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IB AI HL

4. Statistics & Probability 4.2 Correlation & Regression







IB Maths DP

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4.1 Statistics Toolkit

4.1.1 Sampling

Types of Data

What are the different types of data?

- Qualitative data is data that is usually given in words not numbers to describe something
 For example: the colour of a teacher's car
- Quantitative data is data that is given using numbers which counts or measures something
 - · For example: the number of pets that a student has
- Discrete data is quantitative data that needs to be counted
 - Discrete data can only take specific values from a set of (usually finite) values
 - For example: the number of times a coin is flipped until a 'tails' is obtained
- · Continuous data is quantitative data that needs to be measured
 - · Continuous data can take any value within a range of infinite values
 - For example: the height of a student
- Age can be discrete or continuous depending on the context or how it is defined
 - If you mean how many years old a person is then this is discrete
 - If you mean how long a person has been alive then this is continuous

What is the difference between a population and a sample?

- The population refers to the whole set of things which you are interested in
 - For example: if a vet wanted to know how long a typical French bulldog slept for in a day then the population would be all the French bulldogs in the world
- A sample refers to a subset of the population which is used to collect data from
 - For example: the vet might take a sample of French bulldogs from different cities and record how long they sleep in a day
- A sampling frame is a list of all members of the population
 - · For example: a list of employees' names within a company
- Using a sample instead of a population:
 - Is quicker and cheaper
 - Leads to less data needing to be analysed
 - · Might not fully represent the population
 - Might introduce bias



Sampling Techniques

What is a random sample and a biased sample?

- A random sample is where every member of the population has an equal chance of being included in the sample
- A biased sample is one from which misleading conclusions could be drawn about the population
 - Random sampling is an attempt to minimise bias

What sampling techniques do I need to know?

Simple random sampling

- Simple random sampling is where every group of members from the population has an equal probability of being selected for the sample
- To carry this out you would...
 - uniquely number every member of a population
 - randomly select n different numbers using a random number generator or a form of lottery (where numbers are selected randomly)
- Effectiveness:
 - Useful when you have a small population or want a small sample (such as children in a class)
 - It can be time-consuming if the sample or population is large
 - This can not be used if it is not possible to number or list all the members of the population (such as fish in a lake)

Systematic sampling

 Systematic sampling is where a sample is formed by choosing members of a population at regular intervals using a list

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- To carry this out you would...
- calculate the size of the interval $k = \frac{\text{size of population } (N)}{N}$
- choose a random starting point between 1 and k
- select every kth member after the first one
- Effectiveness:
 - Useful when there is a natural order (such as a list of names or a conveyor belt of items)
 - Quick and easy to use
 - · This can not be used if it is not possible to number or list all the members of the population (such as penguins in Antarctica)

Stratified sampling

 Stratified sampling is where the population is divided into disjoint groups and then a random sample is taken from each group



- The proportion of a group that is sampled is equal to the proportion of the population that belong to that group
- To carry this out you would...
 - Calculate the number of members sampled from each stratum
 - size of sample (*n*)
 - $r_{\text{size of population } (N)}$ × number of members in the group
 - Take a random sample from each group
- Effectiveness:
 - Useful when there are very different groups of members within a population
 - The sample will be representative of the population structure
 - The members selected from each stratum are chosen randomly
 - This can not be used if the population can not be split into groups or if the groups overlap

Quota sampling

- · Quota sampling is where the population is split into groups (like stratified sampling) and members of the population are selected until each quota is filled
- To carry this out you would...
 - Calculate how many people you need from each group
 - Select members from each group until that quota is filled
 - The members do not have to be selected randomly
- Effectiveness:
 - Useful when collecting data by asking people who walk past you in a public place or when a sampling frame is not available
 - This can introduce bias as some members of the population might choose not to be included in the sample

Convenience sampling

- Convenience sampling is where a sample is formed using available members of the population who fit the criteria
- To carry this out you would...
 - Select members that are easiest to reach
- Effectiveness:
 - Useful when a list of the population is not possible
 - This is unlikely to be representative of the population structure
 - This is likely to produce biased results

What are the main criticisms of sampling techniques?

- Most sampling techniques can be improved by taking a larger sample
- Sampling can introduce bias so you want to minimise the bias within a sample
 - To minimise bias the sample should be as close to random as possible
- A sample only gives information about those members

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• Different samples may lead to different conclusions about the population





Reliability of Data

How can I decide if data is reliable?

- Data from a sample is reliable if similar results would be obtained from a different sample from the same population
- The sample should be **representative** of the population
- The sample should be **big enough**
 - Sampling a small proportion of a population is unlikely to be reliable

What can cause data to be unreliable?

- If the sample is **biased**
 - It is not random
- If errors are made when collecting data
 - Numbers could be recorded incorrectly, duplicated or missed out
- If the person collecting the data **favours some members** over others
 - They might seek out members who will lead to a desired outcome
 - They might exclude members if they would cause the sample to oppose the desired outcome
- If a significant proportion of data is missing
 - Some data may be unavailable
 - Some members might decide not to be part of the sample
 - This will mean the results are not necessarily representative of the population

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4.1.2 Data Collection

Methods of Data Collection

How do I choose variables to investigate?

- Keep the number of variables you investigate to a minimum
 - Too many variables at once can be **overwhelming**
 - It can be time-consuming to process unnecessary data
- You should choose variables that are linked to what you are investigating
 - If you are investigating the ability of adults to solve puzzles you might use the time it takes them as a variable
 - Consider which variables are likely to have an effect on what you are investigating
 - An adult's reading speed will affect their time to solve a puzzle
 - An adult's height is unlikely to affect their time to solve a puzzle

What makes a good survey?

- A survey is a method of collecting data
- Consider whether the survey needs to be **in-person**
 - A person might be less like<mark>ly to answer questions tru</mark>thfully in person
 - You can quickly survey more people remotely or electronically
 - Such as postal surveys, phone surveys, internet surveys
- Consider whether the interviewer could unintentionally influence participants' responses
 - If a headteacher is asking students whether they enjoy school then they are more likely to say yes as they think that is what the headteacher wants to hear
 - This will introduce bias PAPERS PRACTICE

What makes a good questionnaire?

- A questionnaire is a list of questions
- The questions should be **unbiased**
 - Questions should not be leading
 - For example: "You enjoy school, don't you?"
 - If options are given for the participant to choose from then they should cover all possible responses
- The questions should not be personal
 - This means you should not ask for **unnecessary personal information**
 - Such as date of birth, address, etc
 - The questions should **not reflect your personal opinions**
 - For example: "Do you enjoy watching the boring news on TV?"
 - People can find it difficult to rate personal feeling/qualities
 - For example: "How smart do you think you are?"
- Questions can be structured or unstructured
 - Structured questions usually ask the participants to choose from **options**, give a **rating** or **rank** options
 - These can be quick to analyse
 - The answer choices should be consistent where appropriate
 - Unstructured questions let the participants to express their views in their own words



- These tend to be more **open-ended** questions
- These can take longer to analyse but can give more in-depth views
- Questions should be precise and unambiguous
 - They should be phrased in a way in which the participants understand exactly what you mean
 - For example: "Do you study French or Spanish at school" is not precise
 - Some people might reply with "Yes" or "No"
 - Some people might reply with "French" or "Spanish"





Reliability & Validity

What is reliability & validity of a data collection method?

- Reliability measures how consistent a process is at measuring a variable
 - A process is **reliable** if you would get the **same results** by **repeating** the process with the **same sample** using the **same conditions**
- Validity measures how accurate a process is at measuring a variable
 - A process is valid if it is accurately measuring the variable you want it to measure
- If your process is found not to be reliable or valid then:
 - Adjust the data collection process
 - Change the sampling technique
 - Use a larger sample

What are tests to check reliability?

- Test-retest
 - This is where you use a data collection process with a sample and then repeat the same process with the same sample at a later time
 - The results should show positive correlation if the process is reliable
 - The results might not perfectly match due to external factors during the gap between the data collection
 - Once the sample has been through the process once they will be familiar so this could lead to different results from the second process
- Parallel forms
 - This is where you give the same sample a second set of questions (or second set of experiments) which are similar to the first set **PACTICF**
 - The results should show positive correlation if the process is reliable
 - It can be difficult to make the two processes similar to each other

What are tests to check validity?

- · Content-related validity checks
 - This is where you check how well the **process measures all aspects of the variable**
 - If the process is valid then it should cover all aspects of the variable
 - These checks require knowledge of the variable so experts are often used
 - An example of a process that is valid: A teacher wants to assess how well students understand calculus so they set questions covering differentiation, integration and applications
 - An example of a process that is **not valid**: A restaurant manager wants to assess how good a chef is at cooking steaks so asks the chef to make 10 medium steaks
- Criterion-related validity checks
 - This is where you check how well **one variable predicts the outcome for another variable** (called the criterion variable)
 - If the process is valid then the variable should be a good predictor
 - An example of a process that is valid: Results from a mock exam being used to predict the results in the actual exam

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 An example of a process that is not valid: Results from measuring the heights of meerkats being used to predict the heights of squirrels

Worked Example

Tomas is a dog trainer. Before he agrees to train a dog he assesses the dog's obedience. To do this, he first visits the dog, asks it to perform 10 basic commands and records how many the dog successfully carries out. Two days later, Tomas visits the dog a second time and asks it to do the same 10 commands. Tomas assesses 8 dogs using this process and the table below shows the number of commands performed successfully by each dog on each visit.

Dog	1	2	3	4	5	6	7	8
First visit	3	5	2	3	6	2	0	5
Second visit	3	5	2	4	5	2	1	5

a)

State the reliability test that Tomas is using.

Tomas is using exactly the same process with the same sample Test-retest

Comment on the reliability of Tomas' process.

The number of commands that each dog successfully performed on the second visit was either the same as the first visit or very similar. Therefore the process is reliable.

b)



4.1.3 Statistical Measures

Mean, Mode, Median

What are the mean, mode and median?

- Mean, median and mode are measures of central tendency
 - They describe where the centre of the data is
- They are all types of averages
- · In statistics it is important to be specific about which average you are referring to
- The units for the mean, mode and median are the same as the units for the data

How are the mean, mode, and median calculated for ungrouped data?

- The mode is the value that occurs most often in a data set
 - It is possible for there to be more than one mode
 - It is possible for there to be **no mode**
 - In this case **do not** say the mode is zero
- The median is the middle value when the data is in order of size
 - If there are two values in the middle then the median is the midpoint of the two values
- The mean is the sum of all the values divided by the number of values

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$$_{i=1}^{\overline{x}} \sum_{j=1}^{n} x_{i}$$

- Where $\sum_{i=1}^{n} x_i = x_1 + x_2 + \dots + x_n$ is the sum of the *n* pieces of data
- $\circ~$ The mean can be represented by the symbol μ
- Your GDC can calculate these statistical measures if you input the data using the statistics mode







Quartiles & Range

What are quartiles?

- Quartiles are measures of location
- Quartiles divide a population or data set into four equal sections
 - The lower quartile, Q1 splits the lowest 25% from the highest 75%
 - The median, Q₂ splits the lowest 50% from the highest 50%
 - The upper quartile, Q₃ splits the lowest 75% from the highest 25%
- There are different methods for finding quartiles
 - Values obtained by hand and using technology may differ
- · You will be expected to use your GDC to calculate the quartiles

What are the range and interquartile range?

- The range and interquartile range are both measures of dispersion
 - They describe how spread out the data is
- The range is the largest value of the data minus the smallest value of the data
- The interquartile range is the range of the central 50% of data
 - It is the upper quartile minus the lower quartile



- This is given in the formula booklet
- The units for the range and interquartile range are the same as the units for the data







Standard Deviation & Variance

What are the standard deviation and variance?

- The standard deviation and variance are both measures of dispersion
 - They describe how spread out the data is in relation to the mean
- The variance is the mean of the squares of the differences between the values and the mean
 - Variance is denoted σ²
- The standard deviation is the square-root of the variance
 - Standard deviation is denoted σ
- The units for the standard deviation are the same as the units for the data
- The units for the variance are the square of the units for the data

How are the standard deviation and variance calculated for ungrouped data?

- In the exam you will be expected to use the statistics function on your GDC to calculate the standard deviation and the variance
- Calculating the standard deviation and the variance by hand may deepen your

understanding
• The formula for variance is
$$\sigma^2 = \frac{\sum_{i=1}^{k} f_i (x_i - \mu)^2}{n}$$

• This can be rewritten as **PAPERS PRACTICE**

$$\sigma^2 = \frac{\sum_{i=1}^{k} f_i x_i^2}{n} - \mu^2$$

• The formula for standard deviation is
$$\sigma = \sqrt{\frac{\sum_{i=1}^{k} f_i(x_i - \mu)^2}{n}}$$

This can be rewritten as

$$\sigma = \sqrt{\frac{\sum_{i=1}^{k} f_i x_i^2}{n} - \mu^2}$$

You do not need to learn these formulae as you will use your GDC to calculate these







4.1.4 Frequency Tables

Ungrouped Data

How are frequency tables used for ungrouped data?

- Frequency tables can be used for ungrouped data when you have lots of the same values within a data set
 - They can be used to collect and present data easily
- If the value 4 has a frequency of 3 this means that there are three 4's in the data set

How are measures of central tendency calculated from frequency tables with ungrouped data?

- The mode is the value that has the highest frequency
- The median is the middle value
 - Use cumulative frequencies (running totals) to find the median
- The mean can be calculated by
 - Multiplying each value xi by its frequency fi
 - Summing to get $\Sigma f_i x_i$
 - Dividing by the total frequency $n = \Sigma f_i$
 - This is given in the formula booklet



 Your GDC can calculate these statistical measures if you input the values and their frequencies using the statistics mode

How are measures of dispersion calculated from frequency tables with ungrouped data?

- The range is the largest value of the data minus the smallest value of the data
- The interquartile range is calculated by

$$IQR = Q_3 - Q_1$$

- The **quartiles** can be found by using your GDC and inputting the values and their frequencies
- The standard deviation and variance can be calculated by hand using the formulae
 - Variance

$$\sigma^2 = \frac{\sum_{i=1}^k f_i x_i^2}{n} - \mu^2$$

Standard deviation





- You **do not need to learn** these formulae as you will be expected to use your GDC to find the standard deviation and variance
 - You may want to see these formulae to deepen your understanding

Exam Tip

- Always check whether your answers make sense when using your GDC
 - The value for a measure of central tendency should be within the range of data









Use GDC
$$\sigma_x = 1.159...$$

Standard deviation = 1.16 (3sf)





Grouped Data

How are frequency tables used for grouped data?

- Frequency tables can be used for grouped data when you have lots of the same values within the same interval
 - · Class intervals will be written using inequalities and without gaps
 - $10 \le x < 20$ and $20 \le x < 30$
 - If the class interval $10 \le x \le 20$ has a frequency of 3 this means there are three values in that interval
 - You do not know the exact data values when you are given grouped data

How are measures of central tendency calculated from frequency tables with grouped data?

- The modal class is the class that has the highest frequency
 - This is for equal class intervals only
- The median is the middle value
 - The exact value can not be calculated but it can be estimated by using a **cumulative** frequency graph
- The exact mean can not be calculated as you do not have the raw data
- The mean can be estimated by
 - Identifying the mid-interval value (midpoint) xi for each class
 - Multiplying each value by the class frequency *f*i
 - Summing to get $\Sigma f_i x_i$
 - Dividing by the total frequency $n = \Sigma f_i$
 - This is given in the formula booklet ERS PRACTICE

$$\overline{x} = \frac{\sum_{i=1}^{k} f_i x_i}{n}$$

• Your **GDC** can estimate the mean if you input the mid-interval values and the class frequencies using the statistics mode

How are measures of dispersion calculated from frequency tables with grouped data?

- The exact range can not be calculated as the largest and smallest values are unknown
- The interquartile range can be estimated by

$$IQR = Q_3 - Q_1$$

- Estimates of the quartiles can be found by using a cumulative frequency graph
- The **standard deviation** and **variance** can be estimated using the mid-interval values x_i in the formulae
 - Variance



$$\sigma^2 = \frac{\sum_{i=1}^{k} f_i x_i^2}{n} - \mu^2$$

• Standard deviation

$$\sigma = \sqrt{\frac{\sum_{i=1}^{k} f_i x_i^2}{n} - \mu^2}$$

- You **do not need to learn** these formulae as you will be expected to use your GDC to estimate the standard deviation and variance using the mid-interval values
 - You may want to use these formulae to deepen your understanding



Exam Tip

- As you can only estimate statistical measures from a grouped frequency table it is good practice to indicate that the values are not exact
 - You can do this by rounding values rather than leaving as surds and fractions
 - x = 0.333 (3sf) rather than x = + EXAM PAPERS PRACTICE



Worked Example

The table below shows the heights in cm of a group of 25 students.

Height, <u>h</u>	Frequency		
$150 \le h < 155$	3		
$155 \le h < 160$	5		
$160 \le h < 165$	9		
$165 \le h < 170$	7		
$170 \le h < 175$	1		

a)

Write down the modal class.





4.1.5 Linear Transformations of Data

Linear Transformations of Data

Why are linear transformations of data used?

- Sometimes data might be very large or very small
- You can apply a **linear transformation** to the data to make the values more manageable
 - You may have heard this referred to as:
 - Effects of constant changes
 - Linear coding
- Linear transformations of data can affect the statistical measures

How is the mean affected by a linear transformation of data?

- Let x be the mean of some data
- If you multiply each value by a constant k then you will need to multiply the mean by k
 Mean is kx
- If you add or subtract a constant a from all the values then you will need to add or subtract the constant a to the mean
 - Mean is $\overline{x} \pm a$

How is the variance and standard deviation affected by a linear transformation of data?

- Let σ^2 be the **variance** of some data
 - σ is the standard deviation Δ PERS PRACTICI
- If you multiply each value by a constant k then you will need to multiply the variance by k²
 - \circ Variance is $k^2\sigma^2$
 - You will need to **multiply** the **standard deviation** by the **absolute value** of k
 - Standard deviation is $|k|\sigma$
 - If you add or subtract a constant *a* from all the values then the variance and the standard deviation stay the same
 - Variance is σ^2
 - Standard deviation is σ

Exam Tip

- If you forget these results in an exam then you can look in the HL section of the formula booklet to see them written in a more algebraic way
 - Linear transformation of a single variable



• where E(...) means the mean and Var(...) means the variance

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4.1.6 Outliers

Outliers

What are outliers?

- Outliers are extreme data values that do not fit with the rest of the data
 They are either a lot bigger or a lot smaller than the rest of the data
- Outliers are defined as values that are more than 1.5 × IQR from the nearest quartile
 x is an outlier if x < Q₁ 1.5 × IQR or x > Q₃ + 1.5 × IQR
- Outliers can have a big effect on some statistical measures

Should I remove outliers?

- The decision to remove outliers will depend on the context
- Outliers should be removed if they are found to be errors
 - The data may have been recorded incorrectly
 - For example: The number 17 may have been recorded as 71 by mistake
- Outliers should not be removed if they are a valid part of the sample
 - The data may need to be checked to verify that it is not an error
 - For example: The annual salaries of employees of a business might appear to have an outlier but this could be the director's salary

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4.1.7 Univariate Data

Box Plots

Univariate data is data that is in **one variable**.

What is a box plot (box and whisker diagram)?

- A box plot is a graph that clearly shows key statistics from a data set
 - It shows the median, quartiles, minimum and maximum values and outliers
 - It does not show any other individual data items
- The middle 50% of the data will be represented by the box section of the graph and the lower and upper 25% of the data will be represented by each of the whiskers
- Any outliers are represented with a cross on the outside of the whiskers
 - If there is an outlier then the whisker will end at the value before the outlier
- Only one axis is used when graphing a box plot
- It is still important to make sure the axis has a clear, even scale and is labelled with units



What are box plots useful for?

- · Box plots can clearly show the shape of the distribution
 - If a box plot is symmetrical about the median then the data could be **normally distributed**
- Box plots are often used for comparing two sets of data
 - Two box plots will be drawn next to each other using the same axis
 - They are useful for **comparing data** because it is easy to see the main shape of the distribution of the data from a box plot
 - You can easily compare the medians and interquartile ranges

Exam Tip

- In an exam you can use your GDC to draw a box plot if you have the raw data
 - You calculator's box plot can also include outliers so this is a good way to check







Cumulative Frequency Graphs

What is cumulative frequency?

- The cumulative frequency of x is the running total of the frequencies for the values that are less than or equal to x
- For grouped data you use the upper boundary of a class interval to find the cumulative frequency of that class

What is a cumulative frequency graph?

- A cumulative frequency graph is used with data that has been organised into a grouped frequency table
- Some coordinates are plotted
 - The x-coordinates are the upper boundaries of the class intervals
 - The y-coordinates are the cumulative frequencies of that class interval
- The coordinates are then joined together by hand using a smooth increasing curve

What are cumulative frequency graphs useful for?

