

R tutorial: Beginner to Expert

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About

Welcome to R for SAS Programmers

Course Overview

This comprehensive course is designed specifically for **SAS programmers** who want to learn R programming. We'll leverage your existing knowledge of data manipulation, statistical analysis, and programming concepts to help you become proficient in R.

Learning Objectives

By the end of this course, you will be able to:

- **Understand R fundamentals:** Master R syntax, data types, and programming concepts
- **Data manipulation:** Perform complex data transformations using `dplyr` and `base R`
- **Statistical analysis:** Apply statistical methods and create models in R
- **Data visualization:** Create compelling visualizations using `ggplot2`
- **Reporting:** Generate dynamic reports with R Markdown and Quarto
- **Bridge knowledge:** Map your SAS skills to equivalent R functions and workflows
- **Best practices:** Write clean, efficient, and reproducible R code

Course Structure

Module 1: R Fundamentals for SAS Users

- **Getting Started**
 - R and RStudio installation and setup
 - Understanding the R environment vs SAS environment
 - Package management
- **Basic R Syntax**
 - R syntax compared to SAS syntax
 - Variables and assignment operators
 - Data types and structures
 - Functions and help system

Module 2: Data Import and Export

- CSV, Excel and text files (`readr`, `readxl` packages)
- SAS dataset import (`haven` package) and export xpt files
- other formats (JSON, XML)

Module 3: Data Manipulation - `dplyr`

- **Core `dplyr` Functions**
 - `select()` vs `KEEP/DROP` statements
 - `filter()` vs `WHERE` clause
 - `mutate()` vs `assignment` statements
 - `summarise()` vs `PROC MEANS`
 - `group_by()` vs `BY` statement
- **Advanced Data Manipulation**
 - Joins equivalent to `PROC SQL` joins
 - Reshaping data (`tidyr` vs `PROC TRANSPOSE`)
 - String manipulation (`stringr` vs SAS string functions)

Module 4: Creation of ADSL dataset

- **ADSL and ADVS Dataset Creation**
 - Creating analysis variables
 - Handling missing data and derivations
 - creating ADSL xpt dataset

Module 5: Statistical Analysis

- **Descriptive Statistics**
 - Summary statistics (equivalent to `PROC UNIVARIATE`)
 - Frequency tables (equivalent to `PROC FREQ`)
 - Cross-tabulations and chi-square tests
 - creation of tables like DM and AE outputs
- **Statistical Modeling basics**
 - Linear regression (equivalent to `PROC REG`)
 - Logistic regression (equivalent to `PROC LOGISTIC`)

- ANOVA (equivalent to `PROC ANOVA`)
- Mixed models and advanced techniques

Module 6: Package Development and testing

- **Creating R Packages**
 - Package structure and essential files
 - Documenting functions with `roxygen2`
 - Building and installing packages
- **Testing with `testthat`**
 - Writing unit tests for R functions
 - Running tests and interpreting results

Module 7: Data Visualization

- **Base R Graphics**
 - Basic plots and customization
 - Comparison with SAS/GRAPH
- **ggplot2 - Grammar of Graphics**
 - Understanding the layered approach
 - Creating publication-ready plots
 - Advanced visualization techniques

Module 8: Reporting and Documentation

- **R Markdown and Quarto**
 - Creating dynamic reports (equivalent to ODS output)
 - Integrating code, results, and narrative
 - Output formats: HTML, PDF, Word
- **Reproducible Research**
 - Project organization
 - Version control with Git
 - Best practices for code documentation

SAS to R Translation Guide

| SAS Concept | R Equivalent | Package |
|----------------|---|------------------|
| DATA step | <code>dplyr::mutate()</code> | dplyr |
| PROC SQL | dplyr verbs | dplyr |
| PROC MEANS | <code>dplyr::summarise()</code> | dplyr |
| PROC FREQ | <code>table()</code> , <code>xtabs()</code> | base R |
| PROC REG | <code>lm()</code> | base R |
| PROC LOGISTIC | <code>glm()</code> | base R |
| PROC TRANSPOSE | <code>tidyr::pivot_*()</code> | tidyr |
| ODS OUTPUT | R Markdown/Quarto | rmarkdown/quarto |

Prerequisites

- **SAS Experience:** Familiarity with SAS programming, data steps, and procedures
- **Statistical Knowledge:** Basic understanding of statistical concepts
- **Programming Basics:** Understanding of programming logic and data structures

Course Format

- **Interactive Learning:** Hands-on exercises with real datasets
- **Comparative Examples:** Side-by-side SAS and R code comparisons
- **Practical Projects:** Real-world scenarios mimicking typical SAS workflows
- **Reference Materials:** Quick reference guides and cheat sheets

Getting Started

Required Software

1. **R** (version 4.3+): Download from [CRAN](#)
2. **RStudio**: Download from [Posit](#)
3. **Essential Packages:** We'll install these as needed

```
install.packages(c("tidyverse", "haven", "readxl", "rmarkdown"))
```

Tip: If you don't have sample SDTM/ADaM data yet, the chapters generate **small synthetic data** as a fallback so everything runs end-to-end. ## contact For questions or feedback, reach out to **r2sas2025@gmail.com**

Part I

datatype & structure

1 R Fundamentals for SAS Users

Basic R Syntax - A Comprehensive Guide

2 Introduction

This guide is designed for SAS users transitioning to R. We'll cover fundamental R concepts by comparing them to familiar SAS constructs, providing comprehensive coverage of each topic.

