



This cheat sheet provides a quick reference for essential R programming commands, helping you perform data manipulation, visualization, and statistical analysis with confidence. It covers foundational topics like installing packages and understanding R's data structures, alongside advanced tasks such as building models and applying machine learning techniques.

Each section includes concise syntax and practical examples to illustrate how R commands are used in real-world scenarios. You'll find guidance on working with vectors, lists, matrices, and data frames, performing common data wrangling tasks like filtering and summarizing, and creating visualizations such as histograms, bar plots, and boxplots. The cheat sheet also highlights R's capabilities for statistical analysis with commands like mean, lm, and cor.

Designed for clarity and accessibility, this resource is ideal for data analysts, statisticians, and programmers seeking to enhance their workflows in R. Whether you're exploring data, developing algorithms, or building reproducible reports, this cheat sheet ensures you can quickly apply R's powerful tools to your projects.

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ASSIGNMENT (<-), PRINT, CLASS



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DF\$A OR DF



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Basics

Syntax for How to use

Install Package

```
install.packages("dplyr")
```

Load Package

```
library(dplyr)
```

Assignment

```
x <- 5
```

Print Output

```
print(x)
```

Literals and Data Types

```
TRUE, 125, 12.5, "Hello"
```

Extracting Numbers from Strings

```
library(readr)
data_frame <- mutate(data_frame,
column = parse_number(column))
```

Basic String Indexing

```
str_sub("Dataquest is awesome",
1, 9)
```

Explained

Installs the `dplyr` package.

Loads the `dplyr` package into the current R session.

Assigns value `5` to the variable `x`.

Prints the value of `x` to the console.

Examples of logical, integer, numeric, and character literals in R.

Uses `parse_number` to extract numeric values from string columns.

Extracts “Dataquest” as a substring by specifying start and end indices.



Data Structures

Syntax for How to use

Create Vector

```
c(1, 2, 3)
```

Create List

```
list(a=1, b="two")
```

Create Matrix

```
matrix(1:6, nrow=2)
```

Create Data Frame

```
data.frame(a=1:3, b=4:6)
```

Access Element

```
df$a | df[1, 1]
```

Loading string package

```
library(stringr)
```

Opening a JSON File

```
f <- fromJSON('filename.json')
```

Creating a List

```
new_list <- list("data scientist",
c(50000,40000), "programming
experience")
```

Explained

Combines elements into a vector.

Creates a list with named elements.

Creates a matrix with 2 rows and 3 columns.

Creates a data frame with columns `a` and `b`.

Performs a logical OR operation between a column and a specific element.

Loads the `stringr` library to work with strings in R.

Loads a JSON file into an R dataframe using the `jsonlite` package.

Defines a list containing diverse data types.



Data Manipulation

Syntax for

How to use

Filter Rows

```
filter(df, a > 2)
```

Select Columns

```
select(df, a, b)
```

Mutate Columns

```
mutate(df, c = a + b)
```

Summarize Data

```
summarize(df, avg=mean(a))
```

Arrange Rows

```
arrange(df, desc(a))
```

Importing Data

```
data <-  
read_csv("name_of_file_with_data.cs  
v")
```

Summing Values Across Rows

```
df %>% mutate(new_column_name =  
rowSums(.[1:3]))
```

Summing Values Across Columns

```
df %>% bind_rows(tibble(total =  
colSums(across(everything()))))
```

Importing CSV files

```
dataframe <-  
read_csv("name_of_the_dataset.csv")
```

Explained

Filters rows where column `a` is greater than `2`.

Selects specific columns by name.

Adds a new column `c` as sum of `a` and `b`.

Calculates mean of column `a` and returns as `avg`.

Sorts rows by column `a` in descending order.

Imports dataset into R using the `read_csv` function from `readr`.

Sums specified columns for each row and adds as a new column.

Sums specified rows for each column and adds as a new row.

Read CSV files into R using `readr`'s `read_csv()` for efficient data import.