- They can be used to estimate statistical measures
 - Draw a horizontal line from the y-axis to the curve
 - For the median: draw the line at 50% of the total frequency
 - For the lower quartile: draw the line at 25% of the total frequency
 - For the upper guartile: draw the line at 75% of the total frequency.
 - For the pth percentile: draw the line at p% of the total frequency
 - Draw a vertical line down from the curve to the x-axis
 This x-value is the relevant statistical measure
- · They can used to estimate the number of values that are bigger/small than a given value
 - Draw a vertical line from the given value on the x-axis to the curve
 - Draw a horizontal line from the curve to the y-axis
 - This value is an estimate for how many values are less than or equal to the given value
 - To estimate the number that is greater than the value subtract this number from the total frequency
 - They can be used to estimate the interquartile range IQR = $Q_3 Q_1$
 - They can be used to construct a box plot for grouped data



Cumulative Frequency Graphs

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 - They can be used to construct a **box plot** for grouped data



Worked Example

2

The cumulative frequency graph below shows the lengths in cm, I, of 30 puppies in a training group.











Histograms

What is a (frequency) histogram?

- · A frequency histogram clearly shows the frequency of class intervals
 - The classes will have equal class intervals
 - The frequency will be on the y-axis
 - The bar for a class interval will begin at the lower boundary and end at the upper boundary
- A frequency histogram is similar to a bar chart
 - A bar chart is used for qualitative or discrete data and has gaps between the bars
 - A frequency histogram is used for continuous data and has no gaps between bars

What are (frequency) histograms useful for?

- They show the **modal class** clearly
- They show the shape of the distribution
 - It is important the class intervals are of equal width
- They can show whether the variable can be modelled by a normal distribution
 - If the shape is symmetrical and bell-shaped



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The table below and its corresponding histogram show the mass, in kg, of some new born bottlenose dolphins.

Mass, <i>m</i> kg	Frequency
$4 \le m < 8$	4
$8 \le m < 12$	15
$12 \le m < 16$	19
$16 \le m < 20$	10
$20 \le m \le 24$	6

a)

Draw a frequency histogram to represent the data.





4.1.8 Interpreting Data

Interpreting Data

How do l interpret statistical measures?

- The mode is useful for qualitative data
 - It is not as useful for quantitative data as there is not always a unique mode
- The mean includes all values
 - It is affected by outliers
 - A smaller/larger mean is preferable depending on the scenario
 - A smaller mean time for completing a puzzle is better
 - A bigger mean score on a test is better
- The median is not affected by outliers
 - It does not use all the values
- The range gives the full spread of the all of the data
 - It is affected by outliers
- The interquartile range gives the spread of the middle 50% about the median and is not affected by outliers
 - It does not use all the values
 - A bigger IQR means the data is more spread out about the median
 - A smaller IQR means the data is more centred about the median
- The standard deviation and variance use all the values to give a measure of the average spread of the data about the mean
 - They are affected by outliers
 - A bigger standard deviation means the data is more spread out about the mean
 - A smaller standard deviation means the data is more centred about the mean

How do I choose which diagram to use to represent data?

- Box plots
 - Can be used with ungrouped univariate data
 - Shows the range, interquartile range and quartiles clearly
 - Very useful for comparing data patterns quickly

Cumulative frequency graphs

- Can be used with continuous grouped univariate data
- Shows the running total of the frequencies that fall below the upper bound of each class
- Histograms
 - Can be used with continuous grouped univariate data
 - Used with equal class intervals
 - Shows the frequencies of the group
- Scatter diagrams
 - Can be used with ungrouped **bivariate** data
 - Shows the graphical relationship between the variables

How do I compare two or more data sets?

- Compare a measure of central tendency
 - If the data contains outliers use the median
 - If the data is roughly symmetrical use the mean

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- Compare a measure of dispersion
 - If the data contains outliers use the interquartile range
 - If the data is roughly symmetrical use the standard deviation
- Consider whether it is better to have a smaller or bigger average
 - This will depend on the context
 - A smaller average time for completing a puzzle is better
 - A bigger average score on a test is better
- Consider whether it is better to have a smaller or bigger spread
 - Usually a smaller spread means it is more consistent
- Always relate the **comparisons to the context** and consider reasons
 - Consider the sampling technique and the data collection method





4.2 Correlation & Regression

4.2.1 Bivariate Data

Scatter Diagrams

What does bivariate data mean?

- Bivariate data is data which is collected on two variables and looks at how one of the factors affects the other
 - Each data value from one variable will be **paired** with a data value from the other variable
 - The two variables are often related, but do not have to be

What is a scatter diagram?

- A scatter diagram is a way of graphing bivariate data
 - One variable will be on the x-axis and the other will be on the y-axis
 - The variable that can be **controlled** in the data collection is known as the **independent** or **explanatory variable** and is plotted on the *x*-axis
 - The variable that is measured or discovered in the data collection is known as the dependent or response variable and is plotted on the y-axis
- · Scatter diagrams can contain outliers that do not follow the trend of the data

Exam Tip

- If you use scatter diagrams in your Internal Assessment then be aware that finding outliers for bivariate data is different to finding outliers for univariate data
 - (x, y) could be an outlier for the bivariate data even if x and y are not outliers for their separate univariate data



Correlation

What is correlation?

- Correlation is how the two variables change in relation to each other
 - Correlation could be the result of a **causal relationship** but this is not always the case
- Linear correlation is when the changes are proportional to each other
- Perfect linear correlation means that the bivariate data will all lie on a straight line on a scatter diagram
- When describing correlation mention
 - The type of the correlation
 - Positive correlation is when an increase in one variable results in the other variable increasing
 - Negative correlation is when an increase in one variable results in the other variable decreasing
 - No linear correlation is when the data points don't appear to follow a trend
 - The strength of the correlation
 - Strong linear correlation is when the data points lie close to a straight line
 - Weak linear correlation is when the data points are not close to a straight line
- If there is strong linear correlation you can draw a line of best fit (by eye)
 - The line of best fit will pass through the mean point $(\overline{x}, \overline{y})$
 - If you are asked to draw a line of best fit
 - Plot the mean point
 - Draw a line going through it that follows the trend of the data



What is the difference between correlation and causation?

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- It is important to be aware that just because correlation exists, it does not mean that the change in one of the variables is **causing** the change in the other variable
 - Correlation does not imply causation!
- If a change in one variable **causes** a change in the other then the two variables are said to have a **causal relationship**
 - Observing correlation between two variables does not always mean that there is a causal relationship
 - There could be underlying factors which is causing the correlation
 - Look at the two variables in question and consider the context of the question to decide if there could be a causal relationship
 - If the two variables are temperature and number of ice creams sold at a park then it is likely to be a causal relationship
 - Correlation may exist between global temperatures and the number of monkeys kept as pets in the UK but they are unlikely to have a causal relationship





2

A teacher is interested in the relationship between the number of hours her students spend on a phone per day and the number of hours they spend on a computer. She takes a sample of nine students and records the results in the table below.

Hours spent on a phone per day	7.6	7.0	8.9	3.0	3.0	7.5	2.1	1.3	5.8
Hours spent on a computer per day	1.7	1.1	0.7	5.8	5.2	1.7	6.9	7.1	3.3



b)

Draw a scatter diagram for the data.



Describe the correlation.









4.2.2 Correlation Coefficients

PMCC

What is Pearson's product-moment correlation coefficient?

- Pearson's product-moment correlation coefficient (PMCC) is a way of giving a numerical value to a **linear relationship** of bivariate data
- The PMCC of a sample is denoted by the letter r
 - $r \operatorname{can} take any value such that <math>-1 \le r \le 1$
 - A positive value of r describes positive correlation
 - A negative value of r describes negative correlation
 - r = 0 means there is no linear correlation
 - r=1 means perfect positive linear correlation
 - r = -1 means perfect negative linear correlation
 - The closer to 1 or -1 the stronger the correlation



How do I calculate Pearson's product-moment correlation coefficient (PMCC)?

- · You will be expected to use the statistics mode on your GDC to calculate the PMCC
- · The formula can be useful to deepen your understanding

$$r = \frac{S_{xy}}{S_x S_y}$$



•
$$S_{xy} = \sum_{i=1}^{n} x_i y_i - \frac{1}{n} \left(\sum_{i=1}^{n} x_i \right) \left(\sum_{i=1}^{n} y_i \right)$$
 is linked to the **covariance**
• $S_x = \sqrt{\sum_{i=1}^{n} x_i^2 - \frac{1}{n} \left(\sum_{i=1}^{n} x_i \right)^2}$ and $S_y = \sqrt{\sum_{i=1}^{n} y_i^2 - \frac{1}{n} \left(\sum_{i=1}^{n} y_i \right)^2}$ are linked to the **covariance**

are linked to the

variances

• You do not need to learn this as using your GDC will be expected

When does the PMCC suggest there is a linear relationship?

- Critical values of r indicate when the PMCC would suggest there is a linear relationship
 - In your exam you will be given critical values where appropriate
 - Critical values will depend on the size of the sample
- If the **absolute value** of the **PMCC** is **bigger** than the **critical value** then this suggests a linear model is appropriate





Spearman's Rank

What is Spearman's rank correlation coefficient?

- Spearman's rank correlation coefficient is a measure of how well the relationship between two variables can be described using a **monotonic** function
 - Monotonic means the points are either always increasing or always decreasing
 - This can be used as a way to measure correlation in linear models
 - Though Spearman's Rank correlation coefficient can also be used to assess a nonlinear relationship
- · Each data is ranked, from biggest to smallest or from smallest to biggest
 - For n data values, they are ranked from 1 to n
 - It doesn't matter whether variables are ranked from biggest to smallest or smallest to biggest, but they must be ranked in the **same order for both variables**
- Spearman's rank of a sample is denoted by r
 - $r_{\rm s}$ can take any value such that $-1 \le r_{\rm s} \le 1$
 - A positive value of rs describes a degree of agreement between the rankings
 - A negative value of rs describes a degree of disagreement between the rankings
 - $r_s = 0$ means the data shows **no monotonic behaviour**
 - $r_s = 1$ means the rankings are in complete agreement: the data is strictly increasing
 - An increase in one variable means an increase in the other
 - r_s = -1 means the rankings are in complete disagreement: the data is strictly decreasing
 - An increase in one variable means a decrease in the other
 - The closer to 1 or -1 the stronger the correlation of the rankings



How do I calculate Spearman's rank correlation coefficient (PMCC)?

- · Rank each set of data independently
 - 1 to n for the x-values
 - I to n for the y-values
- If some values are equal then give each the average of the ranks they would occupy



 For example: if the 3rd, 4th and 5th highest values are equal then give each the ranking of 4

$$\frac{3+4+5}{3} = 4$$

- Calculate the PMCC of the **rankings** using your GDC
 - This value is Spearman's rank correlation coefficient





Appropriateness & Limitations

Which correlation coefficient should I use?

- Pearson's PMCC tests for a linear relationship between two variables
 - It will not tell you if the variables have a non-linear relationship
 - Such as exponential growth
 - Use this if you are interested in a linear relationship
- Spearman's rank tests for a monotonic relationship (always increasing or always decreasing) between two variables
 - It will not tell you what function can be used to model the relationship
 - Both linear relationships and exponential relationships can be monotonic
 - Use this if you think there is a non-linear monotonic relationship

How are Pearson's and Spearman's correlation coefficients connected?

- If there is linear correlation then the relationship is also monotonic
 - $\circ r = 1 \Rightarrow r_s = 1$
 - $\circ r = -1 \Rightarrow r_s = -1$
 - However the converse is not true



- It is possible for Spearman's rank to be 1 (or -1) but for the PMCC to be different
 - For example: data that follows an exponential growth model
 - $r_s = 1$ as the points are always increasing
 - r < 1 as the points do not lie on a straight line

Are Pearson's and Spearman's correlation coefficients affected by outliers?

- Pearson's PMCC is affected by outliers
 - as it uses the numerical value of each data point
- Spearman's rank is **not usually** affected by outliers
 - as it only uses the ranks of each data point



Exam Tip

• You can use your GDC to plot the scatter diagram to help you visualise the data



The table below shows the scores of eight students for a maths test and an English test.

Maths (x)	7	18	37	52	61	68	75	82
English (y)	5	3	9	12	17	41	49	97

a)

Write down the value of Pearson's product-moment correlation coefficient, r.

Enter data into GDC. r = 0.79433... r= 0.794 (3sf)

b)

Find the value of Spearman's rank correlation coefficient, r_{e} .

Rank the data x rank 8 7 6 5 4 3 2 1 y rank 7 8 6 5 4 3 2 1 y rank 7 8 6 5 94 3 2 CE Find PMCC of ranks $r_s = 0.97619...$ $r_s = 0.976 (3sf)$

c)

Comment on the values of the two correlation coefficients.

The value of r suggests there is strong positive linear correlation. The value of rs suggests strong positive correlation, which is not necessarily linear.



4.2.3 Linear Regression

Linear Regression

What is linear regression?

- If strong linear correlation exists on a scatter diagram then the data can be modelled by a linear model
 - Drawing lines of best fit by eye is not the best method as it can be difficult to judge the best position for the line
- The least squares regression line is the line of best fit that minimises the sum of the squares of the gap between the line and each data value
 - This is usually called the **regression line of y on x**
 - It can be calculated by looking at the vertical distances between the line and the data values
- The **regression line of y on x** is written in the form y = ax + b
- a is the gradient of the line
 - It represents the change in y for each individual unit change in x
 - If a is **positive** this means y **increases** by a for a unit increase in x
 - If a is **negative** this means y **decreases** by |a| for a unit increase in x
- b is the y intercept
 - It shows the value of y when x is zero
- You are expected to use your GDC to find the equation of the regression line
 - Enter the bivariate data and choose the model "ax + b" T C F
 - Remember the **mean point** $(\overline{x}, \overline{y})$ will lie on the regression line

How do I use a regression line?

- The equation of the regression line can be used to decide what type of correlation there is if there is no scatter diagram
 - If a is positive then the data set has positive correlation
 - If a is negative then the data set has negative correlation
- The equation of the regression line can also be used to **predict** the value of a **dependent variable (y)** from an **independent variable (x)**
 - The equation should **only be used** to make **predictions for y**
 - Using a y on x line to predict x is not always reliable
 - $\circ~$ Making a prediction within the range of the given data is called interpolation
 - This is usually reliable
 - The stronger the correlation the more reliable the prediction
 - $\circ~$ Making a prediction $outside \, of \, the \, range \, of \, the \, given \, data \, is \, called \, extrapolation$
 - This is much less reliable
 - The prediction will be more reliable if the number of data values in the original sample set is bigger





- Once you calculate the values of a and b store then in your GDC
 - This means you can use the full display values rather than the rounded values when using the linear regression equation to predict values
 - This avoids rounding errors





Barry is a music teacher. For 7 students, he records the time they spend practising per week (x hours) and their score in a test (y %).

Time (x)	2	5	6	7	10	11	12
Score (y)	11	49	55	75	63	68	82

a)

Write down the equation of the regression line of y on x, giving your answer in the form y = ax + b where a and b are constants to be found.



Substitute x = 15 $y = (5.5680...) \times 15 + (15.4136...) = 98.93..$ The model predicts a score of 98.9% but this is unreliable as x = 15 is outside the range of data. Therefore extrapolation is being used.



4.3 Further Correlation & Regression

4.3.1 Non-linear Regression

Non-linear Regression

What is non-linear regression?

- You have already seen that **linear regression** is when you can use a straight line to fit bivariate data
- Non-linear regression is when you can use a curve (rather than a straight line) to fit bivariate data
- · In your exam the regression could be:
 - Linear: y = ax + b
 - Quadratic: $y = ax^2 + bx + c$
 - Cubic: $y = ax^3 + bx^2 + cx + d$
 - Exponential: $y = ab^x$ or $y = ae^{bx}$
 - Power: $y = ax^b$
 - Sine: $y = a\sin(bx + c) + d$

How do I find the equation of the non-linear regression model?

PRACIICE

- Using your GDC:
 - Type the two sets of the data into your GDC
 - Select the relevant model
 - The exam question will tell you which model to use
 - Your GDC will calculate the constants
- · You can use logarithms to linearise exponential and power relationships
 - Power: $y = ax^{b}$ then $\ln y = \ln a + b \ln x$
 - Iny and Inx will have a linear relationship
 - Exponential: $y = ab^x$ then $\ln y = \ln a + x \ln b$
 - lny and x will have a linear relationship

Exam Tip

- You can use your GDC to plot the scatter diagram and include the graph of a regression model
 - This will allow you to get a sense of how well the model fits the data



Scarlett and Violet collect data on the length of a film (x minutes) and the audience rating (y %).

X	75	93	101	107	115	124	132	140	171
у	83	75	51	38	47	56	76	91	70

a)

Scarlett claims that there is a cubic relationship. Find the equation of a cubic regression model of the form $y = ax^3 + bx^2 + cx + d$.

Type the data into GDC and choose the cubic regression model a = -0.0005291... b = 0.2030... c = -24.89... d = 1037.7...

b)

Violet claims that there is a sine relationship. Find the equation of a sine regression model of the form $y = a\sin(bx + c) + d$.

C)

Whose model predicts a higher audience rating for a film which is 100 minutes long?





Least Squares Regression Curves

What is a residual?

- Given a set of *n* pairs of data and a **regression model** y = f(x)
- A **residual** is the **actual y-value** (from the data) **minus** the **predicted y-value** (using the regression model)

 $\circ y_i - f(x_i)$

+ The sum of the square residuals is denoted by $SS_{\rm res}$

•
$$SS_{res} = \sum_{i=1}^{n} (y_i - f(x_i))^2$$

- If you have two regression models using the same data then the one with the smaller SS_{res} fits the data better

What is a least squares regression curve?

- The least squares regression curve can be thought of as a "curve of best fit" y = f(x)
- For a given type of model the least squares regression curve minimises the sum of the square residuals
 - · Your GDC calculates the constants for the least squares regression curves

Why is the sum of the square residuals not always a good measure of fit?

- If two models are formed using the **same number of pairs** of data then the sum of the square residuals is a **good measure of fit**
- If two models use different number of pairs of data then SS_{res} is not always a good

measure of fit

- The sum will increase with more pairs of data and so can no longer be compared against a data set with a different number of pairs
- Compare the two scenarios
 - 10 pairs of data and the absolute value of each residual is 15 then $SS_{res} = 10 \times 15^2 = 2250$
 - 2250 pairs of data and the absolute value of each residual is 1 then $SS_{res} = 2250 \times 1^2 = 2250$
- $\circ~$ They have the same value of $S\!S_{\it res}$ but the residuals in the second scenario are much smaller
- Your GDC may give you the mean squared error

•
$$MSe = \frac{1}{n}SS_{res} = \frac{1}{n}\sum_{i=1}^{n}(y_i - f(x_i))^2$$

- This is a better measure of fit
- You do not need to know this for your exam but it might help with your understanding



Jet is the owner of a gym and he is testing different prices options. The table below shows the number of new members per month (M) and the price of a monthly membership ($\pounds p$).

р	10	20	30
М	97	68	55

Jet believes that he can fit the data with either the model $M_1(p) = \frac{2700}{p+20}$ or the

model $M_2(p) = \frac{2100}{p+10}$.

Jet wants to choose the model with the smallest value for the sum of square residuals.

Determine which model Jet should choose.

Calculate the predicted values.

For
$$M_1 : 55_{res} = (97 - 90)^2 + (68 - 67.5)^2 + (55 - 54)^2 = 50.25$$

For $M_2 : 55_{res} = (97 - 105)^2 + (68 - 70)^2 + (55 - 52.5)^2 = 74.25$



The Coefficient of Determination

What is the coefficient of determination?

- The coefficient of determination is a measure of fit for a model
 - If the coefficient of determination is 0.57 this means 57% of the variation of the yvariable can be explained by the variation in the x-variable
 - The other 43% can be explained by other factors
 - The higher this proportion the more the model fits the data
- The coefficient of determination is denoted by R²
 - o R² ≤ 1
 - R² = 1 means the model is a **perfect fit** for the data
 - The closer to 1 the better the fit
 - R² is usually greater than or equal to zero
 - R² can be negative but this is outside the scope of this course
- If the regression model is linear then the coefficient of determination is equal to square of the PMCC
 - $R^2 = r^2$ for linear models
 - Some GDCs will simply denote R² as r² due to its connection to the PMCC for linear models

How do I calculate the coefficient of determination?