2.1 Prerequisites

3 1. R Syntax Compared to SAS Syntax

3.1 Basic Code Structure and Comments

Understanding how R code is structured compared to SAS is fundamental to making the transition.

:: {.panel-tabset}

3.2 SAS

```
/* Multi-line comments in SAS
   can span multiple lines */
* Single line comment with asterisk;

/* Statements must end with semicolons */
DATA mydata;
    SET olddata;
    new_var = old_var * 2;
RUN;

/* Procedures need RUN statements */
PROC PRINT DATA=mydata;
RUN;

/* Case insensitive - these are all the same */
data test;
DATA test;
Data test;
```

3.3 R

```

# Single line comments use hash symbol
# There are no multi-line comments in base R
# (though RStudio supports Ctrl+Shift+C for multiple lines)

# Semicolons are optional but allowed
new_var <- old_var * 2 # No semicolon needed
new_var <- old_var * 2; # Semicolon allowed

# No RUN statements required - code executes immediately
print(mydata)

# R is CASE SENSITIVE - these are different variables
data <- 1
Data <- 2
DATA <- 3

```

...

3.3.1 Key Syntax Differences

| Aspect | SAS | R | Notes |
|-----------------------------|--------------------|------------------------------|------------------------------------|
| Comment | /* */ or *; | # | R comments run to end of line only |
| Statement terminator | ; required | ; optional | Most R code omits semicolons |
| Case sensitivity | Not case-sensitive | Case-sensitive | Major difference! |
| Assignment | = | <- or = | <- is preferred in R |
| Block execution | RUN; required | Automatic | Code executes line by line |
| Line continuation | Automatic | Automatic with open brackets | |

3.3.2 Demonstrating Case Sensitivity

```

# In R, these are THREE DIFFERENT variables
Variable <- 10
variable <- 20

```



```
VARIABLE <- 30
```

```
print(Variable) # Returns 10
```

```
[1] 10
```

```
print(variable) # Returns 20
```

```
[1] 20
```

```
print(VARIABLE) # Returns 30
```

```
[1] 30
```

```
# This would cause confusion in SAS but works in R
```

```
mydata <- data.frame(x = 1:5)
```

```
MyData <- data.frame(x = 6:10)
```

```
MYDATA <- data.frame(x = 11:15)
```

```
# Each is a separate object
```

```
nrow(mydata) # 5
```

```
[1] 5
```

```
nrow(MyData) # 5
```

```
[1] 5
```

```
nrow(MYDATA) # 5
```

```
[1] 5
```

3.4 Procedural vs Functional Programming Paradigm

SAS and R have fundamentally different approaches to data manipulation and analysis.

3.4.1 SAS: Step-by-Step Procedures

```
/* SAS uses distinct procedures with explicit steps */

/* Step 1: Sort data */
PROC SORT DATA=mydata OUT=sorted_data;
  BY age;
RUN;

/* Step 2: Calculate statistics */
PROC MEANS DATA=sorted_data MEAN STD;
  VAR income;
  CLASS gender;
  OUTPUT OUT=summary_stats MEAN=avg_income STD=sd_income;
RUN;

/* Step 3: Print results */
PROC PRINT DATA=summary_stats;
RUN;
```

3.4.2 R: Multiple Functional Approaches

```
# Approach 1: Base R - function chaining
sorted_data <- mydata[order(mydata$age), ]
summary_stats <- aggregate(income ~ gender, data = sorted_data,
                           FUN = function(x) c(mean = mean(x), sd = sd(x)))
print(summary_stats)

# Approach 2: tidyverse - pipe operator (most similar to thinking in steps)
summary_stats <- mydata %>%
  arrange(age) %>%
  group_by(gender) %>%
  summarise(
    avg_income = mean(income, na.rm = TRUE),
    sd_income = sd(income, na.rm = TRUE)
  )
print(summary_stats)

# Approach 3: data.table - high performance
```

```
library(data.table)
dt <- as.data.table(mydata)
summary_stats <- dt[order(age)][, .(avg_income = mean(income),
                                   sd_income = sd(income)),
                             by = gender]
print(summary_stats)
```

