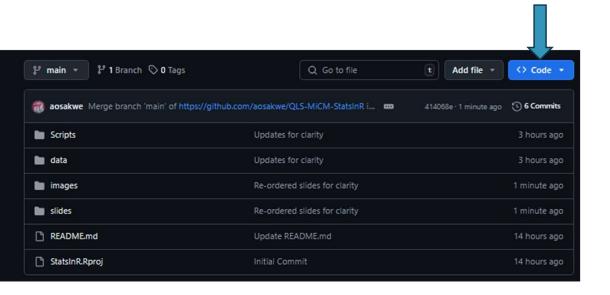
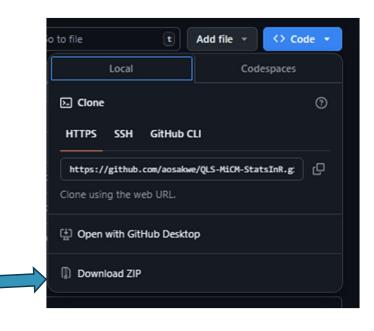


#### Workshop materials

#### https://github.com/aosakwe/QLS-MiCM-StatsInR

#### use 'git clone' command OR











# Statistical Analysis in R

Workshop Lead: Adrien Osakwe

Facilitator: Bangli Cao

February 20, 2025







QLS-MiCM mission statement: deliver quality workshops designed to help biomedical researchers develop the skills they need to succeed.



Location: 550 Sherbrooke Street, Montreal, Quebec

Contact: workshop-micm@mcgill.ca





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#### Workshop Series

Workshop	Date	Location	Registration
How to think in Code	Jan. 28 1PM-3PM	EDUC 133	Closed
Intro to Git & GitHub	Jan. 30 1PM-5PM	EDUC 133	Closed
Intro to Unix	Feb. 6 1PM-5PM	EDUC 133	Closed
Intro to Python (Part 1)	Feb. 11 1PM-5PM	EDUC 133	Closed
Intro to R (Part 1)	Feb. 13 1PM-5PM	EDUC 133	Closed
Exploring MATLAB	Feb. 18 1PM-5PM	EDUC 133	Closed
Statistics in R (Part 2)	Feb. 20 1PM-5PM	EDUC 133	Open
Data Processing in Python	Feb. 25 1PM-5PM	EDUC 133	<u>Open</u>
Intro to Machine Learning	Mar. 13 1PM-5PM	EDUC 133	TBA
Intro to R (Part 1)	TBA	EDUC 133	TBA
Intro to Python (Part 1)	TBA	EDUC 133	TBA

https://www.mcgill.ca/micm/training/workshops-series





#### Outline

- 1. Data Manipulation/Wrangling
- 2. Linear Regression Models
- 3. Logistic Regression Models
- 4. Statistical Testing
- 5. Study Design

#### **Acknowledgements**

Gerardo Martinez - McGill Alex Diaz-Papkovich - Brown Larisa Morales Soto - HMS Lisa Sullivan BUSPH





# Data Wrangling

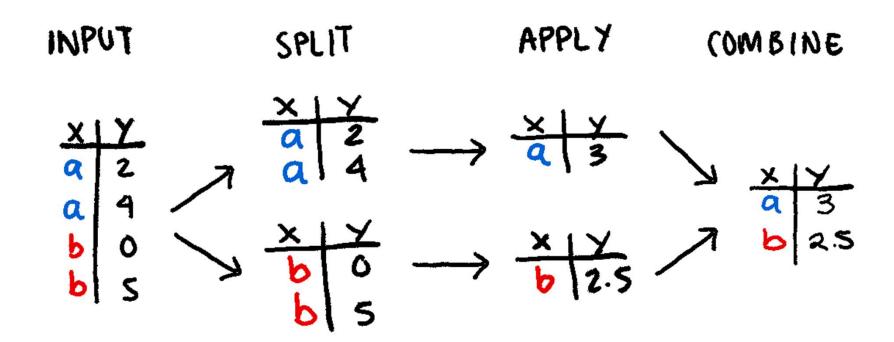
#### **Learning objectives:**

- Become familiar with the dplyr syntax
- Create pipes with the operator %>%
- Perform operations on data frames using dplyr and tidyr functions
- Implement functions from other external packages