- When finding the constants for regression models your GDC might give you the value of R^2
 - · You will only be asked to calculate the coefficient of determination for models for which GDCs give the value of R² EPS PRACTICE
- The coefficient of determination can be calculated by

$$\circ R^2 = 1 - \frac{SS_{res}}{SS_{tot}}$$

• Where
$$SS_{tot} = \sum_{i=1}^{n} (y_i - \overline{y})^2$$

You do not need to know this formula but it might help with your understanding

Does the coefficient of determination determine the validity of a model?

- If R² is close to 1 then the model fits the data well
 - However this alone does not guarantee that it is a good model for the relationship between the two variables
- Consider the scenario where there are big gaps between data points and a model which fits the data well
 - The model only fits the data at the data points
 - As there are gaps between the data points the model might not be a good fit for these areas
- Different types of models have different number of parameters
 - · Therefore using different types of models to fit the same data will have different levels of accuracy
 - Linear models need at least two pairs of data

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- Quadratic models need at least three pairs of data
- Cubic models need at least four pairs of data
 - Using four pairs of data will mean the cubic model will have R² = 1 This is because the cubic graph will go through all four pieces of data – the value is likely to decrease as extra pairs of data are included
 - However this does not mean it is a better fit than the quadratic model
 - The quadratic model could be more accurate as it has one more pair of data than is needed





Data is collected on the lengths of cheetahs (x metres) and their average running speeds (y ms⁻¹).

X	1.21	1.33	1.12	1.45	1.42	1.39	1.24	1.19	1.32
у	24.3	25.1	22.2	35.1	35.1	33.4	27.1	23.1	24.8

a)

Find the equation of the least squares regression curve using:

(i)

a quadratic model $y = ax^2 + bx + c$.

(ii)

an exponential model $y = ab^x$.



b)

Based solely on the coefficients of determination, suggest which model is better fit for the data.

```
Find the coefficients of determination using GDC
Quadratic R^2 = 0.86429...
Exponential R^2 = 0.80157...
Based on the coefficients of determination, the quadratic
regression model as its R^2 value is bigger.
```

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4.3.2 Logarithmic Scales

Logarithmic Scales

What are logarithmic scales?

- Logarithmic scales are scales where intervals increase exponentially
 - A normal scale might go 1, 2, 3, 4, ...
 - A logarithmic scale might go 1, 10, 100, 1000, ...
- Sometimes we can keep the scales with constant intervals by changing the variables
 - If the values of x increase exponentially: 1, 10, 100, 1000, ...
 - Then you can use the variable log x instead which will have the scale: 1, 2, 3, 4, ...
 - This will change the shape of the graph
 - If the graph transforms to a straight line then it is easier to analyse
- Any base can be used for logarithmic scales
 - The most common bases are 10 and e

Why do we use logarithmic scales?

- For variables that have a large range it can be difficult to plot on one graph
 - Especially when a lot of the values are clustered in one region
 - For example: populations of countries
 - This can range from 800 to 1450 000 000
- If we are interested in the **rate of growth** of a variable rather than the actual values then a logarithmic scale is useful



log-log & semi-log Graphs

What is a log-log graph?

- A log-log graph is used when both scales of the original graph are logarithmic
 - You transform both variables by taking logarithms of the values
- log y & log x will be used instead of y & x
- Power graphs ($y = ax^b$) look like straight lines on log-log graphs

What is a semi-log graph?

- A **semi-log graph** is used when **only one scale** (the *y*-axis) of the original graph are logarithmic
 - You transform only the y-variable by taking logarithms of those values
- log y will be used instead of y
- Exponential graphs ($y = ab^x$) look like straight lines on semi-log graphs

How can I estimate values using log-log and semi-log graphs?

- · Identify whether one or both of the scales are logarithmic
- Identify the variable so that the scales have equal intervals
 - x:1,10,100,1000,...uselogx
 - For x: 1, e, e², e³, ... use ln x
- If you are asked to estimate a value:
 - First find the value of any logarithms
 - For example: log y, ln x, etc
 - Use the graph to read off the value
 - If it is a value for a logarithm find the actual value using:
 - $\log x = k \Rightarrow x = 10^k$
 - $\ln x = k \Rightarrow x = e^k$

Exam Tip

• Pay close attention to which base is being used (log or ln)







4.3.3 Linearising using Logarithms

Exponential Relationships

How do I use logarithms to linearise exponential relationships?

- Graphs of exponential functions appear as straight lines on semi-log graphs
- Suppose y = ab^x
 - · You can take logarithms of both sides
 - $\ln y = \ln(ab^x)$
 - · You can split the right hand side into the sum of two logarithms
 - $\ln y = \ln a + \ln(b^x)$
 - You can bring down the power in the final term
 - $\ln y = \ln a + x \ln b$
- $\ln y = \ln a + x \ln b$ is in linear form Y = mX + c
 - $\circ Y = \ln y$
 - $\circ X = x$
 - $\circ m = \ln b$
 - $\circ c = \ln a$



How can I use linearised data to find the values of the parameters in an exponential model $y = ab^{x}$?

- STEP 1: Linearise the data using $Y = \ln y$ and X = x
- STEP 2: Find the equation of the regression line of Y on X: Y = mX + c
- STEP 3: Equate coefficients between Y = mX + c and $\ln y = \ln a + x \ln b$
 - $\circ m = \ln b$
 - $\circ c = \ln a$
- STEP 4: Solve to find a and b
 - $a = e^{c}$
 - $\circ b = e^m$



Hatter has noticed that over the past 50 years there seems to be fewer hatmakers in London. He also knows that global temperatures have been rising over the same time period. He decides to see if there could be any correlation, so he collects data on the number of hatmakers and the global mean temperatures from the past 50 years and records the information in the graph below.



Hatter suggests that the equation for h in terms of t can be written in the form $h = ab^t$

. He linearises the data using x = t and $y = \ln h$ and calculates the regression line of y on x to be y = 4.382 - 1.005x.

Find the values of a and b.



Power Relationships

How do I use logarithms to linearise power relationships?

- Graphs of **power functions** appear as straight lines on **log-log graphs**
- Suppose $y = ax^b$
 - You can take logarithms of both sides
 - $\ln y = \ln(ax^b)$
 - You can split the right hand side into the sum of two logarithms
 - $\ln y = \ln a + \ln(x^b)$
 - You can bring down the power in the final term
 - $\ln y = \ln a + b \ln x$
- $\ln y = \ln a + b \ln x$ is in linear form Y = mX + c
 - $\circ Y = \ln y$
 - $\circ X = \ln x$
 - $\circ m = b$
 - $\circ c = \ln a$

How can I use linearised data to find the values of the parameters in an power model $y = ax^b$?

- STEP 1: Linearise the data using $Y = \ln y$ and $X = \ln x$
- STEP 2: Find the equation of the regression line of Y on X: Y = mX + c
- STEP 3: Equate coefficients between Y = mX + c and $\ln y = \ln a + b \ln x$

$$\circ m = b$$

- $\circ c = \ln a$
- STEP 4: Solve to find a and b
 - $a = e^{c}$
 - $\circ b = m$







4.4 Probability

4.4.1 Probability & Types of Events

Probability Basics

What key words and terminology are used with probability?

- An experiment is a repeatable activity that has a result that can be observed or recorded
 Trials are what we call the repeats of the experiment
- An outcome is a possible result of a trial
- An event is an outcome or a collection of outcomes
 - Events are usually denoted with capital letters: A, B, etc
 - n(A) is the number of outcomes that are included in event A
 - An event can have one or more than one outcome
- A sample space is the set of all possible outcomes of an experiment
 - This is denoted by U
 - n(U) is the total number of outcomes
 - It can be represented as a list or a table

How do I calculate basic probabilities?

- If all outcomes are **equally likely** then probability for each outcome is the same
 - Probability for each outcome is $\frac{1}{n(U)}$
- Theoretical probability of an event can be calculated without using an experiment by dividing the number of outcomes of that event by the total number of outcomes

$$P(A) = \frac{n(A)}{n(U)}$$

- $\circ~$ This is given in the $\mathbf{formula\ booklet}$
- · Identifying all possible outcomes either as a list or a table can help
- Experimental probability (also known as relative frequency) of an outcome can be calculated using results from an experiment by dividing its frequency by the number of trials
 - **Relative frequency** of an outcome is $\frac{\text{Frequency of that outcome from the trials}}{\text{Total number of trials } (n)}$

How do I calculate the expected number of occurrences of an outcome?

- Theoretical probability can be used to calculate the expected number of occurrences of an outcome from *n* trials
- If the probability of an outcome is p and there are n trials then:
 - The expected number of occurrences is **np**
 - This does not mean that there will exactly np occurrences
 - If the experiment is repeated multiple times then we expect the number of occurrences to average out to be *np*

What is the complement of an event?

- The probabilities of all the outcomes add up to 1
- Complementary events are when there are two events and exactly one of them will occur
 One event has to occur but both events can not occur at the same time
- The complement of event A is the event where event A does not happen

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- This can be thought of as not A
- This is denoted A'

$$P(A) + P(A') = 1$$

- This is in the formula booklet
- It is commonly written as P(A') = 1 P(A)

What are different types of combined events?

- The intersection of two events (A and B) is the event where both A and B occur
 - This can be thought of as **A and B**
 - \circ This is denoted as $A \cap B$
- The union of two events (A and B) is the event where A or B or both occur
 - This can be thought of as A or B
 - This is denoted $A \cup B$
- The event where A occurs given that event B has occurred is called conditional probability
 - This can be thought as **A given B**
 - This is denoted $A \mid B$

How do I find the probability of combined events?

• The probability of A or B (or both) occurring can be found using the formula

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

- This is given in the formula booklet
- You subtract the probability of A and B both occurring because it has been included twice (once in P(A) and once in P(B))
- The probability of A and B occurring can be found using the formula

$P(A \cap B) = P(A)P(B|A)$

- A rearranged version is given in the formula booklet
- Basically you multiply the probability of A by the probability of B then happening

\mathbf{O}

Exam Tip

• In an exam drawing a Venn diagram or tree diagram can help even if the question does not ask you to



Dave has two fair spinners, A and B. Spinner A has three sides numbered 1, 4, 9 and spinner B has four sides numbered 2, 3, 5, 7. Dave spins both spinners and forms a two-digit number by using the spinner A for the first digit and spinner B for the second digit.

T is the event that the two-digit number is a multiple of 3.

a)

List all the possible two-digit numbers.





Independent & Mutually Exclusive Events

What are mutually exclusive events?

- Two events are mutually exclusive if they cannot both occur
 - For example: when rolling a dice the events "getting a prime number" and "getting a 6" are mutually exclusive
- If A and B are mutually exclusive events then:
 - $\circ P(A \cap B) = 0$

What are independent events?

- Two events are independent if one occurring does not affect the probability of the other occurring
 - For example: when flipping a coin twice the events "getting a tails on the first flip" and "getting a tails on the second flip" are independent
- If A and B are independent events then:

• P(A|B) = P(A) and P(B|A) = P(B)

- If A and B are independent events then:
 - $\circ P(A \cap B) = P(A)P(B)$
 - This is given in the formula booklet
 - This is a useful formula to test whether two events are statistically independent

How do I find the probability of combined mutually exclusive events?

- If A and B are **mutually exclusive** events then
 - PAPERS PRACTICE $P(A \cup B) = P(A) + P(B)$
 - This is given in the formula booklet
 - This occurs because $P(A \cap B) = 0$
- For any two events A and B the events A ∩ B and A ∩ B' are **mutually exclusive** and A is the **union** of these two events
 - $\circ P(A) = P(A \cap B) + P(A \cap B')$
 - This works for any two events A and B





a)

A student is chosen at random from a class. The probability that they have a dog is 0.8, the probability they have a cat is 0.6 and the probability that they have a cat or a dog is 0.9.

Find the probability that the student has both a dog and a cat.

Let D be event "has a dog" and C be "has a cat" $P(D \cup C) = P(D) + P(C) - P(D \cap C)$ $0.9 = 0.8 + 0.6 - P(D \cap C)$ $P(D \cap C) = 0.5$

b)

Two events, Q and R, are such that P(Q) = 0.8 and $P(Q \cap R) = 0.1$. Given that Q and R are independent, find P(R).



c)

Two events, S and T, are such that P(S) = 2P(T).

Given that S and T are mutually exclusive and that $P(S \cup T) = 0.6$ find P(S) and P(T).

S and T mutually exclusive \Rightarrow P(S u T) = P(S) + P(T) 0.6 = P(S) + P(T) 0.6 = 2P(T) + P(T) P(S) = 2P(T) 0.6 = 3P(T) P(T) = 0.2 and P(S) = 0.4



4.4.2 Conditional Probability

Conditional Probability

What is conditional probability?

- Conditional probability is where the probability of an event happening can vary depending on the outcome of a prior event
- The event A happening given that event B has happened is denoted A|B
- A common example of conditional probability involves selecting multiple objects from a bag without replacement
 - The probability of selecting a certain item changes depending on what was selected before
 - This is because the total number of items will change as they are not replaced once they have been selected

How do I calculate conditional probabilities?

- · Some conditional probabilities can be calculated by using counting outcomes
 - Probabilities without replacement can be calculated like this
 - For example: There are 10 balls in a bag, 6 of them are red, two of them are selected without replacement
 - To find the probability that the second ball selected is red given that the first one is red count how many balls are left:
 - A red one has already been selected so there are 9 balls left and 5 are red so the probability is $\frac{5}{9}$
- You can use sample space diagrams to find the probability of A given B:
 - reduce your sample space to just include outcomes for event B
 - find the proportion that also contains outcomes for event A
- There is a formula for conditional probability that you can use

•
$$P(A | B) = \frac{P(A \cap B)}{P(B)}$$

- This is given in the formula booklet
- This can be rearranged to give $P(A \cap B) = P(B)P(A \mid B)$


Worked Example

In a class of 30 students: 19 students have a dog, 17 students have a cat and 11 have both a dog and a cat. One student is selected at random.

a)

Find the probability that the student has a dog.

Let D be event "has a dog" and C be "has a cat" $P(D) = \frac{n(D)}{n(U)} \leftarrow N$ Number who have dogs $\overline{n(U)} \leftarrow T$ otal number of students $P(D) = \frac{19}{3D}$

b)

Find the probability that the student has a dog given that they have a cat.





4.4.3 Sample Space Diagrams

Venn Diagrams

What is a Venn diagram?

- A Venn diagram is a way to illustrate **events** from an **experiment** and are particularly useful when there is an overlap between possible **outcomes**
- A Venn diagram consists of
 - a rectangle representing the sample space (U)
 - The rectangle is labelled U
 - Some mathematicians instead use S or ξ
 - a circle for each event
 - Circles may or may not overlap depending on which outcomes are shared between events
- The numbers in the circles represent either the **frequency** of that event or the **probability** of that event
 - If the frequencies are used then they should add up to the total frequency
 - If the probabilities are used then they should add up to 1

What do the different regions mean on a Venn diagram?

- A' is represented by the regions that are **not in** the A circle
- $A \cap B$ is represented by the region where the A and B circles overlap
- A U B is represented by the regions that are in A or B or both
- Venn diagrams show 'AND' and 'OR' statements easily
- Venn diagrams also instantly show mutually exclusive events as these circles will not overlap
- Independent events can not be instantly seen
 - You need to use probabilities to deduce if two events are independent









How do I solve probability problems involving Venn diagrams?

- Draw, or add to a given Venn diagram, filling in as many values as possible from the information provided in the question
- It is usually helpful to work from the centre outwards
 - Fill in intersections (overlaps) first
- If two events are independent you can use the formula
 - $\circ P(A \cap B) = P(A)P(B)$
- To find the conditional probability P(A | B)
 - Add together the frequencies/probabilities in the B circle
 - This is your denominator
 - Out of those frequencies/probabilities add together the ones that are also in the A circle
 - This is your numerator
 - Evaluate the fraction PAPERS PRACTICE







Exam Tip

- If you struggle to fill in a Venn diagram in an exam:
 - Label the missing parts using algebra
 - Form equations using known facts such as:
 - the sum of the probabilities should be 1
 - P(A∩B)=P(A)P(B) if A and B are independent events









Tree Diagrams

What is a tree diagram?

- A tree diagram is another way to show the outcomes of combined events
 - They are very useful for intersections of events
- The events on the branches must be **mutually exclusive**
 - Usually they are an event and its complement
- The probabilities on the second sets of branches **can depend** on the outcome of the first event
 - These are conditional probabilities
- When selecting the items from a bag:
 - The second set of branches will be the **same** as the first if the items **are replaced**
 - The second set of branches will be the **different** to the first if the items **are not replaced**

How are probabilities calculated using a tree diagram?

- To find the probability that two events happen together you **multiply** the corresponding probabilities on their branches
 - It is helpful to find the probability of all combined outcomes once you have drawn the tree
- To find the probability of an event you can:
 - add together the probabilities of the combined outcomes that are part of that event
 For example: P(A ∪ B) = P(A ∩ B) + P(A ∩ B') + P(A' ∩ B)
 - **subtract** the probabilities of the combined outcomes that are not part of that event from 1



Do I have to use a tree diagram?

- If there are multiple events or trials then a tree diagram can get big
- You can break down the problem by using the words **AND/OR/NOT** to help you find probabilities without a tree



• You can speed up the process by only drawing parts of the tree that you are interested in

Which events do I put on the first branch?

- If the events A and B are independent then the order does not matter
- If the events A and B are not independent then the order does matter
 - If you have the probability of **A given B** then put **B on the first set** of branches
 - If you have the probability of **B given A** then put **A on the first set** of branches

Exam Tip

- In an exam do not waste time drawing a full tree diagram for scenarios with lots of events unless the question asks you to
 - · Only draw the parts that you are interested in







Worked Example

20% of people in a company wear glasses. 40% of people in the company who wear glasses are right-handed. 50% of people in the company who don't wear glasses are right-handed.

a)

Draw a tree diagram to represent the information.



c)

Given that a person who is right-handed is selected at random, find the probability that they wear glasses.

$$P(A|R) = \frac{P(AR)}{P(R)} = \frac{0.08}{0.48}$$

 $P(A|R) = \frac{1}{6}$



4.5 Probability Distributions

4.5.1 Discrete Probability Distributions

Discrete Probability Distributions

What is a discrete random variable?

- A random variable is a variable whose value depends on the outcome of a random event
 - The value of the random variable is not known until the event is carried out (this is what is meant by 'random' in this case)
- Random variables are denoted using upper case letters (X, Y, etc)
- Particular outcomes of the event are denoted using lower case letters (x, y, etc)
- P(X=x) means "the probability of the random variable X taking the value x"
- A discrete random variable (often abbreviated to DRV) can only take certain values within a set
 - Discrete random variables usually count something
 - Discrete random variables usually can only take a finite number of values but it is possible that it can take an infinite number of values (see the examples below)
- Examples of discrete random variables include:
 - The number of times a coin lands on heads when flipped 20 times
 - this has a finite number of outcomes: {0,1,2,...,20}
 - The number of emails a manager receives within an hour CTICE
 - this has an infinite number of outcomes: {1,2,3,...}
 - The number of times a dice is rolled until it lands on a 6
 - this has an infinite number of outcomes: {1,2,3,...}
 - The number that a dice lands on when rolled once
 - this has a finite number of outcomes: {1,2,3,4,5,6}

What is a probability distribution of a discrete random variable?

- A discrete probability distribution fully describes all the values that a discrete random variable can take along with their associated probabilities
 - This can be given in a **table**
 - Or it can be given as a **function** (called a discrete probability distribution function or "pdf")
 - They can be represented by **vertical line graphs** (the possible values for along the horizontal axis and the probability on the vertical axis)
- The sum of the probabilities of all the values of a discrete random variable is 1
 - This is usually written $\sum P(X=x) = 1$
- A **discrete uniform distribution** is one where the random variable takes a finite number of values each with an **equal probability**

• If there are n values then the probability of each one is $\frac{1}{n}$





How do I calculate probabilities using a discrete probability distribution?

- · First draw a table to represent the probability distribution
 - If it is given as a function then find each probability
 - If any probabilities are unknown then use algebra to represent them
- Form an equation using $\sum P(X = x) = 1$
 - · Add together all the probabilities and make the sum equal to 1
- To find P(X=k)
 - If k is a possible value of the random variable X then P(X = k) will be given in the table
 - If k is not a possible value then P(X = k) = 0
- To find $P(X \le k)$
 - Identify all possible values, x_i , that X can take which satisfy $x_i \le k$
 - Add together all their corresponding probabilities

•
$$P(X \le k) = \sum_{x_i \le k} P(X = x_i)$$

- Some mathematicians use the notation F(x) to represent the cumulative distribution
 F(x) = P(X ≤ x)
- Using a similar method you can find P(X < k), P(X > k) and $P(X \ge k)$
- As all the probabilities add up to 1 you can form the following equivalent equations:
 - P(X < k) + P(X = k) + P(X > k) = 1
 - $\circ P(X > k) = 1 P(X \le k)$
 - $\circ P(X \ge k) = 1 P(X < k)$

How do I know which inequality to use?