Data Visualization

Syntax for

How to use

Creating a Basic Plot

```
data %>% ggplot()
```

Creating Subplots

```
data %>% ggplot(aes(x = variable_1,  
y = variable_2)) + geom_line() +  
facet_wrap(~variable_3)
```

Creating Bar Chart

```
data_frame %>% ggplot(aes(x =  
variable_1, y = variable_2)) +  
geom_col()
```

Plotting multiple columns

```
data %>% ggplot(aes(x =  
variable_1)) + geom_line(aes(y =  
variable_2)) + geom_line(aes(y =  
variable_3))
```

Scatterplots

```
ggplot(data = uber_trips, aes(x =  
distance, y = cost)) + geom_point()
```

Scatterplots with Labels

```
ggplot(data = df, aes(x = predictor,  
y = response)) + geom_point() +  
scale_y_continuous(labels =  
scales::comma)
```

Explained

Initialize a basic `ggplot2` chart without specifying any aesthetics.

Plots subsets of data in separate facets.

Create a bar chart using `ggplot2`, mapping variables to x and y axes.

Plots multiple columns on the same axes using `ggplot2`.

Generate scatterplots to visualize bivariate relationships in `ggplot2`.

Create scatterplots with y-axis labels formatted using commas instead of scientific notation.



Data Visualization

Syntax for

How to use

Scatterplot with Comma Labels

```
ggplot(data = df, aes(x = predictor, y = response)) +  
  scale_y_continuous(labels = scales::comma) + geom_point()
```

Scatterplot with Groups

```
ggplot(data = df, aes(x = predictor, y = response)) +  
  geom_point() + facet_wrap(~ categorical_variable, ncol = 2)
```

Scatterplot with Groups

```
ggplot(data = df, aes(x = predictor, y = response)) +  
  geom_point() + facet_wrap(~ categorical_variable, ncol = 2)
```

Vertical Bar Chart

```
ggplot(data = df, aes(x = col)) +  
  geom_bar()
```

Grouped Bar Plot

```
ggplot(data = df, aes(x = col_1,  
  fill = col_2)) + geom_bar(position = "dodge")
```

Explained

Plots a scatterplot with y-axis labels in comma format.

Creates scatterplots of response vs predictor, grouped by a categorical variable.

Creates scatterplots of response vs predictor, grouped by a categorical variable.

Creates a vertical bar chart to visualize counts of data.

Creates a grouped bar plot to compare frequency distributions of categorical variables.



Statistics & Probability

Syntax for

How to use

Mean

```
mean(x)
```

Median

```
median(x)
```

Weighted Mean

```
mean <- weighted.mean(x = distribution, w = weights)
```

Standard Deviation

```
sd(x)
```

Correlation

```
cor(x, y)
```

Linear Model

```
lm(y ~ x, data=df)
```

Types of Variables

```
# Example Variables: Age  
(Quantitative), Gender  
(Qualitative)
```

P-Value Decision Threshold

```
if (p_value < 0.05) { print('Reject  
null hypothesis') } else {  
  print('Fail to reject null  
hypothesis') }
```

Explained

Calculates the mean of vector `x`.

Calculates the median of vector `x`.

Computes the weighted mean of a numerical vector using specific weights.

Calculates the standard deviation of `x`.

Calculates correlation between `x` and `y`.

Fits a linear regression model.

Classify variables as Quantitative (numerical) or Qualitative (categorical).

Decide on hypothesis rejection using a common p-value threshold of `0.05`.





Statistics & Probability

Syntax for

How to use

Chi-Squared Distribution

```
pchisq(3.84, df = 1)
```

Chi-Squared Test

```
pchisq(q = 10, df = 5)
```

Multi-category Chi-squared Test

```
data <- table(income$sex,  
income$high_income)
```

Computing Mode in R

```
compute_mode <- function(vector)  
{counts_df <- tibble(vector) %>%  
group_by(vector) %>%  
summarise(frequency=n()) %>%  
arrange(desc(frequency));  
counts_df$vector[1]}
```

Calculate Z-score

```
z_score <- function(value, vector)  
{ (value - mean(vector)) /  
sd(vector) }
```

Chi-Squared Distribution

```
pchisq(3.84, df = 1)
```

Explained

Calculates the cumulative probability for a chi-squared distribution with specific degrees of freedom.

Calculate cumulative probability for a chi-squared statistic of 10 with 5 degrees of freedom.

Performs a chi-squared test on the given contingency table.

Defines a function to calculate the mode of a given vector using `dplyr` functions.

This calculates the Z-score for a value relative to a vector's distribution.

Calculates the cumulative probability for a chi-squared distribution with specific degrees of freedom.