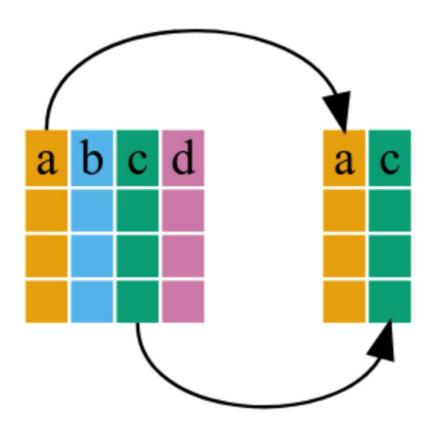
### Split-Apply-Combine problem





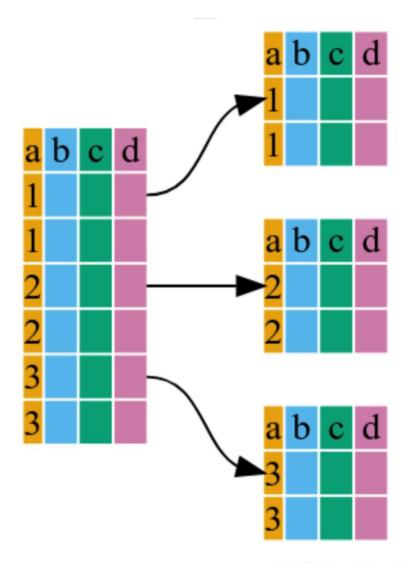


### Select





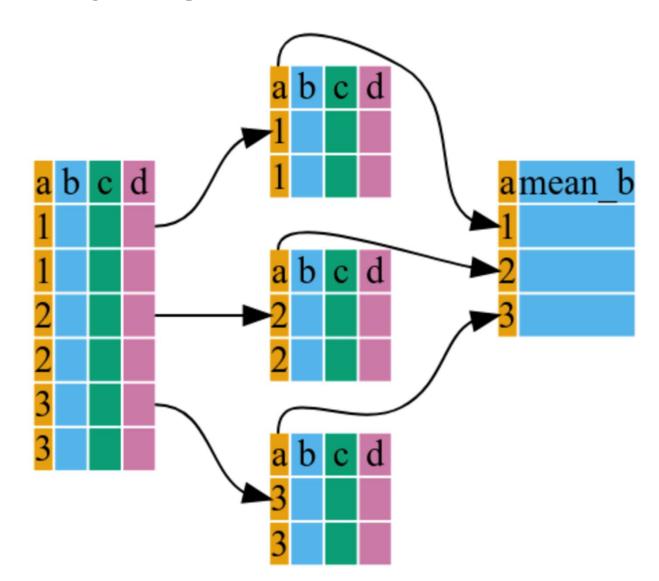
### Group by







#### Summarize







### Managing Missing Data

- Most of the time, samples having missing entries
  - Missed medical appointment
  - Error with measuring instruments etc.
  - Represented as NA in R

#### Simple Solution

Drop samples with missing observations

#### More rigorous solutions

- Fill in with average/median of dataset or condition
- Train a model to 'impute' missing entries based on other features





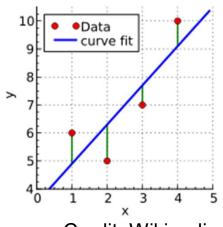
# Linear Regression

#### **Learning objectives:**

- Understand the fundamental regression algorithms
- What is a coefficient/effect size



### The Linear Model



Credit: Wikipedia

- The fundamental model for statistics & machine learning
- Follows a normal distribution

Imagine a dataset with a dependent variable  $\mathbf{y}$  and a set of descriptive features  $\mathbf{x}$ 

We want to learn what features in x are good predictors of y (what is their **effect size/coefficient**  $\beta$ )

$$y = \beta x + \varepsilon$$

$$\varepsilon \sim Normal(0, \sigma^2)$$

$$y \sim Normal(\beta x, \sigma^2)$$





# Model Assumptions

- 1. Outcome is **continuous** and a **linear combination of** predictors
- 2. Outcome is such that  $y \sim Normal(\beta x, \sigma^2)$
- 3. Predictors must not be perfectly correlated (linear combination)
- 4. For every observation *i* the error is:
  - 1. Normally distributed
  - Mean zero
  - 3. Homoskedastic (same variance as other observations)
  - **4. Independent** (not correlated)





