Unit 10.2. MCQs Set 1

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Results



#1. Q1. In descriptive statistics, which of the following is a measure of central tendency that splits the dataset into two equal parts?

(A) Mean
(B) Mode

☐ (C) Median

☐ (D) Variance

The median is the middle value when the data is arranged in order.

#2. Q2. The arithmetic mean is generally suitable for numerical data, but can be misleading if the data:

(A) Is normally distributed

(B) Has no outliers

(C) Is highly skewed or has extreme outliers

(D) Consists only of positive integers

#3. Q3. Which of these is calculated by summing all values and dividing by the number of observations?

(A) Median

(B) Mode

□ (C) Mean

(D) Standard error

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The arithmetic mean is computed by dividing the total sum by the number of data points.
#4. Q4. (Fill in the blank) The is defined as the most frequently occurring value in a dataset.
□ (A) Mean □
(B) Median □
(C) Mode
(D) Range
The mode is the value that appears most frequently in the dataset.
#5. Q5. Descriptive statistics typically do not include:
(A) Calculating measures like mean, median, and mode □
(B) Drawing inferences about a population from a sample □
(C) Displaying data in histograms or boxplots
(D) Summarizing variability with standard deviation or range
Descriptive statistics summarize the data at hand; inferential statistics are used to draw conclusions about a broader population.
#6. Q6. Which statement about Analysis of Variance (ANOVA) is correct?
□ (A) It's used only for two-sample comparisons
(B) It tests if more than two group means differ significantly
(C) It discards within-group variance entirely
(D) It is a commonly used tool in biostatistics
ANOVA is used to test for significant differences among the means of three or more groups.
#7. Q7. A t-test for independent samples is typically used when:
(A) The same group is measured before and after an intervention
□ (B) You have two unrelated groups to compare on a continuous outcome
□ (C) You have more than two groups to compare
□ (D) The data is strictly categorical

#8. Q8. (Fill in the blank) In hypothesis testing, "power" is the probability of correctly

An independent t-test compares the means of two distinct groups on a continuous variable.

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rejecting the null hypothesis when it is
□ (A) True
(B) Valid □ (C) False
(C) False
(D) Undefined Power is defined as $1 - \beta$, the probability of detecting an effect when the null hypothesis is false.
#9. Q9. Sample size calculation often requires:
(A) Desired power, significance level (alpha), and an estimate of the effect size
(B) A fixed sample size regardless of effect size
(C) Ignoring effect size in favor of random sampling
(D) Only the significance level
Accurate sample size calculation depends on power, alpha, and effect size to detect a true effect.
#10. Q10. The p-value in a statistical test typically indicates:
□ (A) The probability that the null hypothesis is true
(B) The probability of observing a result as extreme or more extreme, assuming the null hypothesis is true
(C) The guaranteed correctness of the alternative hypothesis
(D) The margin of error in the measurement
The p-value indicates the probability of obtaining results at least as extreme as the observed ones if the null hypothesis is true.
#11. Q11. Which of the following is not an assumption of a one-way ANOVA?
(A) Each group sample is independent
(B) The data in each group is roughly normally distributed
(C) Homogeneity of variances across groups
□ (D) Each group must contain exactly 100 observations
ANOVA does not require equal sample sizes; each group does not have to have exactly 100 observations.
#12. Q12. (Fill in the blank) A "Type I error" is also known as a in hypothesis testing.
□ (A) False positive
(B) False negative
(C) True positive

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(D) True negative

A Type I error occurs when a true null hypothesis is incorrectly rejected—a false positive.

#13. Q13. The correlation coefficient (Pearson's r) measures: (A) Causal relationships (B) The strength and direction of the linear association between two continuous variables (C) Differences among group means (D) The proportion of variance explained by a regression model Pearson's r quantifies the linear association between two continuous variables, including direction and strength. #14. Q14. If the correlation coefficient r = 0, it implies: (A) No linear relationship between the variables (B) A perfect negative correlation (C) A perfect positive correlation (D) An undefined relationship An r value of 0 indicates no linear association between the variables. #15. Q15. In linear regression, the slope coefficient indicates: (A) The predicted change in the outcome for a one-unit change in the predictor (B) The proportion of variance in the outcome explained by the predictor (C) The value of the intercept (D) The error in predicting the outcome The slope coefficient quantifies the expected change in the dependent variable for a one unit change in the independent variable. #16. Q16. Multiple regression allows for: (A) Only one predictor variable (B) Testing several predictor variables simultaneously to see how they collectively relate to the outcome (C) Ignoring the effects of confounding variables (D) Using only categorical predictors Multiple regression models use more than one predictor to explain the variability in the outcome variable.