Syntax for

How to use

Simulate Coin Toss

```
set.seed(1)  
coin_toss <- function() { if  
(runif(1) <= 0.5) 'HEADS' else  
'TAILS' }
```

Addition Rule for Probability

```
P(A ∪ B) = P(A) + P(B) - P(A ∩ B)
```

Independent Events

```
P(A ∩ B) = P(A) * P(B)
```

Product Rule in Experiments

```
total_outcomes <- a * b
```

Uniform Distribution

```
# Assuming all outcomes have equal  
chance  
outcomes <- c(1, 2, 3, 4, 5, 6)  
probabilities <- rep(1/6, 6)  
paste('Outcome:', outcomes,  
'Probability:', probabilities)
```

Explained

Simulates a random coin toss using R's uniform random numbers.

Formula to calculate probabilities of unions of events, adjusting for overlap in non-exclusive cases.

Probability of independent events occurs as product of individual probabilities.

Calculate the total outcomes for two independent experiments using the product rule.

Demonstrates a uniform distribution for a dice roll, where outcomes equally likely.





Statistics & Probability

Syntax for

How to use

Conditional Probability Calculation

```
P_A_given_B <- P_A_and_B / P_B
```

Conditional Probability

```
P_A_given_B <- length(intersect(A, B)) / length(B)
```

Conditional Probability Definition

```
P_A_given_B <- 1 - P_Ac_given_B
```

Independence

```
P_A_and_B <- P_A * P_B
```

Explained

Compute $P(A|B)$ given the probability of A and B, and probability of B.

Compute $P(A|B)$ using set cardinalities.

Conditional probabilities are interrelated; $P(A|B)$ and its complement $P(Ac|B)$ can be calculated mutually.

Defines independent events: joint probability equals product of individual probabilities.



Programming

Syntax for

How to use

If Statement

```
if (x > 0) print("positive")
```

For Loop

```
for (i in 1:3) print(i)
```

While Loop

```
while (x < 5) x <- x + 1
```

Syntax for functions

```
function_name <- function(input) {  
  # Code to manipulate the input  
  return(output)  
}
```

Define Function

```
f <- function(a, b) a + b
```

Apply Function

```
apply(m, 1, sum)
```

Exponentiation

```
3^5
```

Creating Dates

```
ymd('20/04/21')
```

Creating Dates from Strings

```
ymd("20/04/21")
```

Define Window Frame

```
ROWS BETWEEN 1 PRECEDING AND 1 FOLLOWING
```

Explained

Executes code if condition is true.

Iterates over a sequence.

Repeats code while the `x < 5` condition is true.

Defines a reusable function structure in R.

Defines a function with two arguments.

Applies a function over rows/columns of a matrix.

Calculates 3 raised to the power of 5.

Converts a string into a Date object using 'year-month-day'.

Converts a string to a date object using the specified format.

Defines a window frame including one row before and after the current row for computations.



 Machine Learning

Syntax for

How to use

Fitting a
Linear
Model

```
lm_fit <- lm(response ~ predictor,  
              data = df)
```

Visualize
Residuals

```
library(ggplot2)  
ggplot(data.frame(residuals =  
                  lm_fit$residuals), aes(x =  
                  residuals)) + geom_histogram()
```

Hyperparam
eter Grid
Search

```
knn_grid <- expand.grid(k = 1:20)  
knn_model <- train(tidy_price ~  
                     accommodates + bathrooms +  
                     bedrooms, data = training_data,  
                     method = "knn", trControl =  
                     train_control, preProcess =  
                     c("center", "scale"), tuneGrid =  
                     knn_grid)  
plot(knn_model)
```

Naive Bayes
Algorithm
$$P(\text{Spam} | w_1, \dots, w_n) \propto P(\text{Spam}) * \prod_i P(w_i | \text{Spam})$$

Explained

Fit a linear regression model with a response and a predictor variable.

Visualize the distribution of residuals to check the linear model's fit.

Performs grid search to optimize `k` for k-NN model and visualizes results.

Classifies messages as spam using conditional probabilities.



File I/O

Syntax for

How to use

Read CSV

```
read.csv("file.csv")
```

Write CSV

```
write.csv(df, "file.csv")
```

Read RDS

```
readRDS("file.rds")
```

Write RDS

```
saveRDS(df, "file.rds")
```

List Files

```
list.files()
```

Explained

Reads a CSV file into a data frame.

Writes a data frame to a CSV file.

Reads an RDS file into R.

Saves an object as an RDS file.

Lists files in the current directory.