# Model Fitting

#### **Probability**

Maximum Likelihood Estimation

#### Machine Learning

Mean Squared Error

$$y = \beta x + \epsilon$$
$$\hat{y} = \hat{\beta} x$$

$$MSE = \frac{\sum_{i=1}^{N} (\widehat{y}_i - y_i)^2}{N}$$



### Extensions

#### Interaction effects

Outcomes depend on interactions of (e.g.) age\*sex

#### **Different Types of Data**

- Multivariate regression (multiple, correlated outcomes)
- Logistic regression (Binary outcomes like healthy vs. disease)
- Multinomial/Poisson regression (Count data)
- ARIMA (Time series—data correlated over time)





# Logistic Regression

#### **Learning objectives:**

- What is a logistic function
- How LR extends the linear model
- Cross-Entropy





# Working with categories

Linear Regression could be used to predict categories (technically) BUT it has many flaws

- Formulation isn't bounded (0,1)
- Impractical for predicting multiple classes
- Mean-Squared Error is not an optimal objective function

Propose to use the linear equation with a sigmoid function





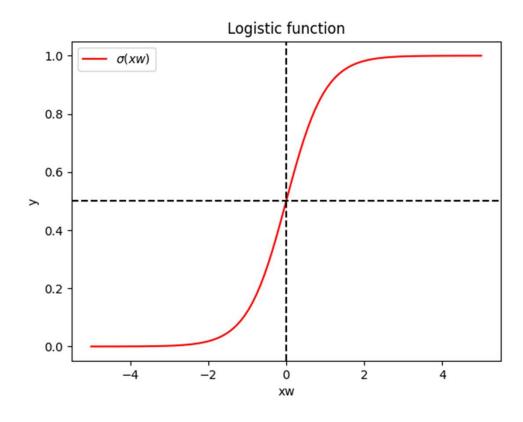
# Logistic Regression

- Sigmoid/Logistic function provides us with a bounded output
- Range is also easy to interpret as a probability

Can also use the **softmax** if working with multiple classes

$$\sigma(\mathbf{z})_i = rac{e^{eta z_i}}{\sum_{j=1}^K e^{eta z_j}}$$

$$\hat{y} = \sigma(\mathbf{x}\mathbf{w}) = \frac{1}{1 + exp(-\mathbf{x}\mathbf{w})}$$







# Cross-Entropy

$$CE(\hat{y}, y) = -ylog(\hat{y}) - (1 - y)log(1 - \hat{y})$$

Most commonly used loss function for classification

- More intuitive meaning for classification than MSE
- Faster model convergence than MSE
- Is compatible with a multi-class dataset (more than two classifications)



# Statistical Testing

#### **Learning objectives:**

- Understand the formulation an assumption of standard statistical tests
- Understand the interpretation of a p-value
- Considerations when designing an experiment





### Hypothesis Tests

#### Goal

- Determine if two sets of data are different
- Can approach this with test statistics
  - Is the difference significant or not?

#### **Many Types of Tests**

- Parametric tests makes an assumption on the data distribution
  - Z-score vs. t-test
  - ANOVA
- Non-parametric does not make an assumption on data dist.
  - Permutation test





### Hypothesis Formulation

#### **Null Hypothesis**

- Effect size is 0
- Difference between conditions is 0

#### **Alternative Hypothesis**

- Effect size is NOT 0
- Difference between conditions is NOT 0

We accept or reject the null based on the generated p-value (< 0.05)





# Understanding Significance

Statistical significance **DOES NOT** imply causality

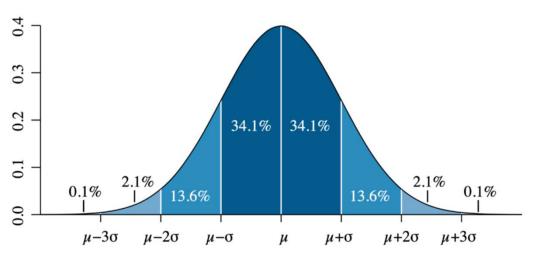
It is a measurement of the likelihood of the observed data happening by chance.

A p-value tells us the probability of seeing the observation if the null hypothesis is true.