#17. Q17. Survival analysis is used to:

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(A) Analyze time-to-event data, such as time until recovery or relapse in clinical studies
(B) Describe the frequency distribution of categorical data
(C) Calculate the mean of continuous data
(D) Compare group medians only
Survival analysis techniques, like Kaplan-Meier and Cox regression, are used to analyze the time until an event occurs including censoring.
#18. Q18. (Fill in the blank) The method estimates and plots survival probabilities
over time for different groups.
□ (A) Log-rank
(B) Cox proportional hazards
C) Kaplan-Meier
(D) Weibull
The Kaplan-Meier method is used to estimate survival probabilities and generate survival curves.
#19. Q19. Genome mapping statistics might involve:
\square (A) Calculating linkage disequilibrium, LOD scores, or association p-values for genetic variants \square
(B) Only describing phenotypic data
(C) Ignoring genetic markers entirely
(D) Aggregating all data into a single average value
Genome mapping employs statistical measures (e.g., LOD scores) to associate genetic variants with traits.
#20. Q20. Bioinformatics often deals with 'big data' from sequencing. The process typicall includes:
□ (A) Data encryption only □
(B) Data cleaning, alignment, variant calling, and statistical association tests $\hfill\Box$
(C) Manual entry of sequence data
(D) Discarding data that are not perfectly clean
Bioinformatics workflows involve multiple steps from data cleaning to statistical analysis.
#21. Q21. Which of these is a nominal type of data?
□ (A) Heights in centimeters
(B) Temperature in Celsius
(C) Blood group categories like A, B, AB, O

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(D) (None)

Nominal data classify observations into distinct categories with no inherent order.

#22. Q22. Ordinal data differs from nominal data in that:
□ (A) They are measured on an interval scale □
(B) Ordinal categories have a ranked order, though differences between ranks are not necessarily equal
(C) They are always numerical
(D) They require no categorization
Ordinal data have an inherent order, unlike nominal data.
#23. Q23. In big data contexts, 'metadata' is:
(A) Data about the analysis methods
□ (B) Data describing other data, such as file name, creation date, and experimental conditions
(C) Irrelevant information
(D) The raw experimental measurements only
Metadata provides essential contextual details that help organize and interpret the primary data.
#24. Q24. 'Multi-dimensional data' might refer to:
(A) Data collected over time only
(B) Datasets with multiple features or variables per observation
□ (C) Data with only one variable □
(D) Data that have no variability
Multi-dimensional data include multiple variables for each observation and require specialized analysis techniques.
#25. Q25. (Fill in the blank) A 'linear algebraic treatment' of data might involve to reduce dimensions or solve large systems.
□ (A) Differential equations □
□ Graph theory □
(C) Matrix operations
□ (D) Logical reasoning
Matrix operations, such as eigenvalue decomposition, are key in reducing dimensions in data analysis.

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#26. Q26. Eigenvalues and eigenvectors are key in:

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(A) Calculating simple averages
(B) Principal Component Analysis (PCA) and dimensionality reduction
LI (C) Determining the median of a dataset
LI (D) Creating pie charts
PCA uses eigen decomposition to identify the principal components that capture the most variance.
#27. Q27. (Fill in the blank) decomposition is a factorization approach used in big data to break a matrix into singular values and orthonormal vectors.
(A) QR
CB) LU
C) SVD (Singular Value Decomposition) C(D) Cholesky
Singular Value Decomposition (SVD) decomposes a matrix into singular values and vectors, widely used for dimensionality reduction.
#28. Q28. Exploratory data analysis (EDA) typically includes:
□ (A) Ignoring data summaries □
(B) Generating summary statistics, histograms, box plots, and scatter plots to detect patterns and outliers
(C) Only applying advanced machine learning models
(D) Discarding any data that do not fit the hypothesis
EDA involves initial data visualization and summarization to understand structure and detect anomalies.
#29. Q29. Descriptive statistics differ from inferential statistics by focusing on:
□ (A) Drawing conclusions about a population
(B) Summarizing datasets using measures like mean and standard deviation without making population inferences
(C) Making predictions for future samples
(D) Testing hypotheses
Descriptive statistics summarize the current dataset without extrapolating to a broader population.
#30. Q30. The 'mean absolute deviation' around the mean is:
\square (A) The average of the squared differences between each data point and the mean
(B) The average of the absolute differences between each data point and the mean
(C) The difference between the maximum and minimum values
□ (D) The standard deviation divided by the mean