- $P(X \le k)$ would be used for phrases such as:
 - At most , no greater than , etc



P(X < k) would be used for phrases such as:

• Fewerthan

- $P(X \ge k)$ would be used for phrases such as:
 - At least , no fewer than , etc
- P(X > k) would be used for phrases such as:
 - Greater than, etc



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 $P(X \leq 3) = \frac{7}{15}$



4.5.2 Expected Values

Expected Values E(X)

What does E(X) mean and how do I calculate E(X)?

- E(X) means the expected value or the mean of a random variable X
 - The expected value does not need to be an obtainable value of X
 - For example: the expected value number of times a coin will land on tails when flipped 5 times is 2.5
- For a **discrete** random variable, it is calculated by:
 - \circ Multiplying each value of X with its corresponding probability
 - Adding all these terms together

$$E(X) = \sum x P(X = x)$$

- This is given in the formula booklet
- Look out for **symmetrical** distributions (where the values of X are symmetrical and their probabilities are symmetrical) as the mean of these is the same as the median
 - For example: if X can take the values 1, 5, 9 with probabilities 0.3, 0.4, 0.3 respectively then by symmetry the mean would be 5

How can I decide if a game is fair?

- Let X be the random variable that represents the gain/loss of a player in a game
 - X will be negative if there is a loss
- Normally the expected gain or loss is calculated by **subtracting** the **cost to play** the game from the **expected value** of the **prize**
- If E(X) is **positive** then it means the player can **expect to make a gain**
- If E(X) is **negative** then it means the player can **expect to make a loss**
- The game is called fair if the expected gain is 0
 - $\circ E(X) = 0$



Worked Example

Daphne pays \$5 to play a game where she wins a prize of \$1, \$5, \$10 or \$100. The random variable W represents the amount she wins and has the probability distribution shown in the following table:

W	1	5	10	100
P(W=w)	0.35	0.5	0.05	0.01

a)

2

Calculate the expected value of Daphne's prize.





4.6 Random Variables

4.6.1 Linear Combinations of Random Variables

Transformation of a Single Variable

What is Var(X)?

- Var(X) represents the variance of the random variable X
- Var(X) can be calculated by the formula
 - $Var(X) = E(X^2) [E(X)]^2$
 - where $E(X^2) = \sum x^2 P(X = x)$
 - You will not be required to use this formula in the exam

What are the formulae for $E(aX \pm b)$ and $Var(aX \pm b)$?

- If a and b are constants then the following formulae are true:
 - $E(aX \pm b) = aE(X) \pm b$
 - $Var(aX \pm b) = a^2 Var(X)$



- These are given in the formula booklet
- This is the same as linear transformations of data
 - The mean is affected by multiplication and addition/subtraction
 - The variance is affected by multiplication but not addition/subtraction
- Remember division can be written as a multiplication

$$\circ \ \frac{X}{a} = \frac{1}{a}X$$







Transformation of Multiple Variables

What is the mean and variance of aX + bY?

- Let X and Y be two random variables and let a and b be two constants
- E(aX + bY) = aE(X) + bE(Y)
 - This is true for **any random variables** X and Y
- $Var(aX+bY) = a^2 Var(X) + b^2 Var(Y)$
 - This is true if X and Y are **independent**
- E(aX bY) = aE(X) bE(Y)
- Var(aX bY) = a² Var(X) + b² Var(Y)
 - Notice that you still add the two terms together on the right hand side
 This is because b² is positive even if b is negative
 - Therefore the variances of aX + bY and aX bY are the same

What is the mean and variance of a linear combination of *n* random variables?

• Let X₁, X₂, ..., X_n be n random variables and a₁, a₂, ..., a_n be n constants

$$E(a_{1}X_{1} \pm a_{2}X_{2} \pm ... \pm a_{n}X_{n}) = a_{1}E(X_{1}) \pm a_{2}E(X_{2}) \pm ... \pm a_{n}E(X_{n})$$

- This is given in the formula booklet
- This can be written as $E(\sum a_i X_i) = \sum a_i E(X_i)$
- This is true for any random variable

$$\operatorname{Var}(a_{1}X_{1} \pm a_{2}X_{2} \pm \dots \pm a_{n}X_{n}) = a_{1}^{2}\operatorname{Var}(X_{1}) + a_{2}^{2}\operatorname{Var}(X_{2}) + \dots + a_{n}^{2}\operatorname{Var}(X_{n})$$

- This is given in the formula booklet
- This can be written as $\operatorname{Var}\left(\sum a_i X_i\right) = \sum a_i^2 \operatorname{Var}(X_i)$
- This is true if the random variables are independent
 - Notice that the constants get squared so the terms on the right-hand side will always be positive

For a given random variable X, what is the difference between 2X and $X_1 + X_2$?

- 2X means one observation of X is taken and then doubled
- X1 + X2 means two observations of X are taken and then added together
- 2X and X₁ + X₂ have the same expected values
 - E(2X) = 2E(X)
 - $E(X_1 + X_2) = E(X_1) + E(X_2) = 2E(X)$
- 2X and X1 + X2 have different variances
 - Var(2X) = 2²Var(X) = 4Var(X)
 - $Var(X_1 + X_2) = Var(X_1) + Var(X_2) = 2Var(X)$
- To see the distinction:
 - Suppose X could take the values 0 and 1
 - 2X could then take the values 0 and 2
 - $X_1 + X_2$ could then take the values 0, 1 and 2

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- · Questions are likely to describe the variables in content
 - For example: The mass of a carton containing 6 eggs is the mass of the carton plus the mass of the 6 **individual** eggs
 - This can be modelled by $M = C + E_1 + E_2 + E_3 + E_4 + E_5 + E_6$ where
 - C is the mass of a carton
 - Eis the mass of an egg
 - It is not C + 6E because the masses of the 6 eggs could be different

Exam Tip

- In an exam when dealing with multiple variables ask yourself which of the two cases is true
 - You are adding together **different observations** using the same variable: $X_1 + X_2 + ... + X_n$
 - You are taking a **single observation** of a variable and multiplying it by a constant: *nX*









4.6.2 Unbiased Estimates

Unbiased Estimates

What is an unbiased estimator of a population parameter?

- An estimator is a random variable that is used to estimate a population parameter
 An estimate is the value produced by the estimator when a sample is used
- · An estimator is called unbiased if its expected value is equal to the population parameter
 - $\circ~$ An estimate from an unbiased estimator is called an unbiased estimate
 - This means that the **mean** of the **unbiased estimates** will get **closer** to the **population parameter** as **more samples** are taken
- The sample mean is an unbiased estimate for the population mean
- The sample variance is not an unbiased estimate for the population variance
 - $\circ~$ On average the sample variance will ${\bf underestimate}$ the population variance
 - As the sample size increases the sample variance gets closer to the unbiased estimate

What are the formulae for unbiased estimates of the mean and variance of a population?

- A sample of *n* data values (x₁, x₂, ... etc) can be used to find unbiased estimates for the mean and variance of the population
- An unbiased estimate for the mean μ of a population can be calculated using

$$\overline{x} = \frac{\sum x}{n}$$

• An unbiased estimate for the variance σ^2 of a population can be calculated using

$$s_{n-1}^2 = \frac{n}{n-1} s_n^2$$

- This is given in the formula booklet
- $\circ s_n^2$ is the variance of the sample data

•
$$s_n^2 = \frac{\sum (x - \bar{x})^2}{n} = \frac{\sum x^2}{n} - (\bar{x})^2$$

• Different calculators can use different notations for s_{n-1}^2

• σ_{n-1}^2 , s^2 , \hat{s}^2 are notations you might see

• You may also see the square roots of these

Is s_{n-1} an unbiased estimate for the standard deviation?

- Unfortunately s_{n-1} is not an unbiased estimate for the standard deviation of the population
- It is better to work with the unbiased variance rather than standard deviation
- There is not a formula for an unbiased estimate for the standard deviation that works for all populations
 - Therefore you will not be asked to find one in your exam

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How do I show the sample mean is an unbiased estimate for the population mean?

- You do not need to learn this proof
 - It is simply here to help with your understanding
- Suppose the population of X has mean μ and variance σ^2
- Take a sample of *n* observations

•
$$E(X_i) = \mu$$

• Using the formula for a linear combination of *n* independent variables:

$$E(\overline{X}) = E\left(\frac{X_1 + X_2 + \dots + X_n}{n}\right)$$
$$= \frac{E(X_1) + E(X_2) + \dots + E(X_n)}{n}$$
$$= \frac{\mu + \mu + \dots + \mu}{n}$$
$$= \frac{n\mu}{n}$$
$$= \mu$$

As E(X) = µ this shows the formula will produce an unbiased estimate for the population mean

RACTICE

Why is there a divisor of n-1 in the unbiased estimate for the variance?

- You do not need to learn this proof
 - It is simply here to help with your understanding
- Suppose the population of X has mean μ and variance σ^2
- Take a sample of *n* observations
 - $\circ X_{1,}X_{2,}...,X_{n}$
 - $E(X_i) = \mu$
 - Var(X_i) = σ^2
- Using the formula for a linear combination of *n* independent variables:

$$\operatorname{Var}(\overline{X}) = \operatorname{Var}\left(\frac{X_1 + X_2 + \dots + X_n}{n}\right)$$
$$= \frac{\operatorname{Var}(X_1) + \operatorname{Var}(X_2) + \dots + \operatorname{Var}(X_n)}{n^2}$$
$$= \frac{\sigma^2 + \sigma^2 + \dots + \sigma^2}{n^2}$$
$$= \frac{n\sigma^2}{n^2}$$
$$= \frac{\sigma^2}{n}$$



- It can be shown that $E(\overline{X}^2) = \mu^2 + \frac{\sigma^2}{n}$
- This comes from rearranging $\operatorname{Var}(\overline{X}) = \operatorname{E}(\overline{X}^2) [\operatorname{E}(\overline{X})]^2$ It can be shown that $\operatorname{E}(X^2) = \operatorname{E}(X_i^2) = \mu^2 + \sigma^2$
 - - This comes from rearranging $Var(X) = E(X^2) [E(X)]^2$
- Using the formula for a linear combination of *n* independent variables:

$$E(S_n^2) = E\left(\frac{\sum X_i^2}{n} - \overline{X}^2\right)$$

= $\frac{\sum E(X_i^2)}{n} - E(\overline{X}^2)$
= $\frac{\sum (\mu^2 + \sigma^2)}{n} - (\mu^2 + \frac{\sigma^2}{n})$
= $\frac{n(\mu^2 + \sigma^2)}{n} - (\mu^2 + \frac{\sigma^2}{n})$
= $\mu^2 + \sigma^2 - (\mu^2 + \frac{\sigma^2}{n})$
= $\sigma^2 - \frac{\sigma^2}{n}$ XAM PAPERS PRACTICE
= $\frac{n\sigma^2 - \sigma^2}{n}$
= $\frac{n-1}{n}\sigma^2$

• As $E(S_n^2) \neq \sigma^2$ this shows that the sample variance is not unbiased

• You need to multiply by
$$\frac{n}{n-1}$$

• E(S²) = σ^2

Exam Tip

- · Check the wording of the exam question carefully to determine which of the following you are given:
 - The population variance:
 - The sample variance:
 - An unbiased estimate for the population variance:







4.7 Binomial Distribution

4.7.1 The Binomial Distribution

Properties of Binomial Distribution

What is a binomial distribution?

- A binomial distribution is a discrete probability distribution
- A discrete random variable X follows a binomial distribution if it counts the number of successes when an experiment satisfies the following conditions:
 - There are a fixed finite number of trials (n)
 - The outcome of each trial is independent of the outcomes of the other trials
 - There are exactly two outcomes of each trial (success or failure)
 - The probability of success is constant (p)
- If X follows a binomial distribution then it is denoted $X \sim B(n, p)$
 - *n* is the **number of trials**
 - p is the probability of success
- The probability of failure is 1 p which is sometimes denoted as q
- The formula for the probability of **r successful trials** is given by:
 - $P(X=r) = {}^{n}C_{r} \times p^{r}(1-p)^{n-r}$ for r = 0, 1, 2, ..., n
 - ${}^{n}C_{r} = \frac{n!}{r!(n-r)!}$ where $n! = n \times (n-1) \times (n-2) \times ... \times 3 \times 2 \times 1$
 - You will be expected to use the distribution function on your GDC to calculate probabilities with the binomial distribution

What are the important properties of a binomial distribution?

• The expected number (mean) of successful trials is

$$E(X) = np$$

- $\circ~$ You are given this in the $formula\,booklet$
- The variance of the number of successful trials is

$$\operatorname{Var}(X) = np(1-p)$$

- You are given this in the formula booklet
- Square root to get the standard deviation
- The distribution can be represented visually using a vertical line graph
 - If p is close to 0 then the graph has a tail to the right
 - If p is close to 1 then the graph has a tail to the left
 - If p is close to 0.5 then the graph is roughly symmetrical
 - If p = 0.5 then the graph is symmetrical







Modelling with Binomial Distribution

How do I set up a binomial model?

- Identify what a trial is in the scenario
 - For example: rolling a dice, flipping a coin, checking hair colour
- Identify what the successful outcome is in the scenario
 - For example: rolling a 6, landing on tails, having black hair
- · Identify the parameters
 - *n* is the number of trials and *p* is the probability of success in each trial
- Make sure you clearly state what your random variable is
 - $\circ\;$ For example, let X be the number of students in a class of 30 with black hair

What can be modelled using a binomial distribution?

- Anything that satisfies the four conditions
- For example: let T be the number of times a fair coin lands on tails when flipped 20 times:
 - A trial is flipping a coin: There are 20 trials so **n = 20**
 - We can assume each coin flip does not affect subsequent coin flips: they are **independent**
 - A success is when the coin lands on tails: **Two outcomes** tails or not tails (heads)
 - The coin is fair: The probability of tails is constant with **p** = 0.5
- Sometimes it might seem like there are more than two outcomes
 - For example: let Y be the number of yellow cars that are in a car park full of 100 cars
 - Although there are more than two possible colours of cars, here the trial is whether a car is yellow so there are two outcomes (yellow or not yellow)
 - Y would still need to fulfil the other conditions in order to follow a binomial distribution
- Sometimes a sample may be taken from a population
 - $\circ\,$ For example: 30% of people in a city have blue eyes, a sample of 30 people from the city is taken and X is the number of them with blue eyes
 - As long as the population is large and the sample is random then it can be assumed that each person has a 30% chance of having blue eyes

What can not be modelled using a binomial distribution?

- Anything where the number of trials is **not fixed** or is **infinite**
 - The number of emails received in an hour
 - The number of times a coin is flipped until it lands on heads
- Anything where the outcome of one trial affects the outcome of the other trials
 - The number of caramels that a person eats when they eat 5 sweets from a bag containing 6 caramels and 4 marshmallows
 - If you eat a caramel for your first sweet then there are less caramels left in the bag when you choose your second sweet
 - Anything where there are more than two outcomes of a trial
 - A person's shoe size
 - The number a dice lands on when rolled
 - Anything where the probability of success changes

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- The number of times that a person can swim a length of a swimming pool in under a minute when swimming 50 lengths
 - The probability of swimming a lap in under a minute will decrease as the person gets tired
 - The probability is not constant



Exam Tip

• An exam question might involve different types of distributions so make it clear which distribution is being used for each variable









4.7.2 Calculating Binomial Probabilities

Calculating Binomial Probabilities

Throughout this section we will use the random variable $X \sim B(n, p)$. For binomial, the probability of X taking a non-integer or negative value is always zero. Therefore any values of Xmentioned in this section will be assumed to be non-negative integers.

How do I calculate P(X = x): the probability of a single value for a binomial distribution?

- You should have a GDC that can calculate binomial probabilities
- · You want to use the "Binomial Probability Distribution" function This is sometimes shortened to BPD, Binomial PD or Binomial Pdf
- You will need to enter:
 - The 'x' value the value of x for which you want to find P(X = x)
 - The 'n' value the number of trials
 - The 'p' value the probability of success
- · Some calculators will give you the option of listing the probabilities for multiple values of x at once
- There is a formula that you can use but you are expected to be able to use the distribution function on your GDC
 - $P(X=x) = {}^{n}C_{x} \times p^{x}(1-p)^{n-x}$ n!

$$nC_x = \frac{n!}{r!(n-r)!}$$

How do I calculate $P(a \le X \le b)$: the cumulative probabilities for a binomial distribution?

- You should have a GDC that can calculate cumulative binomial probabilities
 - Most calculators will find $P(a \le X \le b)$
 - Some calculators can only find $P(X \le b)$
 - The identities below will help in this case
- You should use the "Binomial Cumulative Distribution" function
 - This is sometimes shortened to BCD, Binomial CD or Binomial Cdf
- You will need to enter:
 - The lower value this is the value a
 - This can be zero in the case $P(X \le b)$
 - The upper value this is the value b
 - This can be *n* in the case $P(X \ge a)$
 - The 'n' value the number of trials
 - The 'p' value the probability of success

How do I find probabilities if my GDC only calculates $P(X \le x)$?

- To calculate $P(X \le x)$ just enter x into the cumulative distribution function
- To calculate P(X < x) use:
 - $P(X < x) = P(X \le x 1)$ which works when X is a binomial random variable
 - $P(X < 5) = P(X \le 4)$



- To calculate P(X > x) use:
 - $P(X > x) = 1 P(X \le x)$ which works for any random variable X
 - $P(X > 5) = 1 P(X \le 5)$
- To calculate $P(X \ge x)$ use:
 - $P(X \ge x) = 1 P(X \le x 1)$ which works when X is a binomial random variable
 - $P(X \ge 5) = 1 P(X \le 4)$
- To calculate $P(a \le X \le b)$ use:
 - $P(a \le X \le b) = P(X \le b) P(X \le a 1)$ which works when X is a binomial random variable
 - $P(5 \le X \le 9) = P(X \le 9) P(X \le 4)$

What if an inequality does not have the equals sign (strict inequality)?

- For a binomial distribution (as it is discrete) you could **rewrite all strict inequalities** (< and >) as **weak inequalities** (≤ and ≥) by using the identities for a binomial distribution
 - $P(X < x) = P(X \le x 1)$ and $P(X > x) = P(X \ge x + 1)$
 - For example: $P(X < 5) = P(X \le 4)$ and $P(X > 5) = P(X \ge 6)$
- It helps to think about the range of integers you want
 - Identify the smallest and biggest integers in the range
- If your range has no minimum or maximum then use 0 or n
 - $P(X \le b) = P(0 \le X \le b)$ • $P(X \ge a) = P(a \le X \le n)$ **PAPERS PRACTICE**
- P(a < X ≤ b) = P(a + 1 ≤ X ≤ b)
 P(5 < X ≤ 9) = P(6 ≤ X ≤ 9)
- P(a≤X<b) = P(a≤X≤b-1)
 P(5≤X<9) = P(5≤X≤8)
- $P(a < X < b) = P(a + 1 \le X \le b 1)$ • $P(5 < X < 9) = P(6 \le X \le 8)$

Exam Tip

- If the question is in context then write down the inequality as well as the final answer
 - This means you still might gain a mark even if you accidentally type the wrong numbers into your GDC







4.8 Normal Distribution

4.8.1 The Normal Distribution

Properties of Normal Distribution

The binomial distribution is an example of a discrete probability distribution. The normal distribution is an example of a **continuous** probability distribution.

What is a continuous random variable?

- A continuous random variable (often abbreviated to CRV) is a random variable that can take **any value** within a range of infinite values
 - Continuous random variables usually measure something
 - For example, height, weight, time, etc

What is a continuous probability distribution?

- A continuous probability distribution is a probability distribution in which the random variable X is continuous
- The probability of X being a particular value is always zero
 - P(X=k)=0 for any value k
 - Instead we define the **probability density function** f(x) for a specific value
 - This is a function that describes the relative likelihood that the random variable would be close to that value
 - $\circ~$ We talk about the **probability** of X being within a **certain range**
- A continuous probability distribution can be represented by a continuous graph (the values for X along the horizontal axis and probability **density** on the vertical axis)
- The area under the graph between the points x = a and x = b is equal to P(a ≤ X ≤ b)
 The total area under the graph equals 1
- As P(X = k) = 0 for any value k, it does not matter if we use strict or weak inequalities
 P(X ≤ k) = P(X < k) for any value k when X is a continuous random variable

What is a normal distribution?