### Z-score Test



Based on the Normal Distribution

- Credit: Wikipedia
- Generate a Z-score from our effect size estimate
- Compare to Normal(0,1) distribution
  - One-tail or Two-tail
- Used when we have a large sample size (Central Limit Theorem)

$$z = \frac{\beta - \mu}{\sigma} = \frac{\beta}{SE(\beta)}$$

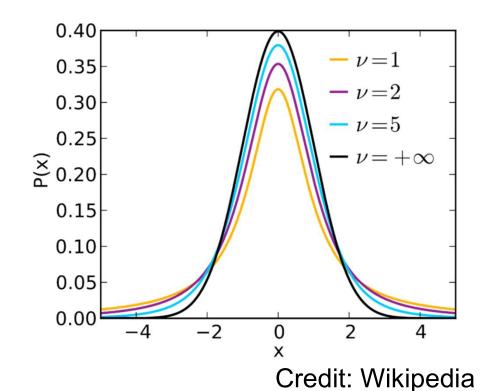




### t-test

#### Based on the student t distribution

- Defined by degrees of freedom (d.f)
  - n-1 d.f.
- More appropriate for smaller sample sizes (most cases)
- Use our sample mean  $\overline{X}$  and sample s.d  $\hat{\sigma}$  to calculate test statistic
- Value of  $\mu$  depends on your null hypothesis
  - One-Sample
  - Two-sample



$$t = \frac{\overline{X} - \mu}{s} = \frac{\overline{X} - \mu}{\hat{\sigma}/\sqrt{n}}$$



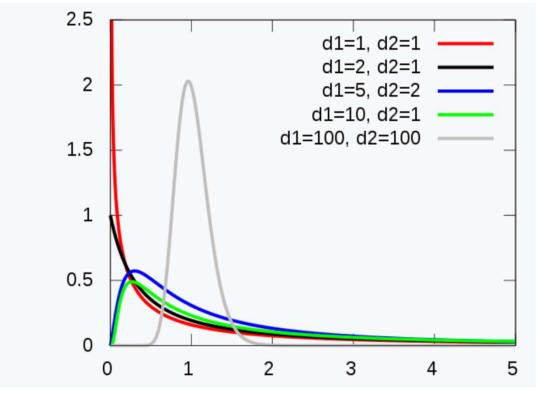
### ANOVA

#### Based on the F-distribution

- Similar to t-statistic
- Statistic is the ratio of two Chi-squareds

Null hypothesis: multiple means are all equal

Alternative: At least one group is different



Credit: Wikipedia





### ANOVA

How much variation is explained by our treatment condition?

$$SST = \sum_{i=1}^{\infty} (y_i - \overline{y})^2$$

Total sum of squares

$$SST = SS(Treatments) + SS(Residuals)$$

Take the ratio of the variance between treatments over the variance within treatments

High variance between treatments will give a low p-value





### Non-Parametric Test

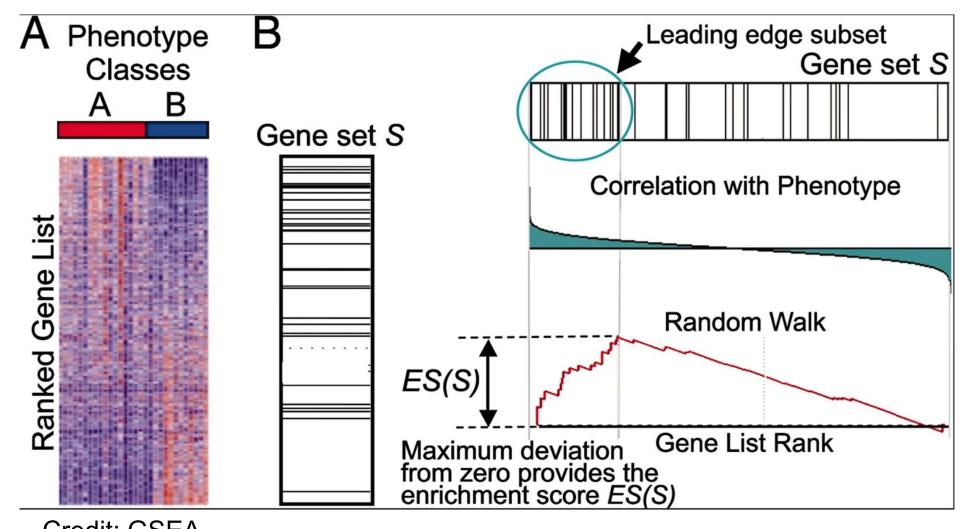
Similar to what is done for enrichment analyses

- Avoids needing to make assumptions on distributions
- Can be very slow
- 1. Shuffle labels in your data to generate estimates (random baseline)
- 2. See where your true observation lies on this baseline