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Mean absolute deviation is calculated as the average of the absolute deviations from the mean, providing a measure of dispersion.

#31.	031.	Match	the	following	data	types	with	their	descriptions	::
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- 1. Nominal data;
- 2. Ordinal data;
- 3. Interval data;
- 4. Ratio data
- (a) Rank-ordered categories (e.g., mild, moderate, severe)
- (b) Named categories with no inherent order (e.g., hair color)
- (c) Numeric scale with a meaningful absolute zero (e.g., weight, height)
- (d) Numeric scale with equal intervals but an arbitrary zero (e.g., temperature in °C)

(A) 1-b, 2-a, 3-d, 4-c

(B) 1-a, 2-b, 3-c, 4-d

□ (C) 1-c, 2-d, 3-a, 4-b

(C) 1-C, 2-d, 3-a, 4-

(D) 1-d, 2-c, 3-b, 4-a

Nominal data are categories with no order (b), ordinal data are rank-ordered (a), interval data have equal spacing (d), and ratio data have a true zero (c).

#32. Q32. Survival analysis methods may incorporate 'censoring,' which arises when:

(A) Data collection is halted abruptly

(B) A participant does not experience the event by the study end or is lost to follow-up

(C) The data are missing completely at random

(D) None of the above

Censoring occurs when the event of interest is not observed in some participants, due to dropout or study end.

#33. Q33. The standard error of the mean (SEM) indicates:

(A) The median's variability

(B) How far the sample mean is likely to be from the population mean

(C) The range of the data

(D) The overall sample size

SEM estimates the precision of the sample mean as an estimate of the population mean.

#34. Q34. In correlation analysis, r2 (the coefficient of determination) represents:

(A) The strength of the linear relationship only

(B) The proportion of the variance in one variable that is explained by another

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□ (C) The p-value of the correlation	
(D) The slope of the regression line	
The coefficient of determination (r^2) quantifies the proportion of variance in one variable accounted for by the other.	
#35. Q35. (Fill in the blank) The distribution is often used to model count data, such as the number of events in a fixed time interval.	ch
(A) Exponential	
(B) Normal	
(C) Poisson	
(D) Uniform	
The Poisson distribution is commonly used for modeling the number of occurrences of an event in a fixed interval of times or space.	nε
#36. Q36. The Wilcoxon Rank-Sum (Mann-Whitney) test compares:	
□ (A) Mean values of two groups assuming normality	
□ (B) Two independent groups on ordinal or non-normally distributed continuous data using ranks □	
C) Variances across multiple groups □	
(D) Paired data from the same group	
This non-parametric test compares two independent groups based on rank-order, suitable when data do not follow normal distribution.	v a
#37. Q37. Which statement is incorrect about standard deviation (SD)?	
□ (A) It measures data spread around the mean	
(B) A larger SD indicates greater variability	
C) SD is robust to outliers	
(D) SD is the square root of the variance	
SD is sensitive to outliers, which can significantly affect its value.	
#38. Q38. Chi-square tests are commonly used for:	
□ (A) Comparing means of continuous variables	
□ (B) Examining associations between two categorical variables in contingency tables	
□ C) Analyzing correlation coefficients □ □	
(D) Predicting continuous outcomes	
Chi-square tests analyze categorical data to determine if there is a significant association between variables.	

