normality. (ii) Constant variance. (iii) Independence.

Logistic Regression:

- > Take the response variable, Y_i as a Bernoulli random variable. In this case notice that E(Y) = P(Y = 1).
- \succ The logit response function has the form

$$E(Y) = \frac{\exp(\beta_0 + \beta_1 x)}{1 + \exp(\beta_0 + \beta_1 x)}.$$

- > Fitting a logistic regression model to data yields estimates of β_0 and β_1 .
- \succ The following formula is called the **odds**

$$\frac{E(Y)}{1 - E(Y)} = \exp(\beta_0 + \beta_1 x).$$

 $\begin{array}{c} -1.9 \\ -1.7 \\ -1.5 \end{array}$ -1.0This table was generated using the "CDF" command in Minitab. -0.1-0.9-0.6-2.0-2.4-2.3-2.1-2.6-2.8 -2.9 -2.7 ώ 0 N .3446 .3821 .4207 .4602 .0179 .0139 0107 .2119 .0968 .0668 .0548 .0287 .0359 .0228 .0082 .0047 .0035 .0007 5000. 308 .2743 .2420 .1357 .1151 8080. 2000 .0026 6100 £100. 0100 184 158 .00 .0174 .0136 .0104 .4168 .4562 .4960 .1131 1560 .0793 .0281 .0351 .0436 .0537 .0045 .0025 0007 .3783 0800 .0034 8100 £100. 6000 3409 .2389 .1814 .1335 .0655 0000 3050 2709 1562 0060 0 .0102 .4129 .4522 .4920 .2358 .1314 .1112 .0934 .0778 .0427 .0344 .0170 .0132 .0044 9000 .3745 .3372 .2676 .2061 .1539 .0643 .0526 .0274 .0217 8700 .0059 .0033 .0024 8100 .0013 6000 3015 1788 0005 000 .02 .0166 .0129 .3336 .3707 .4090 .4488 .0764 .0516 .0268 .0336 .0418 .0212 6600 .0023 9000 .1515 .1292 .1093 8160 .0630 .0075 .0057 .0043 .0032 7100 .0012 6000 .2327 .1762 000 298 .03 .44052 .44443 .4840 .1271 .1492 .1075 1060 .0749 8190 .0505 .0409 .0329 .0262 .0207 .0162 .0125 9600 .0073 .0041 0031 .0023 0016 8000 .0006 .3669 .3300 .0012 261 2005 0055 0003 2940 1736 . 04 .0735 .0495 .0256 .0322 .0158 .0122 .0094 .0006 .4013 .4404 .2578 .1056 C880. 0606 .0202 .0071 .0040 .0030 .0022 8000 3632 2912 .2266 .1977 171 0054 .0016 001 0001 3264 1469 125 .05 .3974 .4364 .4761 .0154 .0091 .0869 .0485 .0392 .0250 .0197 6110 .0069 .0029 0015 .0008 .0006 .0721 .0039 .0021 3594 2546 2236 1038 0594 0052 001 000 3228 2877 1949 1685 1446 1230 .06 .0475 .0244 .0307 .0192 .0150 .0116 0089 .3936 .4325 .4721 .3557 .2514 .1210 .1020 .0708 .0582 8900 .0051 .0038 .0028 .0021 .0015 8000 .0005 3192 .1660 .1423 001 0004 284 9 .2483 .2177 8580. .0694 .0465 .0571 .0375 .0239 8810 .0146 .0113 0087 .0066 .0005 .3520 .3156 .2810 .1635 .1190 .1003 .0049 .0037 .0027 .0020 .0014 .0010 .0007 0004 4681 3897 1894 140 000 .08 .3483 .3859 4247 6860 .0455 .0367 .0183 .0143 .0110 .0084 .3121 .2451 .2148 .1170 .0823 .0681 .0559 .0233 .0064 .0048 .0036 .0026 .0019 .0014 .0010 .0007 0005 404 2776 1867 0003 161 1379 2000 . 09 2.5 2.6 2.7 2.2 2.3 2.4 0.7 0.5 0.00.20.32.0 0.4 ωωωω 4ω2-3.0 2.8 1.0 0.9 0.8 1.94 <u>ω</u> 1 N .9332 .9452 .9554 .9641 .9713 .8849 .9032 .9192 .9938 .9953 .9965 .9974 .9773 .9821 .9861 .9893 .8413 8159 .6915 .6179 .5793 .5000 788 .7580 6554 8 .9564 .9463 .9345 .8869 8665 .8186 .5832 .9995 5666 £666 .9975 .9966 .9955 .9940 .9896 .9920 .9864 .9826 .9778 .9719 .9649 .9207 .9049 .8438 .6950 .6591 .6217 .5438 .7910 .7291 866 .9982 .7611 5040 <u>0</u> .9956 .9967 .9976 .9922 .9898 .9868 .9783 .9726 .9656 .9573 .9357 .9474 .9222 .9066 8888. 9898 .8212 .7939 .7642 .7324 .6985 .6628 .6255 .5871 .5080 .999 9999 .9999. 1999. .9941 .8461 .02 .8485 .9943 .9957 .9968 .9977 .9901 .9871 .9788 .9370 .9484 .9582 .9664 .9732 .9236 .9082 .8907 .7357 .7019 .6664 .6293 .5910 .5120 7666 1666 1866 7967 7673 1666 1666 .<u>0</u> .9793 .9495 .9969 .9959 .9945 .9927 .9904 .9875 .9838 .9591 .9671 .9738 .9382 .9251 9099 .8925 .8729 .6331 .5948 .5557 .9999 .9992 .9994 .9977 .7054 .6700 8866 9984 8508 .8264 .7995 7704 .7389 5160 04 .9970 .9960 .9946 .9906 .9878 .9842 .9798 .9744 .9678 .9599 .9394 .9265 .9115 .8944 .8531 .8749 .8023 .7422 .6736 .6368 .5987 .7734 99992 99994 99996 9978 9984 8289 .7088 5199 . G .9961 .9971 .9979 .9406 .9131 .8962 .8554 .6026 .9948 .9909 .9846 .9803 .9686 .9608 .9279 .6772 .6406 .99985 .99992 .9881 .8051 .5239 999 8315 7764 .7454 7123 . 06 .9525 .9147 0868 .8790 .6064 .9992 .9995 .9979 .9972 .9962 .9949 .9932 .9911 .9884 .9850 .9808 .9693 .9756 .9616 .9418 9292 .8577 .8340 .8078 .7486 .7157 .6808 .6443 .5675 2866 .7794 999 900 5279 97 .9812 .9854 .9162 .6103 .9995 09993 .9951 .9963 .9934 .9913 .9887 .9699 .9761 .9625 .9429 .9535 .9306 .8997 .8599 .8365 9018 .7517 .7190 .6844 .6480 .5714 999 866 998(7823 5319 .08 .9936 0686 .9633 .9706 .9767 .9441 .9545 .9319 .9177 .9015 .8133 .6141 .9974 .9964 .9952 9166 .9817 .9857 .8621 .8830 .7549 6879 .6517 8066 666 5666 E666 00666 9866 866 8389 7852 .5753 5359 7224 .09