- · A normal distribution is a continuous probability distribution
- The continuous random variable X can follow a normal distribution if:
 - The distribution is symmetrical
 - The distribution is **bell-shaped**
- If X follows a normal distribution then it is denoted $X\!\sim\!\mathrm{N}(\mu,\,\sigma^2)$
 - μ is the mean
 - ² is the variance
 - σ is the standard deviation
- If the mean changes then the graph is translated horizontally
- If the variance increases then the graph is widened horizontally and made taller vertically to maintain the same area
 - A small variance leads to a tall curve with a narrow centre
 - A large variance leads to a short curve with a wide centre





What are the important properties of a normal distribution?

- The mean is μ
- The variance is σ²
 - If you need the standard deviation remember to square root this
- · The normal distribution is symmetrical about
 - Mean = Median = Mode = μ
- · There are the results:
 - Approximately two-thirds (68%) of the data lies within one standard deviation of the mean (μ± σ)
 - Approximately 95% of the data lies within two standard deviations of the mean ($\mu \pm 2\sigma$)
 - Nearly all of the data (99.7%) lies within three standard deviations of the mean ($\mu \pm 3\sigma$)



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Modelling with Normal Distribution

What can be modelled using a normal distribution?

- A lot of real-life continuous variables can be modelled by a normal distribution provided that the population is large enough and that the variable is **symmetrical** with **one mode**
- For a normal distribution X can take any real value, however values far from the mean (more than 4 standard deviations away from the mean) have a probability density of **practically zero**
 - This fact allows us to model variables that are not defined for all real values such as height and weight

What can not be modelled using a normal distribution?

- Variables which have more than one mode or no mode
 - For example: the number given by a random number generator
- Variables which are not symmetrical
 - For example: how long a human lives for

Exam Tip

• An exam question might involve different types of distributions so make it clear which distribution is being used for each variable

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4.8.2 Calculations with Norma IDistribution

Calculating Normal Probabilities

Throughout this section we will use the random variable $X \sim N(\mu, \sigma^2)$. For X distributed normally, X can take any real number. Therefore any values mentioned in this section will be assumed to be real numbers.

How do I find probabilities using a normal distribution?

- The area under a normal curve between the points x = a and x = b is equal to the probability P(a < X < b)
 - Remember for a normal distribution you do not need to worry about whether the inequality is strict (< or >) or weak (≤ or ≥)
 - $P(a < X < b) = P(a \le X \le b)$
- You will be **expected to use** distribution functions on your **GDC** to find the probabilities when working with a normal distribution

How do I calculate P(X = x): the probability of a single value for a normal distribution?

- The probability of a single value is always zero for a normal distribution
 You can picture this as the area of a single line is zero
- P(X=x)=0
- Your GDC is likely to have a "Normal Probability Density" function
 - This is sometimes shortened to NPD, Normal PD or Normal Pdf
 - IGNORE THIS FUNCTION for this course!
 - This calculates the probability density function at a point NOT the probability

How do I calculate P(a < X < b): the probability of a range of values for a normal distribution?

- You need a GDC that can calculate cumulative normal probabilities
- You want to use the "Normal Cumulative Distribution" function
 - This is sometimes shortened to NCD, Normal CD or Normal Cdf
- You will need to enter:
 - The 'lower bound' this is the value a
 - The 'upper bound' this is the value b
 - $\circ~$ The ' μ' value this is the mean
 - The 'o' value this is the standard deviation
- Check the order carefully as some calculators ask for standard deviation before mean
 - Remember it is the standard deviation
 - so if you have the variance then square root it
- Always sketch a quick diagram to visualise which area you are looking for

How do I calculate P(X > a) or P(X < b) for a normal distribution?

- You will still use the "Normal Cumulative Distribution" function
- P(X > a) can be estimated using an upper bound that is sufficiently bigger than the mean
 - Using a value that is more than 4 standard deviations **bigger than the mean** is quite accurate
 - Or an easier option is just to input lots of 9's for the upper bound (99999999... or 10⁹⁹)
- P(X < b) can be estimated using a lower bound that is sufficiently smaller than the mean

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- Using a value that is more than 4 standard deviations **smaller than the mean** is quite accurate
- Or an easier option is just to input lots of 9's for the lower bound with a negative sign (-99999999... or -10⁹⁹)

Are there any useful identities?

- $P(X < \mu) = P(X > \mu) = 0.5$
- As P(X = a) = 0 you can use:
 - $\circ P(X < a) + P(X > a) = 1$
 - $\circ P(X > a) = 1 P(X < a)$
 - P(a < X < b) = P(X < b) P(X < a)
- These are useful when:
 - The mean and/or standard deviation are unknown
 - You only have a diagram
 - You are working with the inverse distribution

Exam Tip

Check carefully whether you have entered the standard deviation or variance into your GDC

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Inverse Normal Distribution

Given the value of P(X < a) how do I find the value of a?

- Your GDC will have a function called "Inverse Normal Distribution"
 - Some calculators call this InvN
- Given that P(X < a) = p you will need to enter:
 - The 'area' this is the value p
 - Some calculators might ask for the 'tail' this is the left tail as you know the area to the left of a
 - The 'µ' value this is the mean
 - The 'o' value this is the standard deviation

Given the value of P(X > a) how do I find the value of a?

- If your calculator **does** have the **tail option** (left, right or centre) then you can use the "Inverse Normal Distribution" function straightaway by:
 - Selecting 'right' for the tail
 - Entering the area as 'p'
- If your calculator **does not** have the **tail option** (left, right or centre) then:
 - Given P(X > a) = p
 - Use P(X < a) = 1 P(X > a) to rewrite this as
 - P(X < a) = 1 p

• Then use the method for P(X < a) to find a

Exam Tip

- Always check your answer makes sense
 - If P(X < a) is less than 0.5 then a should be smaller than the mean

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- If P(X < a) is more than 0.5 then a should be bigger than the mean
- A sketch will help you see this







4.9 Further Normal Distribution (incCentral Limit Theorem)

4.9.1 Sample Mean Distribution

Combinations of Normal Variables

What is a linear combination of normal random variables?

- Suppose you have *n* independent normal random variables $X_i \sim N(\mu_i, \sigma_i^2)$ for *i* = 1,2,3, ..., *n*
- A linear combination is of the form $X = a_1X_1 + a_2X_2 + \dots + a_nX_n + b$ where a_i and b are constants
- The mean and variance can be calculated using results from random variables

•
$$E(X) = a_1 \mu_1 + a_2 \mu_2 + \dots + a_n \mu_n + b$$

- $\operatorname{Var}(X) = a_1^2 \sigma_1^2 + a_2^2 \sigma_2^2 + \dots + a_n^2 \sigma_n^2$
 - The variables need to be independent for this result to be true
- A linear combination of n independent normal random variables is also a normal random variable itself
 - $X \sim N(a_1\mu_1 + a_2\mu_2 + \dots + a_n\mu_n + b, a_1^2\sigma_1^2 + a_2^2\sigma_2^2 + \dots + a_n^2\sigma_n^2)$
 - This can be used to find probabilities when combining normal random variables

What is meant by the sample mean distribution?

- Suppose you have a population with distribution X and you take a random sample with n
 observations X₁, X₂, ..., X_n
- The sample mean distribution is the distribution of the values of the sample mean

$$\circ \overline{X} = \frac{X_1 + X_2 + \dots + X_n}{n}$$

- For an individual sample the sample mean \overline{x} can be calculated
 - This is also called a point estimate
 - $\circ \overline{X}$ is the distribution of the point estimates

What does the sample mean distribution look like when X is normally distributed?

If the population is normally distributed then the sample mean distribution is also normally distributed

•
$$E(\overline{X}) = E\left(\frac{X_1 + X_2 + \dots + X_n}{n}\right) = \frac{E(X_1) + E(X_2) + \dots + E(X_n)}{n} = \frac{\mu + \mu + \dots + \mu}{n} = \frac{n\mu}{n} = \mu$$

• $Var(\overline{X}) = Var\left(\frac{X_1 + X_2 + \dots + X_n}{n}\right) = \frac{Var(X_1) + Var(X_2) + \dots + Var(X_n)}{n^2} = \frac{\sigma^2 + \sigma^2 + \dots + \sigma^2}{n^2} = \frac{n\sigma^2}{n^2} = \frac{\sigma^2}{n}$

• Therefore you divide the variance of the population by the size of the sample to get the variance of the sample mean distribution

•
$$X \sim N(\mu, \sigma^2) \Rightarrow \overline{X} \sim N\left(\mu, \frac{\sigma^2}{n}\right)$$



Amber makes a cup of tea using a hot drink vending machine. When the hot water button is pressed the machine dispenses Wml of hot water and when the milk button is pressed the machine dispenses M ml of milk. It is known that $W \sim N(100, 15^2)$ and $M \sim N(10, 2^2)$.

To make a cup of tea Amber presses the hot water button three times and the milk button twice. The total amount of liquid in Amber's cup is modelled by C ml.

a)

2

Write down the distribution of C. $(=W_1+W_2+W_3+M_1+M_2)$ $E(C) = E(W_1) + E(W_2) + E(W_3) + E(M_1) + E(M_2)$ $\mu = 100 + 100 + 100 + 10 + 10 = 320$ $Var(C) = Var(W_1) + Var(W_2) + Var(W_3) + Var(M_1) + Var(M_2)$ $\sigma^2 = 15^2 + 15^2 + 15^2 + 2^2 + 2^2 = 683$ A linear combination of normal variables is also a normal variable (~ N(320, 683)

Find the probability that the total amount of liquid in Amber's cup exceeds 360 ml.

F

Use normal distribution on GDC

$$\mu = 320$$
 $\sigma = \sqrt{683}$
Lower = 360 Upper = 9999...
P(c > 360) = 0.062939...
P(c > 360) = 0.0629 (3sf)

c)

b)

Amber makes 15 cups of tea and calculates the mean \overline{C} . Write down the distribution of \overline{C} .

$$\overline{C} \sim N(\mu, \frac{\sigma^2}{n})$$
$$\overline{C} \sim N(320, \frac{683}{15})$$



Central Limit Theorem

What is the Central Limit Theorem?

- The Central Limit Theorem says that if a sufficiently large random sample is taken from any distribution X then the sample mean distribution \overline{X} can be approximated by a normal distribution
 - In your exam n > 30 will be considered sufficiently large for the sample size

• Therefore
$$\overline{X}$$
 can be modelled by $N\left(\mu, \frac{\sigma^2}{n}\right)$

- μ is the mean of X
- σ^2 is the variance of X
- n is the size of the sample

Do I always need to use the Central Limit Theorem when working with the sample mean distribution?

- No the Central Limit Theorem is not needed when the population is normally distributed
- If X is normally distributed then \overline{X} is normally distributed
 - This is true regardless of the size of the sample
 - The Central Limit Theorem is not needed
- If X is **not normally distributed** then \overline{X} is approximately normally distributed
 - Provided the sample size is large enough
 - The Central Limit Theorem is needed

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The integers 1 to 29 are written on counters and placed in a bag. The expected value when one is picked at random is 15 and the variance is 70. Susie randomly picks 40 integers, returning the counter after each selection.

a)

Find the probability that the mean of Susie's 40 numbers is less than 13.



Explain whether it was necessary to use the Central Limit Theorem in your calculation.





4.9.2 Confidence Interval for the Mean

Confidence Interval for μ

What is a confidence interval?

- It is impossible to find the exact value of the population mean when taking a sample
 - The mean of a sample is called a **point estimate**
 - The best we can do is find an **interval** for which the **exact value is likely to lie**
 - This is called the confidence interval for the mean
- The **confidence level** of a confidence interval is the **probability** that the **interval contains the population mean**
 - Be careful with the wording the population mean is a fixed value so it does not make sense to talk about the probability that it lies within an interval
 - Instead we talk about the probability of an interval containing the mean
 - Suppose samples were collected and a 95% confidence interval for the population mean was constructed for each sample then for every 100 intervals we would expect on average 95 of them to contain the mean
 - 95 out of 100 is not guaranteed it is possible that all of them could contain the mean
 - It is also possible (though very unlikely) that none of them contains the mean

How do I find a confidence interval for the population mean (μ) ?

- You will be given data using a sample from a population
 - The population will be **normally distributed**
 - If not then the sample size should be large enough so you can use the Central Limit Theorem
- You will use the interval functions on your calculator
- Use a z-interval if the population variance is known σ^2
 - On your GDC enter:
 - the standard deviation σ and the confidence level a%
 - EITHER the raw data
 - OR the sample mean \overline{x} and the sample size n
- Use a t-interval if the population variance is unknown
 - In this case the test uses the **unbiased estimate** for the variance s_{n-1}^2
 - On your GDC enter:
 - the confidence level α %
 - EITHER the raw data
 - OR the sample mean \overline{x} , the value of s_{n-1} and the sample size n
- Your GDC will give you the lower and upper bounds of the interval
 - It can be written as $a < \mu < b$

What affects the width of a confidence interval?

- The width of a confidence interval is the range of the values in the interval
- The confidence level affects the width



- Increasing the confidence level will increase the width
- Decreasing the confidence level with decrease the width
- The size of the sample affects the width
 - Increasing the sample size will decrease the width
 - Decreasing the sample size will increase the width

How can l interpret a confidence interval?

- After you have found a confidence interval for μ you might be expected to **comment on the claim** for a value of μ
- If the claimed value is within the confidence interval then there is not enough evidence to reject the claim
 - Therefore the claim is supported
- If the claimed value is **outside** the interval then there is **sufficient evidence to reject the claim**
 - The value is unlikely to be correct





Cara wants to check the mean weight of burgers sold by a butcher. The weights of the burgers are assumed to be normally distributed. Cara takes a random sample of 12 burgers and finds that the mean weight is 293 grams and the standard deviation of the sample is 5.5 grams.

a)

Find a 95% confidence interval for the population mean, giving your answer to 4 significant figures.

The population variance is unknown so use a t-interval Unbiased estimate of population variance s_{u-1}^2 Formula booklet $s_{n-1}^2 = \frac{n}{n-1}s_n^2$ $S_{n-1}^{2} = \frac{12}{11} \times 5.5^{2} = 33$ Enter data into GDC Confidence level = 0.95 $\bar{x} = 293$ $s_{n_1} = 12$ Lower: 289.35. Upper: 296.64 ... 2894 Apr 296 A R43FRS PRACTICE

b)

The butcher claims the burgers weigh 300 grams. Comment on this claim with reference to the confidence interval.





4.10 Poisson Distribution

4.10.1 Poisson Distribution

Properties of Poisson Distribution

What is a Poisson distribution?

- A Poisson distribution is a discrete probability distribution
- A discrete random variable X follows a Poisson distribution if it counts the number of occurrences in a fixed time period given the following conditions:
 - Occurrences are independent
 - Occurrences occur at a uniform average rate for the time period (m)
- If X follows a Poisson distribution then it is denoted $X \sim Po(m)$
 - m is the average rate of occurrences for the time period
- The formula for the probability of roccurrences is given by:

•
$$P(X=r) = \frac{e^{-m}m^r}{r!}$$
 for $r = 0, 1, 2, ...$

- $r! = r \times (r-1) \times ... \times 2 \times 1$ and 0! = 1
- There is no upper bound for the number of occurrences
- You will be expected to use the distribution function on your GDC to calculate probabilities with the Poisson distribution

What are the important properties of a Poisson distribution?

- The expected number (mean) of occurrences is m
 - You are given this in the formula booklet
- The **variance** of the number of occurrences is *m*
 - You are given this in the **formula booklet**
 - Square root to get the standard deviation
- The mean and variance for a Poisson distribution are equal
- The distribution can be represented visually using a vertical line graph
 - The graphs have tails to the right for all values of m
 - As m gets larger the graph gets more symmetrical
- If $X \sim Po(m)$ and $Y \sim Po(\lambda)$ are independent then $X + Y \sim Po(m + \lambda)$
 - This extends to *n* independent Poisson distributions $X_i \sim Po(m_i)$
 - $X_1 + X_2 + \dots + X_n \sim Po(m_1 + m_2 + \dots + m_n)$







Modelling with Poisson Distribution

How do I set up a Poisson model?

- Identify what an occurrence is in the scenario
 - For example: a car passing a camera, a machine producing a faulty item
- Use proportion to find the mean number of occurrences for the given time period
 - For example: 10 cars in 5 minutes would be 120 cars in an hour if the Poisson model works for both time periods
- Make sure you clearly state what your random variable is
 - For example: let X be the number of cars passing a camera in 10 minutes

What can be modelled using a Poisson distribution?

- Anything that satisfies the two conditions
- For example, Let C be the number of calls that a helpline receives within a 15-minute period:
 C~Po(m)
 - An occurrence is the helpline receiving a call and can be considered independent
 - The helpline receives calls at an average rate of *m* calls during a 15-minute period
- Sometimes a measure of space will be used instead of a time period
 - For example, the number of daisies that exist on a square metre of grass
- If the mean and variance of a discrete variable are equal then it might be possible to use a Poisson model



Exam Tip EXAM PAPERS PRACTICE

• An exam question might involve different types of distributions so make it clear which distribution is being used for each variable







4.10.2 Calculating Poisson Probabilities

Calculating Poisson Probabilities

Throughout this section we will use the random variable $X \sim Po(m)$. For a Poisson distribution X, the probability of X taking a non-integer or negative value is always zero. Therefore, any values mentioned in this section for X will be assumed to be non-negative integers. The value of m can be any real positive value.

How do I calculate P(X = x): the probability of a single value for a Poisson distribution?

- You should have a GDC that can calculate Poisson probabilities
- You want to use the "Poisson Probability Distribution" function
 - This is sometimes shortened to PPD, Poisson PD or Poisson Pdf
- You will need to enter:
 - The 'x' value the value of x for which you want to find P(X = x)
 - The '\lambda' value the mean number of occurrences (m)
- Some calculators will give you the option of **listing the probabilities** for **multiple values** of x at once
- There is a formula that you can use but you are expected to be able to use the distribution function on your GDC

$$\circ P(X=x) = \frac{e^{-m_{II}}}{x!}$$

- where e is Euler's constant
- x!=x×(x-1)×...×2×1and0!=1 PRACTICE

How do I calculate $P(a \le X \le b)$: the cumulative probabilities for a Poisson distribution?

- You should have a GDC that can calculate cumulative Poisson probabilities
 - Most calculators will find $P(a \le X \le b)$
 - Some calculators can only find $P(X \le b)$
 - The identities below will help in this case
- You should use the "Poisson Cumulative Distribution" function
 - This is sometimes shortened to PCD, Poisson CD or Poisson Cdf
- You will need to enter:
 - The lower value this is the **value a**
 - This can be zero in the case $P(X \le b)$
 - The upper value this is the **value b**
 - This can be a very large number (9999...) in the case $P(X \ge a)$
 - \circ The ' λ ' value the mean number of occurrences (m)

How do I find probabilities if my GDC only calculates $P(X \le x)$?

- To calculate $P(X \le x)$ just enter x into the cumulative distribution function
- To calculate P(X < x) use:
 - $P(X < x) = P(X \le x 1)$ which works when X is a Poisson random variable

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- $P(X < 5) = P(X \le 4)$
- To calculate P(X > x) use:
 - $P(X > x) = 1 P(X \le x)$ which works for any random variable X
 - $P(X > 5) = 1 P(X \le 5)$
- To calculate $P(X \ge x)$ use:
 - $P(X \ge x) = 1 P(X \le x 1)$ which works when X is a Poisson random variable
 - $P(X \ge 5) = 1 P(X \le 4)$
- To calculate $P(a \le X \le b)$ use:
 - $P(a \le X \le b) = P(X \le b) P(X \le a 1)$ which works when X is a Poisson random variable
 - $P(5 \le X \le 9) = P(X \le 9) P(X \le 4)$

What if an inequality does not have the equals sign (strict inequality)?

- For a Poisson distribution (as it is discrete) you could **rewrite all strict inequalities** (< and >) as **weak inequalities** (≤ and ≥) by using the identities for a Poisson distribution
 - $P(X < x) = P(X \le x 1)$ and $P(X > x) = P(X \ge x + 1)$
 - For example: P(X < 5) = P(X ≤ 4) and P(X > 5) = P(X ≥ 6)
- It helps to think about the range of integers you want
 Identify the smallest and biggest integers in the range
- If your range has no minimum then use 0
 - $P(X \le b) = P(0 \le X \le b)$ PAPERS PRACTICE
- $P(a < X \le b) = P(a + 1 \le X \le b)$ • $P(5 < X \le 9) = P(6 \le X \le 9)$
- P(a≤X<b) = P(a≤X≤b-1)
 P(5≤X<9) = P(5≤X≤8)
- $P(a < X < b) = P(a + 1 \le X \le b 1)$ • $P(5 < X < 9) = P(6 \le X \le 8)$







4.11 Hypothesis Testing

4.11.1 Hypothesis Testing

Language of Hypothesis Testing

What is a hypothesis test?