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#39. Q39. Degrees of freedom in statistical tests generally indicate:
☐ (A) The number of values in the final calculation that are free to vary
□ (B) The total sample size
(C) The number of missing data points
(D) The reliability of the data
Degrees of freedom represent the number of independent values that can vary in a calculation without breaking an constraints.
#40. Q40. If the data is heavily skewed to the right, which measure of central tendency is typically preferred?
□ (A) Mean
(B) Median
(C) Mode
(D) Range
The median is less affected by extreme values and is preferred in skewed distributions.
#41. Q41. Match the following: 1. Parametric test; 2. Non-parametric test; 3. One-sample t-test; 4. Kruskal-Wallis test
(a) Compares the median among more than two groups (a rank-based alternative to ANOVA)
(b) Assumes normal distribution, using means and standard deviations
(c) Compares a single sample mean against a hypothesized value(d) Distribution-free test often using ranks
(A) 1-b, 2-d, 3-c, 4-a (B) 1-d, 2-b, 3-a, 4-c (C) 1-c, 2-b, 3-d, 4-a (D) 1-a, 2-c, 3-b, 4-d
Parametric tests assume a normal distribution (b), non-parametric tests are distribution-free (d), the one-sample t-test compares a sample mean with a hypothesized value (c), and the Kruskal-Wallis test compares medians across groups (a).
#42. Q42. (Fill in the blank with multiple-choice) A distribution typically describes the sum of independent Bernoulli trials—for example, the number of successes in n trials.
□ (A) Poisson
□ (B) Exponential

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SVD decomposes a matrix into singular values and vectors, facilitating dimensionality reduction.
(D) SVD (Singular Value Decomposition)
(C) Cholesky decomposition □
(B) LU decomposition □
□ (A) QR decomposition □
#46. Q46. (Fill in the blank) is the method used to factor a rectangular matrix M into $U \cdot \Sigma \cdot V^T$, which is widely employed for dimensionality reduction in big data.
PCA reduces the number of variables while preserving the most variance in the data.
(D) To calculate simple averages of multiple variables
(C) To standardize data to have zero mean □
(B) Reducing the dimensionality of the dataset while retaining as much variance as possible □
□ (A) To increase the dimensionality of the dataset □
#45. Q45. The primary purpose of PCA (Principal Component Analysis) in multi-dimensional data is:
Eigenvalues are defined by the equation $Av = \lambda v$, where v is a non-zero eigenvector.
(D) The inverse of a matrix
(C) The diagonal elements of the identity matrix
(B) The sum of the elements in a matrix
\Box (A) Scalars λ for which there is a non-zero vector \mathbf{v} satisfying $A\mathbf{v} = \lambda \mathbf{v}$
#44. Q44. In matrix algebra, 'eigenvalues' are best described as:
Metadata comprises additional information that provides context for the primary data.
(D) The raw experimental measurements only
(C) Irrelevant information □
(B) Contextual details such as creation date, experimental conditions, and sample descriptions □
□ (A) Detailed raw measurements only □
#43. Q43. "Metadata" in big data analysis commonly includes:
The binomial distribution models the number of successes in a fixed number of independent Bernoulli trials.
(D) Binomial
(C) Normal
п

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#47. Q47. Exploratory data analysis (EDA) typically includes:
□ (A) Ignoring data summaries completely
(B) Generating summary statistics, histograms, box plots, and scatter plots to identify patterns and outliers
(C) Immediately building predictive models
(D) Discarding any data that do not match expectations
EDA involves a preliminary analysis of data through visualizations and summary statistics to understand its structure.
#40 O40 A (greened in wood to)
#48. Q48. A 'z-score' is used to:
(A) Standardize an individual value by subtracting the mean and dividing by the standard deviation
(B) Calculate the range of the dataset
(C) Determine the median position
(D) Estimate the mode from the data
A z-score standardizes data, showing how many standard deviations an observation is from the mean.
#49. Q49. Which statement is incorrect about inferential statistics?
(A) They allow conclusions about a population based on sample data
(B) Confidence intervals and hypothesis tests are commonly used
(C) They always produce a zero margin of error
(D) They often require assumptions about the data distribution
Inferential statistics always involve some level of uncertainty; a zero margin of error is not possible.
#50. Q50. If the correlation coefficient between two continuous variables is $r = -0.85$, it
implies:
(A) No linear relationship
(B) A strong negative linear relationship, indicating that as one variable increases, the other tends to decrease
□ (C) A strong positive linear relationship □

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A correlation coefficient of -0.85 indicates a strong inverse linear relationship between the variables.

(D) A weak relationship with high variability

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