Standard Normal Cumulative Probabilities

t-Distribution Quantiles

ν	Q(.9)	Q (.95)	Q(.9 75)	Q(.99)	Q(.995)	Q(.999)	Q(.9995)
1	3.078	6.314	12.706	31.821	63.657	318.317	636.607
2	1.886	2.920	4.303	6.965	9.925	22.327	31.598
3	1.638	2.353	3.182	4.541	5.841	10.215	12.924
4	1.533	2.132	2.776	3.747	4.604	7.173	8.610
5	1.476	2.015	2.571	3.365	4.032	5.893	6.869
6	1.440	1.943	2.447	3.143	3.707	5.208	5.959
7	1.415	1.895	2.365	2.998	3.499	4.785	5.408
8	1.397	1.860	2.306	2.896	3.355	4.501	5.041
9	1.383	1.833	2.262	2.821	3.250	4.297	4.781
10	1.372	1.812	2.228	2.764	3.169	4.144	4.587
11	1.363	1.796	2.201	2.718	3.106	4.025	4.437
12	1.356	1.782	2.179	2.681	3.055	3.930	4.318
13	1.350	1.771	2.160	2.650	3.012	3.852	4.221
14	1.345	1.761	2.145	2.624	2.977	3.787	4.140
15	1.341	1.753	2.131	2,602	2.947	3.733	4.073
16	1.337	1.746	2.120	2.583	2.921	3.686	4.015
17	1.333	1.740	2.110	2.567	2.898	3.646	3.965
18	1.330	1.734	2.101	2.552	2.878	3.610	3.922
19	1.328	1.729	2.093	2.539	2.861	3.579	3.883
20	1.325	1.725	2.086	2.528	2.845	3.552	3.849
21	1.323	1.721	2.080	2.518	2.831	3.527	3.819
22	1.321	1.717	2.074	2.508	2.819	3.505	3.792
23	1.319	1.714	2.069	2.500	2.807	3.485	3.768
24	1.318	1.711	2.064	2.492	2.797	3.467	3.745
25	1.316	1.708	2.060	2.485	2.787	3.450	3.725
26	1.315	1.706	2.056	2.479	2.779	3.435	3.707
27	1.314	1.703	2.052	2.473	2.771	3.421	3.690
28	1.313	1.701	2.048	2.467	2.763	3.408	3.674
29	1.311	1.699	2.045	2.462	2.756	3.396	3.659
30	1.310	1.697	2.042	2.457	2.750	3.385	3.646
40	1.303	1.684	2.021	2.423	2.704	3,307	3.551
60	1.296	1.671	2.000	2.390	2.660	3.232	3.460
120	1.289	1.658	1.980	2.358	2.617	3.160	3.373
œ	1.282	1.645	1.960	2.326	2.576	3.090	3.291

This table was generated using the "INVCDF" command in Minitab.