- A hypothesis test uses a **sample of data** in an experiment to test a **statement** made about the **population**
 - The statement is either about a **population parameter** or the distribution of the **population**
- The hypothesis test will look at the probability of observed outcomes happening under set conditions
- The probability found will be compared against a given **significance level** to determine whether there is **evidence to support** the statement being made

What are the key terms used in statistical hypothesis testing?

- Every hypothesis test **must** begin with a clear **null hypothesis** (what we believe to already be true) and **alternative hypothesis** (how we believe the data pattern or probability distribution might have changed)
- A hypothesis is an assumption that is made about a particular population parameter or the distribution of the population
 - A population parameter is a numerical characteristic which helps define a population
 Such as the mean value of the population
 - $\circ~$ The **null hypothesis** is denoted $H^{}_0$ and sets out the assumed population parameter or distribution given that no change has happened
 - $\circ~$ The $\mbox{alternative hypothesis}$ is denoted $H^{}_1$ and sets out how we think the population parameter or distribution could have changed
 - When a hypothesis test is carried out, the null hypothesis is **assumed to be true** and this assumption will either be **accepted** or **rejected**
 - When a null hypothesis is accepted or rejected a **statistical inference** is made
- A hypothesis test will always be carried out at an appropriate significance level
 - The significance level sets the **smallest probability** that an event could have occurred by chance
 - Any probability smaller than the significance level would suggest that the event is unlikely to have happened by chance
 - The significance level must be set before the hypothesis test is carried out
 - The **significance level** will usually be 1%, 5% or 10%, however it may vary



One-tailed Tests

What are one-tailed tests?

- A one-tailed test is used for testing:
 - Whether a distribution can be used to model the population
 - Whether the population parameter has increased
 - Whether the population parameter has decreased
- One-tailed tests can be used with:
 - Chi-squared test for independence
 - Chi-squared goodness of fit test
 - Test for proportion of a binomial distribution
 - Test for population mean of a Poisson distribution
 - Test for population mean of a normal distribution
 - Test to compare population means of two distributions

Two-tailed Tests

What are two-tailed tests?

- A two-tailed test is used for testing:
 - Whether the population parameter has changed
- Two-tailed tests can be used with:
 - Test for population mean of a normal distribution
 - Test to compare population means of two distributions



Conclusions of Hypothesis Testing

How do I decide whether to reject or accept the null hypothesis?

- A sample of the population is taken and the **test statistic** is calculated **using the observations** from the sample
 - Your GDC can calculate the test statistic for you (if required)
- To decide whether or not to reject the null hypothesis you first need either the *p-value* or the critical region
- The **p** value is the probability of a value being at least as extreme as the test statistic, assuming that the null hypothesis is true
 - Your GDC will give you the p-value (if required)
 - If the *p*-value is less than the significance level then the null hypothesis would be rejected
- The critical region is the range of values of the test statistic which will lead to the null hypothesis being rejected
 - If the **test statistic** falls within the **critical region** then the **null hypothesis** would be **rejected**
- The critical value is the boundary of the critical region
 - It is the least extreme value that would lead to the rejection of the null hypothesis
 - The critical value is determined by the significance level

How should a conclusion be written for a hypothesis test?

- Your conclusion **must** be written in the **context** of the question
- Use the wording in the question to help you write your conclusion C
 - If **rejecting the null** hypothesis your conclusion should state that there is **sufficient evidence** to suggest that the null hypothesis is unlikely to be true
 - If **accepting the null** hypothesis your conclusion should state that there is **not enough evidence** to suggest that the null hypothesis is unlikely to be true
- Your conclusion **must not** be definitive
 - There is a chance that the test has led to an **incorrect conclusion**
 - The outcome is **dependent on the sample**
 - a different sample might lead to a different outcome
- The conclusion of a two-tailed test can state if there is evidence of a change
 - You should not state whether this change is an increase or decrease
 - If you are testing the difference between the means of two populations then you can **only conclude that the means are not equal**
 - You can not say which population mean is bigger
 - You'd need to use a **one-tailed** test for this

Exam Tip

- Accepting the null hypothesis does **not** mean that you are saying it is true
 - You are simply saying there is not enough evidence to reject it



4.11.2 Chi-squared Test for Independence

Chi-Squared Test for Independence

What is a chi-squared test for independence?

- A chi-squared (χ²) test for independence is a hypothesis test used to test whether two variables are independent of each other
 - This is sometimes called a χ^2 two-way test
- This is an example of a goodness of fit test
 - We are testing whether the data fits the model that the variables are independent
- The chi-squared (χ^2) distribution is used for this test
- You will use a contingency table
 - This is a two-way table that shows the **observed frequencies** for the different combinations of the two variables
 - For example: if the two variables are hair colour and eye colour then the contingency table will show the frequencies of the different combinations

Why might I have to combine rows or columns?

- The observed values are used to calculate expected values
 - These are the expected frequencies for each combination assuming that the variables are independent
 - Your GDC can calculate these for you after you input the observed frequencies
- The expected values must all be bigger than 5
- If one of the expected values is less than 5 then you will have to combine the corresponding row or column in the matrix of observed values with the adjacent row or column
 - The decision between row or column will be based on which seems the **most** appropriate
 - For example: if the two variables are age and favourite TV genre then it is more appropriate to combine age groups than types of genre

What are the degrees of freedom?

- There will be a **minimum number of expected values** you would need to know in order to be able to calculate all the expected values
- + This minimum number is called the degrees of freedom and is often denoted by u
- For a test for independence with an $m \times n$ contingency table
 - $\circ v = (m-1) \times (n-1)$
 - For example: If there are 5 rows and 3 columns then you only need to know **2 of the** values in **4 of the rows** as the rest can be calculated using the totals

What are the steps for a chi-squared test for independence?

- STEP 1: Write the hypotheses
 - \circ H₀: Variable X is independent of variable Y
 - H1: Variable X is not independent of variable Y
 - Make sure you clearly write what the variables are and don't just call them X and Y
- STEP 2: Calculate the degrees of freedom for the test
 - For an *m* × *n* contingency table
 - Degrees of freedom is $v = (m-1) \times (n-1)$

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- STEP 3: Enteryour observed frequencies into your GDC using the option for a 2-way test
 - Enter these as a matrix
 - Your GDC will give you a matrix of the expected values (assuming the variables are independent)
 - If any values are 5 or less then combine rows/columns and repeat step 2
 - Your GDC will also give you the x² statistic and its p-value
 - The χ^2 statistic is denoted as χ^2_{calc}
- STEP 4: Decide whether there is evidence to reject the null hypothesis
 - EITHER compare the 2² statistic with the given critical value
 - If χ^2 statistic > critical value then reject H₀
 - If x² statistic < critical value then accept H₀
 - OR compare the *p*-value with the given significance level
 - If p-value < significance level then reject H₀
 - If p-value > significance level then accept H₀
- STEP 5: Write your conclusion
 - If you reject H₀
 - There is sufficient evidence to suggest that variable X is not independent of variable Y
 - Therefore this suggests they are associated
 - If you accept H₀
 - There is insufficient evidence to suggest that variable X is not independent of variable Y A PAPERS PRACTICE
 Therefore this suggests they are independent

How do I calculate the chi-squared statistic?

- You are expected to be able to use your GDC to calculate the χ^2 statistic by inputting the matrix of the observed frequencies
- · Seeing how it is done by hand might deepen your understanding but you are not expected to use this method
- STEP 1: For each observed frequency O_i calculate its expected frequency E_i
 - Assuming the variables are independent
 - $E_i = P(X = x) \times P(Y = y) \times Total$
 - Which simplifies to $E_i = \frac{\text{Row Total} \times \text{Column Total}}{\text{Overall Total}}$

• STEP 2: Calculate the χ^2 statistic using the formula

$$\circ \chi^2_{calc} = \sum \frac{(O_i - E_i)^2}{E_i}$$

- · You do not need to learn this formula as your GDC calculates it for you
- To calculate the p-value you would find the probability of a value being bigger than your χ^2 statistic using a χ^2 distribution with v degrees of freedom





Exam Tip

Note for Internal Assessments (IA)

- If you use a χ^2 test in your IA then beware that the outcome may not be accurate if there is only 1 degree of freedom
 - This means it is a 2 x 2 contingency table





At a school in Paris, it is believed that favourite film genre is related to favourite subject. 500 students were asked to indicate their favourite film genre and favourite subject from a selection and the results are indicated in the table below.

	Comedy	Action	Romance	Thriller
Maths	51	52	37	55
Sports	59	63	41	33
Geography	35	31	28	15

It is decided to test this hypothesis by using a χ^2 test for independence at the 1% significance level.

The critical value is 16.812.

a)

State the null and alternative hypotheses for this test.

Ho: Favourite	subject is	independent of	tavourite film genre
H. Favourite	subject is	not independent	of favourite film genre

b) Write down the number of degrees of freedom for this table.

```
y = (rows - 1) \times (columns - 1) = (3 - 1) \times (4 - 1)
```

y=6

c)

Calculate the χ^2 test statistic for this data.

```
Type matrix into GDC
\chi^2 statistic = 12.817...
\chi^2_{calc} = 12.8 (3 sf)
```

d)

Write down the conclusion to the test. Give a reason for your answer.



12.8 < 16.812

Accept H_0 as χ^2 statistic < critical value. There is insufficient evidence to suggest that favourite subject is not independent of favourite film genre. Therefore this suggests they are independent.





4.11.3 Goodness of Fit Test

Chi-Squared GOF: Uniform

What is a chi-squared goodness of fit test for a given distribution?

- A chi-squared (χ^2) goodness of fit test is used to test data from a sample which suggests that the population has a given distribution
- This could be that:
 - the proportions of the population for different categories follows a given ratio
 - the population follows a **uniform distribution**
 - This means all outcomes are equally likely

What are the steps for a chi-squared goodness of fit test for a given distribution?

- STEP 1: Write the hypotheses
 - \circ H₀: Variable X can be modelled by the given distribution
 - \circ H₁: Variable X cannot be modelled by the given distribution
 - Make sure you clearly write what the variable is and don't just call it X
- STEP 2: Calculate the degrees of freedom for the test
 - Forkoutcomes
 - Degrees of freedom is v = k 1
- STEP 3: Calculate the expected frequencies
 - Split the total frequency using the given ratio
 - For a uniform distribution: divide the total frequency N by the number of outcomes k
- STEP 4: Enter the frequencies and the degrees of freedom into your GDC
 - Enter the observed and expected frequencies as two separate lists
 - Your GDC will then give you the χ^2 statistic and its p-value
 - The χ^2 statistic is denoted as χ^2_{calc}
- STEP 5: Decide whether there is evidence to reject the null hypothesis
 - EITHER compare the χ^2 statistic with the given critical value
 - If χ^2 statistic > critical value then reject H₀
 - If X² statistic < critical value then accept H₀
 - OR compare the *p-value* with the given significance level
 - If p-value < significance level then reject H₀
 - If p-value > significance level then accept H₀
- STEP 6: Write your conclusion
 - If you reject H₀
 - There is sufficient evidence to suggest that variable X does not follow the given distribution
 - Therefore this suggests that the data is **not distributed as claimed**
 - If you accept H₀
 - There is insufficient evidence to suggest that variable X does not follow the given distribution
 - Therefore this suggests that the data is **distributed as claimed**

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A car salesman is interested in how his sales are distributed and records his sales results over a period of six weeks. The data is shown in the table.

Week	1	2	3	4	5	6
Number of sales	15	17	11	21	14	12

A χ^2 goodness of fit test is to be performed on the data at the 5% significance level to find out whether the data fits a uniform distribution.

a)

Find the expected frequency of sales for each week if the data were uniformly distributed.





d) Calculate the p-value.



e)

State the conclusion of the test. Give a reason for your answer.

0.493 > 0.05

Accept Ho as p-value > significance level There is insufficient evidence to suggest that number of sales can not be modelled by a uniform distribution. Therefore this suggests it is uniformly distributed

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Chi-Squared GOF: Binomial

What is a chi-squared goodness of fit test for a binomial distribution?

- A chi-squared (χ^2) **goodness of fit test** is used to test data from a sample suggesting that the population has a **binomial distribution**
 - You will either be **given a precise binomial distribution** to test **B**(*n*, *p*) with an assumed value for *p*
 - Or you will be asked to test whether a binomial distribution is **suitable without being given an assumed value** for p
 - In this case you will have to calculate an estimate for the value of p for the binomial distribution
 - To calculate it divide the mean by the value of n

•
$$p = \frac{\overline{x}}{n} = \frac{1}{n} \times \frac{\sum fx}{\sum f}$$

What are the steps for a chi-squared goodness of fit test for a binomial distribution?

- STEP 1: Write the hypotheses
 - H₀: Variable X can be modelled by a binomial distribution
 - H₁: Variable X cannot be modelled by a binomial distribution
 - Make sure you clearly write what the variable is and don't just call it X
 - If you are given the assumed value of p then state the precise distribution B(n, p)
- STEP 2: Calculate the expected frequencies
 - If you were not given the assumed value of p then you will first have to estimate it using the observed data M DADEDS DDACTICE
 - Find the probability of the outcome using the binomial distribution P(X = x)
 - Multiply the probability by the total frequency $P(X = x) \times N$
 - You will have to combine rows/columns if any expected values are 5 or less
- STEP 3: Calculate the degrees of freedom for the test
 - For k outcomes (after combining expected values if needed)
 - Degrees of freedom is
 - v = k 1 if you were **given** the assumed value of p
 - v = k 2 if you had to estimate the value of p
- STEP 4: Enter the frequencies and the degrees of freedom into your GDC
 - Enter the observed and expected frequencies as two separate lists
 - $\circ~$ Your GDC will then give you the χ^2 statistic and its p-value
 - The χ^2 statistic is denoted as χ^2_{calc}
- STEP 5: Decide whether there is evidence to reject the null hypothesis
 - EITHER compare the χ^2 statistic with the given critical value
 - If χ^2 statistic > critical value then reject H₀
 - If χ^2 statistic < critical value then accept H₀
 - OR compare the p-value with the given significance level
 - If p-value < significance level then reject H₀
 - If p-value > significance level then accept H₀
- STEP 6: Write your conclusion



- If you reject H₀
 - There is sufficient evidence to suggest that variable X does not follow the binomial distribution B(n, p)
 - Therefore this suggests that the data **does not follow** B(n, p)
- If you accept H₀
 - There is insufficient evidence to suggest that variable X does not follow the binomial distribution B(n, p)
 - Therefore this suggests that the data follows B(n, p)





A stage in a video game has three boss battles. 1000 people try this stage of the video game and the number of bosses defeated by each player is recorded.

Number of bosses defeated	0	1	2	3
Frequency	490	384	111	15

A χ^2 goodness of fit test at the 5% significance level is used to decide whether the number of bosses defeated can be modelled by a binomial distribution with a 20% probability of success.

a)

State the null and alternative hypotheses.



b)

Assuming the binomial distribution holds, find the expected number of people that would defeat exactly one boss.

```
Let X \sim B(3, 0.2)
Using GD( P(X = 1) = 0.384
Expected 1000 x 0.384 = 384
Expected frequency of 1 = 384
```

c)

Calculate the p-value for the test.







Chi-Squared GOF: Normal

What is a chi-squared goodness of fit test for a normal distribution?

- A chi-squared (χ^2) goodness of fit test is used to test data from a sample suggesting that the population has a normal distribution
 - You will either be given a precise normal distribution to test $N(\mu, \sigma^2)$ with assumed values for μ and σ
 - $\circ~$ Or you will be asked to test whether a normal distribution is **suitable without being** given assumed values for μ and/or σ
 - In this case you will have to calculate an estimate for the value of μ and/or σ for the normal distribution
 - Either use your GDC or use the formulae

•
$$\overline{x} = \frac{\sum fx}{\sum f}$$
 and $s_{n-1}^2 = \frac{n}{n-1}s_n^2$

What are the steps for a chi-squared goodness of fit test for a normal distribution?

- STEP 1: Write the hypotheses
 - H₀: Variable X can be modelled by a normal distribution
 - H₁: Variable X cannot be modelled by a normal distribution
 - Make sure you clearly write what the variable is and don't just call it X
 - If you are given the assumed values of μ and σ then state the precise distribution $N(\mu, \sigma^2)$
- STEP 2: Calculate the expected frequencies
 - If you were not given the assumed values of μ or σ then you will first have to estimate them
 - Find the probability of the outcome using the normal distribution P(a < X < b)
 - Multiply the probability by the total frequency $P(a < X < b) \times N$
 - You will have to combine rows/columns if any expected values are 5 or less
- STEP 3: Calculate the degrees of freedom for the test
 - For k class intervals (after combining expected values if needed)
 - Degrees of freedom is
 - v = k 1 if you were **given** the assumed values for **both** μ and σ
 - v = k 2 if you had to estimate either μ or σ but not both
 - v = k 3 if you had to estimate both μ and σ
- STEP 4: Enter the frequencies and the degrees of freedom into your GDC
 - Enter the observed and expected frequencies as two separate lists
 - Your GDC will then give you the χ^2 statistic and its p-value
 - The χ^2 statistic is denoted as χ^2_{calc}
- STEP 5: Decide whether there is evidence to reject the null hypothesis
 - EITHER compare the χ^2 statistic with the given critical value
 - If \chi² statistic > critical value then reject H₀
 - If χ^2 statistic < critical value then **accept H**₀
 - OR compare the *p*-value with the given significance level

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- If p-value < significance level then reject H₀
- If p-value > significance level then accept H₀
- STEP 6: Write your conclusion
 - If you reject H₀
 - There is sufficient evidence to suggest that variable X does not follow the normal distribution $N(\mu, \sigma^2)$
 - Therefore this suggests that the data does not follow $N(\mu, \sigma^2)$
 - If you accept H₀
 - There is insufficient evidence to suggest that variable X does not follow the normal distribution $N(\mu, \sigma^2)$
 - Therefore this suggests that the data follows $N(\mu, \sigma^2)$





300 marbled ducks in Quacktown are weighed and the results are shown in the table below.

Mass (g)	Frequency
<i>m</i> < 450	1
$450 \le m < 470$	9
$470 \le m < 520$	158
$520 \le m < 570$	123
<i>m</i> ≥570	9

A χ^2 goodness of fit test at the 10% significance level is used to decide whether the mass of a marbled duck can be modelled by a normal distribution with mean 520 g and standard deviation 30 g.

a)

Explain why it is necessary to combine the groups m < 450 and $450 \le m < 470$ to create the group m < 470 with frequency 10.

Combine categories if expected frequencies are 5 or less 300 × P(X < 450 | X~N(520,30²)) = 300 × 0.00981...=2.944... The expected frequency is less than 5 so combine with the next category.

b)

Calculate the expected frequencies, giving your answers correct to 2 decimal places.

Let X ~ N(520, 30²) 300 × probability

Mass (9)	Probability	Expected frequency
m < 470	0.047790	14.34
470×m×520	0.452209	135.66
520 ≤ m < 570	0.452209	135.66
m≥ 570	0.047790	14.34



C) Write down the null and alternative hypotheses.



d)

Calculate the χ^2 statistic. Enter the observed and expected frequencies into GD(y = 4 - 1 = 3) λ^2 statistic = 8.162 ... $\chi^{2}_{colc} = 8.16$ (3sf) e) Given that the critical value is 6.251, state the conclusion of the test. Give a reason for your answer. EMANAPAPERS PRACTICE Reject Ho as X° statistic > critical value. There is sufficient evidence to suggest that the mass of the marbled ducks can not be modelled by the normal distribution N(520, 302).


Chi-squared GOF: Poisson

What is a chi-squared goodness of fit test for a Poisson distribution?

- A chi-squared (χ^2) goodness of fit test is used to test data from a sample suggesting that the population has a Poisson distribution
 - You will either be **given a precise Poisson distribution** to test **Po**(*m*) with an assumed value for *m*
 - Or you will be asked to test whether a Poisson distribution is **suitable without being given an assumed value** for *m*
 - In this case you will have to calculate an estimate for the value of m for the Poisson distribution
 - To calculate it just calculate the mean

$$m = \frac{\sum fx}{\sum f}$$

What are the steps for a chi-squared goodness of fit test for a Poisson distribution?