### Gene Set Enrichment







# Multiple Testing

- May test multiple hypothesis (example, effect sizes in linear model)
- If null distribution is true, expect a uniform distribution of p-values
  - 5% of tests will give a significant result
- Multiple test correction methods are used to address this
  - Bonferonni Correction
  - Benjamini & Hochberg Correction
  - Found in results as FDR, p-adjusted, q-value



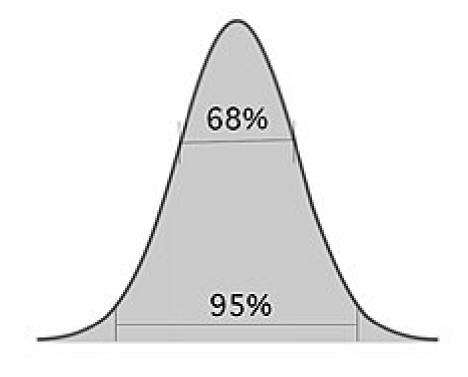


### Confidence Intervals

- We may want to attribute a range to our estimate
- Can be calculated with the margin of error

$$E = Z \frac{\sigma}{\sqrt{n}}$$

 Where Z is the z-score that defines our confidence level



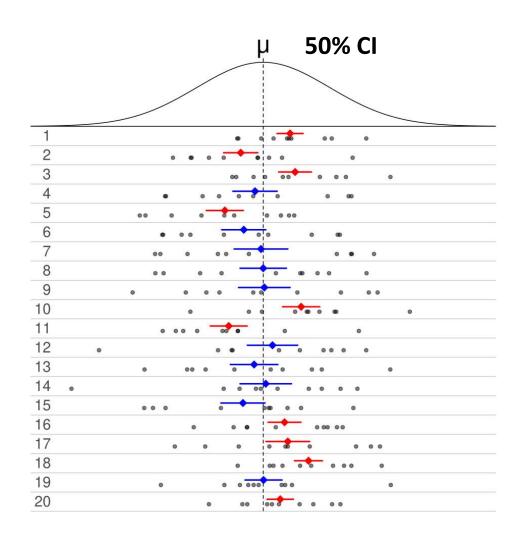


### PSA: Confidence Intervals

A 95% CI **DOES NOT mean** there is a 95% chance the true value is in that range

For multiple experiments, 95% of CIs will overlap with the true value

Consider a Bayesian formulation instead (beyond scope)



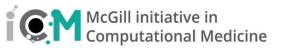




#### **Learning objectives:**

- Understand how simulations can be used to help with study design
- How to determine the required sample size

### Study Design





### Study Design

Data collection can be expensive and timeconsuming

Need to ensure that the collected data has enough **power** to avoid false negatives (type II error)

Can simulate data to help decide on the statistical test used and determine the appropriate sample size





# Determining sample size

- Statistical test results depend on multiple factors
  - Difference in Samples
  - Sample Size
- We want to make sure we have enough samples to avoid False Negatives (type II errors)
- Can leverage the assumptions used for our statistical test to determine sample size





### Confidence Intervals

In practice you will want your CI to cover a small range to be informative. Can achieve this by solving for the **margin of error E** 

$$E = Z \frac{\sigma}{\sqrt{n}}$$

$$n = \left(\frac{Z\sigma}{E}\right)^2$$

Note that we may not have a value for  $\sigma$ 

- Use estimate from past study
- Make an assumptive guess
- Run a small pilot study





# Sample Comparison

$$n_i = 2\left(\frac{Z_{1-\frac{\alpha}{2}} + Z_{1-\beta}}{ES}\right)^2$$

Effect Size (ES) = 
$$\frac{|\mu_1 - \mu_2|}{\sigma}$$

 $\alpha \rightarrow Significance Level$ 

Probability of rejecting null when it is true

 $1-\beta \rightarrow Power$ 

Probability of rejecting null when it is false





#### To summarize

- ✓ Basics for data manipulation & cleaning
- ✓ Explored basic linear modeling for regression & classification
- ✓ Explored statistical testing and intuition behind study design

#### Now you are ready to:

- Use dplyr to facilitate reproducible data manipulation
- Apply and interpret linear model results
- Interpret confidence intervals and p-values
- Explore more complex models and tests!





### Future Statistics Workshops

- Dimension Reduction
- Bayesian Inference
- Time-series Analysis
- And more...





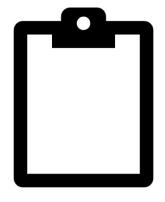
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