- STEP 1: Write the hypotheses
 - H₀: Variable X can be modelled by a Poisson distribution
 - \circ H₁: Variable X cannot be modelled by a Poisson distribution
 - Make sure you clearly write what the variable is and don't just call it X
 - If you are given the assumed value of m then state the precise distribution Po(m)
- STEP 2: Calculate the expected frequencies
 - If you were not given the assumed value of *m* then you will first have to **estimate it** using the **observed data APERS PRACTICE**
 - Find the probability of the outcome using the Poisson distribution P(X = x)
 - Multiply the probability by the total frequency $P(X = x) \times N$
 - If a is the smallest observed value then calculate $P(X \le a)$
 - If b is the largest observed value then calculate $P(X \ge b)$
 - You will have to combine rows/columns if any expected values are 5 or less
- STEP 3: Calculate the degrees of freedom for the test
 - For k outcomes (after combining expected values if needed)
 - Degree of freedom is
 - v = k 1 if you were **given** the assumed value of m
 - v = k 2 if you had to estimate the value of m
- STEP 4: Enter the frequencies and the degree of freedom into your GDC
 - Enter the observed and expected frequencies as two separate lists
 - $\circ~$ Your GDC will then give you the χ^2 statistic and its p-value
 - The χ^2 statistic is denoted as χ^2_{calc}
- STEP 5: Decide whether there is evidence to reject the null hypothesis
 - EITHER compare the 2² statistic with the given critical value
 - If \(\chi^2\) statistic > critical value then reject H_0
 - If \(\chi^2\) statistic < critical value then accept H₀
 - OR compare the *p*-value with the given significance level
 - If p-value < significance level then reject H₀

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- If p-value > significance level then accept H₀
- STEP 6: Write your conclusion
 - If you reject H₀
 - There is sufficient evidence to suggest that variable X does not follow the Poisson distribution Po(m)
 - Therefore this suggests that the data **does not follow** Po(m)
 - If you accept Ho
 - There is insufficient evidence to suggest that variable X does not follow the Poisson distribution Po(m)
 - Therefore this suggests that the data follows Po(m)





A parent claims the number of messages they receive from their teenage child within an hour can be modelled by a Poisson distribution. The parent collects data from 100 one hour periods and records the observed frequencies of the messages received from the child. The parent calculates the mean number of messages received from the sample and uses this to calculate the expected frequencies if a Poisson model is used.

Number of messages	Observed frequency	Expected frequency
0	9	7.28
1	16	а
2	23	24.99
3	22	21.82
4	16	14.29
5	14	7.49
6 or more	0	b

A χ^2 goodness of fit test at the 10% significance level is used to test the parent's claim. a)

Write down null and alternative hypotheses to test the parent's claim.

We are not given a specific Poisson distribution

- Ho: Number of messages received con be modelled by a Poisson distribution
- H,: Number of messages received con not be modelled by a Poisson distribution

b)

Show that the mean number of messages received per hour for the sample is 2.62.

$$m = \frac{\Sigma f_{x}}{\Sigma f} = \frac{0 \times 9 + 1 \times 16 + 2 \times 23 + 3 \times 22 + 4 \times 16 + 5 \times 14}{9 + 16 + 23 + 22 + 16 + 14} = 2.62$$







4.12 Further Hypothesis Testing

4.12.1 Hypothesis Testing for Mean (One Sample)

One-Sample z-tests

What is a one-sample z-test?

- A one-sample z-test is used to test the mean (μ) of a normally distributed population
 You use a z-test when the population variance (σ²) is known
- The **mean of a sample** of size *n* is calculated \overline{x} and a normal distribution is used to test the **test statistic**
- \overline{x} can be used as the test statistic
 - In this case you would use the distribution $\overline{X} \sim N\left(\mu, \frac{\sigma^2}{n}\right)$
 - Remember when using this distribution that the standard deviation is $\frac{\sigma}{\sqrt{n}}$
- $z = \frac{\overline{x} \mu}{\frac{\sigma}{\overline{x}}}$ can be used as the test statistic
 - In this case you would use the distribution $Z \sim N(0, 1^2)$
 - This is a more old-fashioned approach but your GDC still might tell you the z-value when you do the test
 - You will not need to use this method in the exam as your GDC should be capable of doing the other method

What are the steps for performing a one-sample z-test on my GDC?

STEP 1: Write the hypotheses

- $\circ H_0: \mu = \mu_0$
 - Clearly state that μ represents the **population mean**
 - μ₀ is the assumed population mean
- For a **one-tailed** test $H_1: \mu < \mu_0$ or $H_1: \mu > \mu_0$
- For a **two-tailed** test: $H_1: \mu \neq \mu_0$
 - The alternative hypothesis will depend on what is being tested
- STEP 2: Enter the data into your GDC and choose the one-sample z-test
 - If you have the raw data
 - Enter the data as a list
 - Enter the value of σ
 - If you have summary statistics
 - Enter the values of \overline{X} , σ and n
 - Your GDC will give you the p-value
- STEP 3: Decide whether there is evidence to reject the null hypothesis
 - \circ If the p-value < significance level then reject H₀
- STEP 4: Write your conclusion



- If you reject H₀ then there is evidence to suggest that...
 - The mean has decreased (for $H_1: \mu < \mu_0$)
 - The mean has increased (for $H_1: \mu > \mu_0$)
 - The mean has changed (for $H_1: \mu \neq \mu_0$)
- If you **accept H₀** then there is **insufficient evidence** to reject the null hypothesis which suggests that...
 - The mean has not decreased (for $H_1: \mu < \mu_0$)
 - The mean has not increased (for $H_1: \mu > \mu_0$)
 - The mean has not changed (for $H_1: \mu \neq \mu_0$)

How do I find the p-value for a one-sample z-test using a normal distribution?

- The p-value is determined by the test statistic \overline{x}
- For H₁: $\mu < \mu_0$ the p-value is $P(\overline{X} < \overline{x} | \mu = \mu_0)$
- For H₁: $\mu > \mu_0$ the p-value is $P(\overline{X} > \overline{X} | \mu = \mu_0)$
- For $H_1: \mu \neq \mu_0$ the p-value is $P(|\overline{X} \mu_0| > |x \mu_0| | \mu = \mu_0)$
 - If $\overline{x} < \mu_0$ then this can be calculated easier by $2 \times P(\overline{X} < \overline{x} | \mu = \mu_0)$
 - If $\overline{x} > \mu_0$ then this can be calculated easier by $2 \times P(\overline{X} > \overline{x} | \mu = \mu_0)$

How do I find the critical value and critical region for a one-sample *z*-test?

- The critical region is determined by the significance level α%
 - For $H_1: \mu < \mu_0$ the critical region is $\overline{X} < c$ where $P(\overline{X} < c | \mu = \mu_0) = \alpha\%$
 - For $H_1: \mu > \mu_0$ the critical region is $\overline{X} > c$ where $P(\overline{X} > c | \mu = \mu_0) = \alpha\%$
 - For $H_1: \mu \neq \mu_0$ the critical regions are $\overline{X} < c_1$ and $\overline{X} > c_2$ where

$$P(\overline{X} < c_1 | \mu = \mu_0) = P(\overline{X} > c_2 | \mu = \mu_0) = \frac{1}{2} \alpha\%$$

- The critical value(s) can be found using the inverse normal distribution function
 - When rounding the critical value(s) you should choose:
 - The lower bound for the inequalities $\overline{X} < c$
 - The upper bound for the inequalities X > c
 - This is so that the probability does not exceed the significance level

Exam Tip

- Exam questions might specify a method for you to use so practise all methods (using GDC, p-values, critical regions)
- If the exam question does not specify a method then use whichever method you want
 - Make it clear which method you are using
 - You can always use a second method as a way of checking your answer



The mass of a Burmese cat, C, follows a normal distribution with mean 4.2 kg and a standard deviation 1.3 kg. Kamala, a cat breeder, claims that Burmese cats weigh more than the average if they live in a household which contains young children. To test her claim, Kamala takes a random sample of 25 cats that live in households containing young children.

a)

2

State the null and alternative hypotheses to test Kamala's claim.



```
b)
```

Using a 5% level of significance, find the critical region for this test.



c)

Kamala calculates the mean of the 25 cats included in her sample to be 4.65 kg. Determine the conclusion of the test.

4.65 > 4.6276... so 4.65 is in critical region

Reject Ho as test statistic is in critical region. There is sufficient evidence to suggest that Burmese cats weigh more if they live in a household which contains young children.



One-Sample t-tests

What is a one-sample t-test?

- A one-sample t-test is used to test the mean (µ) of a normally distributed population
 - You use a t-test when the **population variance (***o*²**) is unknown**
 - You need to use the unbiased estimate for the population variance (s_{n-1}^2)
- The mean of a sample of size n is calculated \overline{x} and a t-distribution is used to test it
 - t-distributions are similar to normal distributions
 - As the sample size gets larger the t-distribution tends towards the standard normal distribution
- You won't be expected to find the critical value
 - The p-value can be found using the test function on your GDC

What are the steps for performing a one-sample t-test on my GDC?

- STEP 1: Write the hypotheses
 - $\circ H_0: \mu = \mu_0$
 - Clearly state that μ represents the **population mean**
 - μ₀ is the assumed population mean
 - For a **one-tailed** test $H_1: \mu < \mu_0$ or $H_1: \mu > \mu_0$
 - For a **two-tailed** test: $H_1: \mu \neq \mu_0$
 - The alternative hypothesis will depend on what is being tested
- STEP 2: Enter the data into your GDC and choose the one-sample t-test
 - If you have the raw data APERS PRACTICE
 - Enter the data as a list
 - If you have summary statistics
 - Enter the values of \overline{x} , s_{n-1} (sometimes written as s_x on a GDC) and n
 - Your GDC will give you the p-value
- STEP 3: Decide whether there is evidence to reject the null hypothesis
 - If the p-value < significance level then reject H₀
- STEP 4: Write your conclusion
 - If you reject H₀ then there is evidence to suggest that...
 - The mean has decreased (for $H_1: \mu < \mu_0$)
 - The mean has increased (for $H_1: \mu > \mu_0$)
 - The mean has changed (for $H_1: \mu \neq \mu_0$)
 - If you **accept H₀** then there is **insufficient evidence** to reject the null hypothesis which suggests that...
 - The mean has not decreased (for $H_1: \mu < \mu_0$)
 - The mean has not increased (for $H_1: \mu > \mu_0$)
 - The mean has not changed (for $H_1: \mu \neq \mu_0$)



The IQ of a student at Calculus High can be modelled as a normal distribution with mean 126. The headteacher decides to play classical music during lunchtimes and suspects that this has caused a change in the average IQ of the students.

a)

State the null and alternative hypotheses to test the headteacher's suspicion.



b)

The headteacher selects 15 students and asks them to complete an IQ test. The mean score for the sample is 127.1 and the sample variance is 14.7. Calculate the unbiased estimate for the population variance s_{n-1}^2 .

Formula booklet
Unbiased estimate of
population variance
$$s_{n-1}^2 = \frac{n}{n-1}s_n^2$$

 $s_{n-1}^2 = \frac{15}{14} \times 14.7$ APERS PRACTICE
 $s_{n-1}^2 = 15.75$

c)

Calculate the p-value for the test.

The population variance is unknown so use a t-test Enter summary statistics into GDC using one-sample t-test $\overline{x} = 127.1$ $S_{n-1} = \sqrt{15.75}$ n = 15 p = 0.3012...p = 0.301 (3sf)

d)

State whether the headteacher's suspicion is supported by the test.



```
0.3012... > 0.1
```

Accept H₀ as p-value > significance level. There is insufficient evidence to support the headteacher's suspicion.





4.12.2 Hypothesis Testing for Mean (Two Sample)

Two-Sample Tests

What is a two-sample test?

- A two-sample test is used to compare the means (μ₁ & μ₂) of two normally distributed populations
 - You use a *z*-test when the population variances ($\sigma_1^2 \& \sigma_2^2$) are known
 - You use a t-test when the population variances are unknown
 - In this course you will assume the variances are equal and use a pooled sample for a t-test
 - In a pooled sample the data from both samples are used to estimate the population variance

What are the steps for performing a two-sample test on my GDC?

• STEP 1: Write the hypotheses

- $H_0: \mu_1 = \mu_2$
 - Clearly state that µ1& µ2 represent the population means
 - Make sure you make it clear which mean corresponds to which population
 - In words this means that the two population means are equal
- For a **one-tailed** test $H_1: \mu_1 < \mu_2$ or $H_1: \mu_1 > \mu_2$
- For a **two-tailed** test: $H_1: \mu_1 \neq \mu_2$
- The alternative hypothesis will depend on what is being tested
- STEP 2: Decide if it is a z-test or a t-test ERS PRACTICE
 - If the populations variances are **known** then use a **z-test**
 - If the populations variances are **unknown** then use a **t-test**
 - Assume the variances are equal and use a pooled sample
- STEP 3: Enter the data into your GDC and choose two-sample z-test or two-sample ttest
 - If you have the raw data
 - Enter the data as a list
 - Enter the values of $\sigma_1 \& \sigma_2$ if a z-test
 - Choose the pooled option if a t-test
 - If you have summary statistics (only for a z-test)
 - Enter the values of \overline{x}_1 , \overline{x}_2 , σ_1 , σ_2 , $n_1 \& n_2$
 - Your GDC will give you the p-value
- STEP 4: Decide whether there is evidence to reject the null hypothesis
 - If the p-value < significance level then reject H₀
- STEP 5: Write your conclusion
 - If you reject H₀ then there is evidence to suggest that...
 - The mean of the 1st population is smaller (for H₁: $\mu_1 < \mu_2$)
 - The mean of the 1st population is bigger (for $H_1: \mu_1 > \mu_2$)
 - The means of the two populations are different (for $H_1: \mu_1 \neq \mu_2$)

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- If you accept H₀ then there is insufficient evidence to reject the null hypothesis which suggests that...
 - The mean of the 1st population is not smaller (for H₁: $\mu_1 < \mu_2$)
 - The mean of the 1st population is not bigger (for H₁: $\mu_1 > \mu_2$)
 - The means of the two populations are not different (for $H_1: \mu_1 \neq \mu_2$)





The times (in minutes) for children and adults to complete a puzzle are recorded below.

Children	3.1	2.7	3.5	3.1	2.9	3.2	3.0	2.9		
Adults	3.1	3.6	3.5	3.6	2.9	3.6	3.4	3.6	3.7	3.0

The creator of the puzzle claims children are generally faster at solving the puzzle than adults. A t-test is to be performed at a 1% significance level.

a)

b)

C)

Write down the null and alternative hypotheses.



Reject Ho as p-value < significance level. There is sufficient evidence to suggest that children are generally faster at solving the puzzle than adults. This supports the creator's claim.



Paired t-tests

What is a paired t-test?

- A paired test is where you take **two samples** but **each data point from one sample can be paired with a data point from the other sample**
 - These are used when one group of members are used twice and the two results for each member are paired
 - It could be to compare the sample before and after introducing a new factor
 - It could be to compare the sample under two different conditions
- For this test you use the differences between the pairs and treat them as one sample
 - As the variance of the differences is unlikely to be known you will use a t-test
 - For a paired test you need to assume the differences are normally distributed
 - You don't need to assume the populations are normally distributed

What are the steps for performing a paired t-test on my GDC?

• STEP 1: Write the hypotheses

- $H_0: \mu_D = 0$
 - Clearly state that µ_D represents the population mean of the differences
 - In words this means the population mean has not changed
- For a **one-tailed** test $H_1: \mu_D < 0$ or $H_1: \mu_D > 0$
- For a two-tailed test: $H_1: \mu_D \neq 0$
 - The alternative hypothesis will depend on what is being tested
- STEP 2: Enter the data into your GDC and choose the one-sample t-test
 - Enter the differences as a list
 - Be consistent with the order in which you subtract paired values
 - Your GDC will give you the p-value
- STEP 3: Decide whether there is evidence to reject the null hypothesis
 - $\circ~$ If the p-value < significance level then reject H_0
- STEP 4: Write your conclusion
 - If you reject H₀ then there is evidence to suggest that...
 - The mean has decreased (for $H_1: \mu_D < 0$)
 - The mean has increased (for $H_1: \mu_D > 0$)
 - The mean has changed (for $H_1: \mu_D \neq 0$)
 - If you **accept H₀** then there is **insufficient evidence** to reject the null which suggests that...
 - The mean has not decreased (for $H_1: \mu_D < 0$)
 - The mean has not increased (for $H_1: \mu_D > 0$)
 - The mean has not changed (for $H_1: \mu_D \neq 0$)

Exam Tip

- If an exam question has two samples with the same number of members then consider carefully whether it makes sense to do a paired test or a two sample test
- The examiner might make it look like it is a paired test to trick you!



2

In a school all students must study French and Spanish. 9 students are selected and complete a test in both subjects, the standardised scores are shown below

Student	1	2	3	4	5	6	7	8	9
French score	61	82	77	80	99	69	75	71	81
Spanish score	74	79	83	66	95	79	82	81	85

The headteacher wants to investigate whether there is a difference in the students' scores between the two subjects. A paired t-test is to be performed at a 10% significance level.

```
a)
```

Write down the null and alternative hypotheses. Let D be the French score minus the Spanish score for the students. Let μ_D be the mean difference for the whole population of students. Ho: $\mu_D = 0$ Ho: $\mu_D = 0$ Testing for a difference in scores

b)

C)

Find the p-value for this test.

Calcu	late the	diffi	eren	e fo	r ea	ch	stud	ent	d=	Frenc	h -Spanish
	Student	1	2	3	4	5	6	7	8	9	
	Student d	-13	3	-6	14	4	-10	-7	-10	-4	
	the dif		les	into	the	<u> </u>		ind	use	a	t-test
p = 0	2958										
· · ·											

Write down the conclusion to the test. Give a reason for your answer.



0.2958... > 0.1

Accept H_0 as p-value > significance level. There is insufficient evidence to suggest that there is a difference in scores.





4.12.3 Binomial Hypothesis Testing

Binomial Hypothesis Testing

What is a hypothesis test using a binomial distribution?

- You can use **a binomial distribution** to test whether the **proportion** of a population with a specified characteristic has **increased** or **decreased**
 - These tests will always be **one-tailed**
 - You will not be expected to perform a two-tailed hypothesis test with the binomial distribution
- A sample will be taken and the **test statistic** *x* will be the **number of members with the characteristic** which will be tested using the distribution $X \sim B(n, p)$
 - This can be thought of as the number of successes

What are the steps for a hypothesis test of a binomial proportion?

- STEP 1: Write the hypotheses
 - $\circ H_0: p = p_0$
 - Clearly state that p represents the population proportion
 - p₀ is the assumed population proportion
 - $H_1: p < p_0 \text{ or } H_1: p > p_0$
- STEP 2: Calculate the p-value or find the critical region
 - See below
- STEP 3: Decide whether there is evidence to reject the null hypothesis
 - \circ If the p-value < significance level then reject H₀
 - If the test statistic is in the critical region then reject H_0 CT [CE
- STEP 4: Write your conclusion
 - If you reject H₀ then there is evidence to suggest that...
 - The population proportion has decreased (for H₁: p < p₀)
 - The population proportion has increased (for H₁: p > p₀)
 - If you **accept H**₀ then there is **insufficient evidence** to reject the null hypothesis which suggests that...
 - The population proportion has not decreased (for $H_1: p < p_0$)
 - The population proportion has not increased (for H₁: p > p₀)

How do I calculate the p-value?

- The p-value is determined by the **test statistic** x
- The p-value is the probability that 'a value being **at least as extreme** as the test statistic' would occur if **null hypothesis were true**
 - For H₁: $p < p_0$ the p-value is $P(X \le x | p = p_0)$
 - For H₁: $p > p_0$ the p-value is $P(X \ge x | p = p_0)$

How do I find the critical value and critical region?

- The critical value and critical region are determined by the significance level α%
- Your calculator might have an **inverse binomial function** that works just like the inverse normal function



- You need to use this value to find the critical value
- The value given by the inverse binomial function is normally one away from the actual critical value
- For $H_1: p < p_0$ the critical region is $X \le c$ where c is the critical value
 - c is the largest integer such that $P(X \le c | p = p_0) \le \alpha\%$
 - Check that $P(X \le c+1 | p = p_0) > \alpha\%$
- For $H_1: p > p_0$ the critical region is $X \ge c$ where c is the critical value
 - c is the smallest integer such that $P(X \ge c | p = p_0) \le \alpha\%$
 - Check that $P(X \ge c 1 | p = p_0) > \alpha\%$





The existing treatment for a disease is known to be effective in 85% of cases. Dr Sabir develops a new treatment which she claims is more effective than the existing one. To test her claim she uses the new treatment on a random sample of 60 patients with the disease and finds that the treatment was effective for 57 of them.

a)

b)

State null and alternative hypotheses to test Dr Sabir's claim.

```
Let p be the proportion of the population for which the new
         treatment is effective.
         Ho: p=0.85
H1: p>0.85 Testing for an increase
Perform the test using a 1% significance level. Clearly state the conclusion in
context.
         Let X \sim B(60, p) be the number of people in the sample
         for which the new treatment is effective
         Find the p-values and compare to the significance level
          p = P(X \ge 57 | p = 0.85) = 0.01483... > 0.01
          Accept Ho as p-value > significance level.
          There is insufficient evidence to suggest that
           the new treatment is more effective than the
           existing treatment.
```



4.12.4 Poisson Hypothesis Testing

Poisson Hypothesis Testing

What is a hypothesis test using a Poisson distribution?

- You can use a Poisson distribution to test whether the mean number of occurrences for a given time period within a population has increased or decreased
 - These tests will always be **one-tailed**
 - You will not be expected to perform a two-tailed hypothesis test with the Poisson distribution
- A sample will be taken and the **test statistic** *x* will be the **number of occurrences** which will be tested using the distribution *X* ~ Po(*m*)

What are the steps for a hypothesis test of a Poisson proportion?

STEP 1: Write the hypotheses

- $H_0: m = m_0$
 - Clearly state that *m* represents the mean number of occurrences for the given time period
 - m₀ is the assumed mean number of occurrences
 - You might have to use proportion to find m₀
- $H_1: m < m_0 \text{ or } H_1: m > m_0$
- STEP 2: Calculate the *p*-value or find the critical region
 - See below
- STEP 3: Decide whether there is evidence to reject the null hypothesis
 - If the p-value < significance level then reject H₀
 - $\circ~$ If the test statistic is in the critical region then reject ${\rm H}_{\rm O}$
- STEP 4: Write your conclusion
 - If you **reject H₀** then there is evidence to suggest that...
 - The mean number of occurrences has decreased (for $H_1: m < m_0$)
 - The mean number of occurrences has increased (for H₁: m > m₀)
 - If you **accept H₀** then there is **insufficient evidence** to reject the null hypothesis which suggests that...
 - The mean number of occurrences has not decreased (for H₁: m < m₀)
 - The mean number of occurrences has not increased (for H₁:m > m₀)

How do I calculate the p-value?

- The p-value is determined by the test statistic x
- The *p*-value is the probability that 'a value being **at least as extreme** as the test statistic' would occur if **null hypothesis were true**
 - For H₁: $m < m_0$ the p-value is $P(X \le x \mid m = m_0)$
 - For H₁: $m > m_0$ the p-value is $P(X \ge x | m = m_0)$

How do I find the critical value and critical region?

• The critical value and critical region are determined by the significance level $\alpha\%$



- Your calculator might have an **inverse Poisson function** that works just like the inverse normal function
 - You need to use this value to find the critical value
 - The value given by the inverse Poisson function is normally one away from the actual critical value
- For $H_1: m < m_0$ the critical region is $X \le c$ where c is the critical value
 - c is the largest integer such that $P(X \le c \mid m = m_0) \le \alpha\%$
 - Check that $P(X \le c+1 \mid m = m_0) > \alpha\%$
- For $H_1: m > m_0$ the critical region is $X \ge c$ where c is the critical value
 - c is the smallest integer such that $P(X \ge c \mid m = m_0) \le \alpha\%$
 - Check that $P(X \ge c 1 \mid m = m_0) > \alpha\%$



Exam Tip

- In an examit is very important to state the time period for your variable
- Make sure the mean used in the null hypothesis is for the stated time period





The owner of a website claims that his website receives an average of 120 hits per hour. An interested purchaser believes the website receives on average fewer hits than they claim. The owner chooses a 10-minute period and observes that the website receives 11 hits. It is assumed that the number of hits the website receives in any given time period follows a Poisson Distribution.

```
a)
```

State null and alternative hypotheses to test the purchaser's claim.



```
b)
```

Find the critical region for a hypothesis test at the 5% significance level.

Let $X \sim Po(m)$ be the number of hits in a 10-minute period The critical value c is the largest value such that $P(X \le c \mid m=20) \le 0.05$ You can use the inverse Poisson function on the GDC to decide which value to check first $P(X \le 13 \mid m=20) = 0.0661... > 0.05$ Too big so reduce the region $P(X \le 12 \mid m=20) = 0.0390... \le 0.05$

(ritical region X≤12

c)

Perform the test using a 5% significance level. Clearly state the conclusion in context.

11 < 12 so 11 is in the critical region

Reject Ho as test statistic is in critical region. There is sufficient evidence to suggest that the website receives on average fewer hits than they claim.



4.12.5 Hypothesis Testing for Correlation

Hypothesis Testing for Correlation

What is a hypothesis test for correlation?

- You can use a **t-test** to test whether there is **linear correlation** between two normally distributed variables
 - If specifically testing for positive (or negative) linear correlation then a **one-tailed test** is used
 - If testing for any linear correlation then a **two-tailed test** is used
- A sample will be taken and the **raw data** will be given
 - You might be asked to calculate the **PMCC (Pearson's product-moment correlation coefficient)**

What are the steps for a hypothesis test for correlation?

• STEP 1: Write the hypotheses

- ∘ H₀:ρ=Ο
 - Clearly state that prepresents population correlation coefficient between the two variables
 - In words this means there is no correlation
- $H_1: \rho < 0, H_1: \rho > 0 \text{ or } H_1: \rho \neq 0$
- STEP 2: Calculate the p-value or the PMCC
 - Choose a t-test for linear regression
 - Enter the data as two lists into GDCERS PRACTIC
- STEP 3: Decide whether there is evidence to reject the null hypothesis
 - \circ If the *p*-value < significance level then reject H₀
 - $\circ~$ If the absolute value of the PMCC is bigger than the absolute value of the critical value then reject H_0
 - If you are expected to use the PMCC you will be given the critical value in the exam
- STEP 4: Write your conclusion
 - If you reject H₀ then there is evidence to suggest that...
 - There is a negative linear correlation between the two variables (for $H_1: \rho < 0$)
 - There is a positive linear correlation between the two variables (for $H_1: \rho > 0$)
 - There is a linear correlation between the two variables (for $H_1: \rho \neq 0$)
 - If you **accept H₀** then there is **insufficient evidence** to reject the null hypothesis which suggests that...
 - There is not a negative linear correlation between the two variables (for $H_1: \rho < 0$)
 - There is not a positive linear correlation between the two variables (for $H_1: \rho > 0$)
 - There is not a linear correlation between the two variables (for $H_1: \rho \neq 0$)



Jessica wants to test whether there is any linear correlation between the distance she runs in a day, d km, and the amount of sleep she has the night after her run, t hours. Over the period of a month she takes a random sample of 9 days, the results are recorded in the table.

Distance (<i>d</i> km)	1.2	2.3	1.5	1.3	2.5	1.8	1.9	2.0	1.1
Sleep (<i>t</i> hours)	7.9	8.1	7.6	7.3	8.1	8.4	7.8	7.9	6.8

a)

Write down null and alternative hypotheses that Jessica can use for her test.

Let ρ be the correlation coefficient between Jessica's distances and the hours of sleep she gets. H₀: $\rho = 0$ H₁: $\rho \neq 0$ Testing for any linear correlation

b)

Perform the hypothesis test for linear correlation using a 5% significance level. Clearly state your conclusion.

```
Type the data in GDC and select a t-test for linear regression

p = 0.03833... < 0.05

Reject H<sub>0</sub> as p-value < significance level.
```

There is sufficient evidence to suggest that there is a linear correlation between the distance that Jessica runs and the hours she sleeps.



4.12.6 Type I & Type II Errors

Type I & Type II Errors

What are Type I & Type II errors?

- There are four possible outcomes of a hypothesis test:
 - H₀ is false and H₀ is rejected
 - $\circ~H_0$ is true and H_0 is not rejected
 - The test is accurate for these two outcomes
 - $\circ~H_0$ is true and H_0 is rejected
 - $\circ~H_0$ is false and H_0 is not rejected
 - The test has led to an error for these two outcomes
- A Type I error occurs when a hypothesis test gives sufficient evidence to reject H₀ despite it being true
 - This is sometimes called a "false positive"
 - In a court case this would be when the defendant is found **guilty despite being** innocent
- A Type II error is when a hypothesis test gives insufficient evidence to reject H₀ despite it being false
 - This is sometimes called a "false negative"
 - In a court case this would be when the defendant is found innocent despite being guilty



How do I find the probabilities of a Type I or Type II error occurring?

- You should calculate the probability of errors occurring before a sample is taken
- The probabilities are determined by the critical region
 - Equally it is determined by the significance level α%
 - · Critical regions are determined such that:
 - They keep the probability of a Type I error less than or equal to the significance level
 - They maximise the probability of a Type l error
- The probability of a **Type I error** occurring is equal to the probability of **being in the critical region** if H₀ were true
 - P(Type I error) = P(being in the critical region | H₀ is true)
 - For a continuous distribution (normal, t, χ^2)
 - P(Type | error) = α %
 - · For a discrete distribution (binomial, Poisson)
 - P(Type | error) $\leq \alpha\%$

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- The probability of a **Type II** error occurring is equal to the probability of **not being in the critical region** given the actual value of the population parameter
 - P(Type II error) = P(not being in critical region | actual population parameter)
 - You need to know the actual population parameter in order to find the probability of a Type II error
- Once a sample has been taken you can determine which error could have occurred
 - $\circ~$ If you $rejected\,H_0$ then you could have made a $Type\,l\,error$
 - $\circ~$ If you $accepted\,H_0$ then you could have made a $Type\,II\,error$

Can I reduce the probabilities of making a Type I or Type II error?

- You can reduce the probability of a Type I error by reducing the significance level
 However this will increase the probability of a Type II error
- You can reduce the probability of a Type II error by increasing the significance level
 However this will increase the probability of a Type I error
- The only way to reduce both probabilities is by increasing the size of the sample

Exam Tip

- If an exam question asks you to find the probability of a Type I or II error then double check that the test has not been carried out yet
- The examiner could test your understanding of errors by asking you to state which error could not have occurred once the test has been carried out

EXAM PAPERS PRACTICE







4.13 Transition Matrices & Markov Chains

4.13.1 Markov Chains

Markov Chains

What is meant by a "state"?

- States refer to mutually exclusive events with the current event able to change over time
- Examples of states include:
 - Daily weather conditions
 - The states could be: "sunny" and "not sunny"
 - Countries visited by an inspector each day
 - The states could be: "France", "Spain" and "Germany"
 - Store chosen for weekly grocery shop:
 - The states could be: "Foods-U-Like", "Smiley Shoppers" and "Better Buys"

What is a Markov chain?

- A Markov chain is a model that describes a sequence of states over a period of time
 - Time is measured in discrete steps
 - Such as days, months, years, etc
- The conditions for a Markov chain are:
 - The probability of a state being the next state in the sequence only depends on the current state
 - For example PAPERS PRACTICE
 The 11th state only depends on the 10th state
 - The first 9 states **do not affect** the 11th state
 - This probability is called a transition probability
 - The transition probabilities do not change over time
 - For example
 - The probability that the 11th state is A given that the 10th state is B is equal to the probability that the 12th state is A given that the 11th state is B
- A Markov chain is said to be **regular** if it possible to reach any state after a finite period of time regardless of the initial state

What is a transition state diagram?

- A transition diagram is a directed graph
 - The vertices are the states
 - The edges represent the transition probabilities between the states
- The graph can contain
 - Loops
 - These will be the transition probabilities of the next state being the same as the current state
 - Two edges between each pair of vertices
 - The edges will be in opposite directions

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- Each edge will show the transition probability of the state changing in the given direction
- The probabilities on the edges coming out of a vertex add up to 1



Exam Tip

• Drawing a transition state diagram (even when the question does not ask for one) can help you visualise the problem





Fleur travels to work by car, bike or bus. Each day she chooses her mode of transport based on the transport she chose the previous day.

- If Fleur travels by car then there is a 40% chance that she will travel by car the following day and a 10% chance that she will travel by bike.
- If Fleur travels by bike then there is a 60% chance that she will travel by bike the following day and a 25% chance that she will travel by bus.
- If Fleur travels by bus then there is an 80% chance that she will travel by bike the following day and a 20% chance that she will travel by car.

Represent this information as a transition state diagram.





4.13.2 Transition Matrices

Transition Matrices

What is a transition matrix?

- A transition matrix *T* shows the transition probabilities between the current state and the next state
 - The columns represent the current states
 - The rows represent the next states
- The element of **T** in the *i*th row and *j*th column gives the transition probability *t_{ij}* of :
 - the next state being the state corresponding to row i
 - given that the current state is the state corresponding to column j
- The probabilities in each column must add up to 1
- The transition matrix depends on how you assign the states to the columns
 - Each transition matrix for a Markov chain will contain the same elements
 - The rows and columns may be in different orders though
 - E.g. Sunny (S) & Cloudy (C) could be in the order S then C or C then S

What is an initial state probability matrix?

- An initial state probability matrix s₀ is a column vector which contains the probabilities of each state being chosen as the initial state
 - If you know which state was chosen as the initial state then that entry will be 1 and the others will all be zero
- You can find the **state probability matrix** s₁ which contains the probabilities of each state being chosen after **one interval of time**
 - \circ s₁ = Ts₀

How do I find expected values after one interval of time?

- Suppose the Markov change represents a population moving between states
 Examples include:
 - People in a town switching gyms each year
 - Children choosing a type of sandwich for their lunch each day
- Suppose the total population is fixed and equals N
- You can **multiply the state probability matrix s₁** by *N* to find the expected number of members of the population at each state

Exam Tip

- If you are asked to find a transition matrix, check that all the probabilities within a column add up to 1
- Drawing a transition state diagram can help you to visualise the problem



Each year Jamie donates to one of three charities: A, B or C. At the start of each year, the probabilities of Jamie continuing donate to the same charity or changing charities are represented by the following transition state diagram:



a)

Write down a transition matrix \boldsymbol{T} for this system of probabilities.

		Cur	rent st	ate						
EXA	Ņ		ß	DE	RS	Ρ	10.5	62	0.6	ŀ
tate	A	0.5	0.2	0.6		т.	1		1	
t s	B	0.2	0.7	0.4		13	0.2	0.7	0.4	
Nex	C	0.3	0.1	0			0.3	0.1	0 /	

b)

There is a 10% chance that charity A is the first charity that Jamie chooses, a 10% chance for charity B and an 80% chance for charity C. Find the charity which has the highest probability of being picked as the second charity after the first year.



Write down the initial state vector
$$s_0 = \begin{pmatrix} 0.1 \\ 0.1 \\ 0.8 \end{pmatrix}$$

 $s_1 = Ts_0$ $s_1 = \begin{pmatrix} 0.5 & 0.2 & 0.6 \\ 0.2 & 0.7 & 0.4 \\ 0.3 & 0.1 & 0 \end{pmatrix} \begin{pmatrix} 0.1 \\ 0.1 \\ 0.8 \end{pmatrix} = \begin{pmatrix} 0.55 \\ 0.41 \\ 0.04 \end{pmatrix}$
Charity A has the highest probability of bei

Charity A has the highest probability of being the second charity picked.





Powers of Transition Matrices

How do I find powers of a transition matrix?

- You can simply use your GDC to find given powers of a matrix
- The power could be left in terms of an **unknown** n
 - In this case it would be more helpful to write the transition matrix in diagonalised form
 - (see section 1.8.2 Applications of Matrices) T = PDP⁻¹ where
 - D is a diagonal matrix of the eigenvalues
 - P is a matrix of corresponding eigenvectors
 - Then *Tⁿ* = *PDⁿP⁻¹*
 - This is given in the formula booklet
 - Every transition matrix always has an eigenvalue equal to 1

What is represented by the powers of a transition matrix?

- The powers of a transition matrix also represent probabilities
- The element of **T**ⁿ in the *i*th row and *j*th column gives the **probability** tⁿ*ij* of :
 - the future state after n intervals of time being the state corresponding to row i
 - given that the current state is the state corresponding to column j
- For example: Let **T** be a transition matrix with the element t_{2,3} representing the probability that tomorrow is sunny given that it is raining today
 - The element t⁵_{2,3} of the matrix T⁵ represents the probability that it is sunny in 5 days' time given that it is raining today
- The probabilities in each column must still add up to 1 ACT CE

How do I find the column state matrices?

- The column state matrix **s**_n is a column vector which contains the **probabilities** of each state being chosen after *n* intervals of time given the current state
 - $\circ s_n$ depends on s_0
- To calculate the column state matrix you raise the transition matrix to the power *n* and multiply by the initial state matrix
 - $\circ \quad \boldsymbol{T}^n \boldsymbol{s}_0 = \boldsymbol{s}_n$
 - You are given this in the formula booklet
- You can multiply **s**_n by the fixed population size to find the expected number of members of the population at each state after *n* intervals of time



At a cat sanctuary there are 1000 cats. If a cat is brushed on a given day, then the probability it is brushed the following day is 0.2. If a cat is not brushed on a given day, then the probability that is will be brushed the following day is 0.9.

The transition matrix **T** is used to model this information with $T = \begin{pmatrix} 0.2 & 0.9 \\ 0.8 & 0.1 \end{pmatrix}$.

a)

On Monday Hippo the cat is brushed. Find the probability that Hippo will be brushed on Friday.





On Monday 700 cats were brushed. Find the expected number of cats that will be brushed on the following Monday.

> On Monday 700 brushed $S_0 = \begin{pmatrix} 0.7\\ 0.3 \end{pmatrix}$ Expected numbers after 7 days Total × Sz = Total × T^{*}sn $|000 \times \begin{pmatrix} 0.2 & 0.9 \\ 0.6 & 0.1 \end{pmatrix}^{\frac{3}{2}} \begin{pmatrix} 0.7 \\ 0.3 \end{pmatrix} = \begin{pmatrix} 0.2 & 0.9 \\ 0.6 & 0.1 \end{pmatrix}^{\frac{3}{2}} \begin{pmatrix} 700 \\ 300 \end{pmatrix} = \begin{pmatrix} 515 \cdot 36309 \\ 484 \cdot 63691 \end{pmatrix} = \begin{pmatrix} 616 - 616 \\ 616 \end{pmatrix} = \begin{pmatrix} 616 - 616 \\ 616 \\ 616 \end{pmatrix} = \begin{pmatrix} 616 - 616 \\ 6$ 515 cats



Steady State & Long-term Probabilities

What is the steady state of a regular Markov chain?

- The vector **s** is said to be a **steady state** vector if it does not change when multiplied by the transition matrix
 - ∘ **Ts**=s
- Regular Markov chains have steady states
 - A Markov chain is said to be regular if there exists a positive integer k such that none of the entries are equal to 0 in the matrix T^k
 - For this course all Markov chains will be regular
- The transition matrix for a regular Markov chain will have **exactly one** eigenvalue equal to 1 and the **rest will all be less than 1**
- As n gets bigger **T**ⁿ tends to a matrix where **each column is identical**
 - The column matrix formed by using **one of these columns** is called the steady state column matrix **s**
 - This means that the long-term probabilities tend to fixed probabilities
 - s_n tends to s

How do I use long-term probabilities to find the steady state?

- As **T**ⁿ tends to a matrix whose columns equal the steady state vector
 - Calculate Tⁿ for a large value of n using your GDC
 - If the columns are identical when rounded to a required degree of accuracy then the column is the steady state vector
 - If the columns are not identical then choose a higher power and repeat

How do I find the exact steady state probabilities?

- As **Ts** = **s** the steady state vector **s** is the eigenvector of **T** corresponding to the eigenvalue equal to 1 whose elements sum to 1:
 - Let **s** have entries x₁, x₂, ..., x_n
 - Use **Ts** = **s** to form a system of linear equations
 - There will be an infinite number of solutions so choose a value for one of the unknowns
 - For example: let x_n = 1
 - Ignoring the last equation solve the system of linear equations to find $x_1, x_2, ..., x_{n-1}$
 - Divide each value x_i by the sum of the values
 - This makes the values add up to 1
- You might be asked to **show this result using diagonalisation**
 - Write T = PDP⁻¹ where D is the diagonal matrix of eigenvalues and P is the matrix of eigenvectors
 - Use $T^n = PD^nP^{-1}$
 - As *n* gets large **D**^{*n*} tends to a matrix where all entries are 0 apart from one entry of 1 due to the eigenvalue of 1
 - Calculate the limit of **T**ⁿ which will have **identical columns**
 - You can calculate this by multiplying the three matrices (**P**, **D**[∞], **P**⁻¹) together





Exam Tip

- If you calculate T^∞ by hand then a quick check is to see if the columns are identical







If a cat is brushed on a given day, then the probability it is brushed the following day is 0.2. If a cat is not brushed on a given day, then the probability that is will be brushed the following day is 0.9.

The transition matrix **T** is used to model this information with $T = \begin{pmatrix} 0.2 & 0.9 \\ 0.8 & 0.1 \end{pmatrix}$.

a)

Find an eigenvector of T corresponding to the eigenvalue 1.



b)

Hence find the steady state vector.

Scale the elements so that they add to $\begin{bmatrix} q \\ 17 \\ \frac{8}{17} \end{bmatrix}$ The eigenvector corresponding to the eigenvalue I, whose elements add to I, is the steady state vector.