3.4.3 Understanding the Pipe Operator (%>% and |>)

The pipe operator makes R code read more like SAS procedures:

```
# Create sample data
employees <- data.frame(
  name = c("John", "Jane", "Bob", "Alice", "Charlie", "Diana"),
  department = c("Sales", "IT", "IT", "Sales", "HR", "Sales"),
  salary = c(50000, 75000, 68000, 52000, 48000, 55000),
  years = c(2, 5, 3, 1, 4, 3)
)

# Without pipes (nested functions - hard to read)
result1 <- head(arrange(filter(employees, department == "Sales"), desc(salary)), 3)

# With pipes (reads left to right, top to bottom)
result2 <- employees %>%
  filter(department == "Sales") %>%
  arrange(desc(salary)) %>%
  head(3)

print(result2)
```

| | name | department | salary | years |
|---|-------|------------|--------|-------|
| 1 | Diana | Sales | 55000 | 3 |
| 2 | Alice | Sales | 52000 | 1 |
| 3 | John | Sales | 50000 | 2 |

```
# Native pipe |> (R 4.1+) - similar but slightly different
result3 <- employees |>
  filter(department == "Sales") |>
  arrange(desc(salary)) |>
  head(3)
```

```
print(result3)
```

| | name | department | salary | years |
|---|-------|------------|--------|-------|
| 1 | Diana | Sales | 55000 | 3 |
| 2 | Alice | Sales | 52000 | 1 |
| 3 | John | Sales | 50000 | 2 |

3.5 Execution and Evaluation Models

Understanding how R evaluates code differently from SAS helps avoid common pitfalls.

3.5.1 Sequential vs Expression-Based Execution

SAS: Executes in clearly defined DATA and PROC steps with explicit boundaries

```
/* SAS processes entire DATA step before moving to next step */
DATA step1;
    SET input_data;
    x = 10;
RUN; /* Everything above completes before proceeding */

DATA step2;
    SET step1;
    y = x + 5; /* x is available because step1 completed */
RUN;
```

R: Expression-based evaluation, code executes immediately

```
# R executes line by line
x <- 10          # Executes immediately
y <- x + 5       # Can use x immediately
z <- x + y       # Can use both x and y

print(paste("x:", x, "y:", y, "z:", z))
```

```
[1] "x: 10 y: 15 z: 25"
```

```
# Functions execute when called
my_calculation <- function() {
  a <- 100
  b <- 200
  return(a + b)
}

# Function is defined but not executed yet
result <- my_calculation() # Now it executes
print(result)
```

```
[1] 300
```

3.5.2 Lazy Evaluation in R

R uses “lazy evaluation” - function arguments are only evaluated when actually used:

```
# Demonstrating lazy evaluation
my_function <- function(x, y, z) {
  # If we return early, unused arguments never get evaluated
  if (x > 0) {
    return(x * 2)
  }
  # y and z are never evaluated if x > 0
  return(y + z)
}

# This works even though second and third arguments would error
result <- my_function(5, stop("Error in y!"), stop("Error in z!"))
print(result) # Returns 10, no errors
```

```
[1] 10
```

```
# But this would cause an error
# result <- my_function(-1, stop("Error in y!"), 10) # ERROR!
```

3.5.3 Vectorized Operations

R operates on entire vectors at once (unlike SAS’s row-by-row processing):

```
# Create vectors
vector1 <- c(1, 2, 3, 4, 5)
vector2 <- c(10, 20, 30, 40, 50)

# Vectorized operation - all elements at once
result <- vector1 + vector2
print(result) # 11 22 33 44 55
```

```
[1] 11 22 33 44 55
```

```
# Comparison to SAS approach
# In SAS, you'd typically process row by row:
# DATA result;
#     SET input;
#     new_value = value1 + value2;
# RUN;

# In R, you can also do row-wise operations on data frames
df <- data.frame(value1 = vector1, value2 = vector2)
df$result <- df$value1 + df$value2
print(df)
```

| | value1 | value2 | result |
|---|--------|--------|--------|
| 1 | 1 | 10 | 11 |
| 2 | 2 | 20 | 22 |
| 3 | 3 | 30 | 33 |
| 4 | 4 | 40 | 44 |
| 5 | 5 | 50 | 55 |

3.5.4 Recycling Rules

R automatically “recycles” shorter vectors to match longer ones:

```
# Vector recycling
short_vec <- c(1, 2)
long_vec <- c(10, 20, 30, 40, 50, 60)

# short_vec is recycled: 1, 2, 1, 2, 1, 2
result <- short_vec + long_vec
print(result) # 11 22 31 42 51 62
```

```
[1] 11 22 31 42 51 62
```

```
# Warning when lengths don't divide evenly
vec1 <- c(1, 2, 3)
vec2 <- c(10, 20, 30, 40, 50)
# result <- vec1 + vec2 # Would give warning

# Practical use: add constant to all elements
values <- c(100, 200, 300, 400)
values_plus_10 <- values + 10 # 10 is recycled
print(values_plus_10)
```

```
[1] 110 210 310 410
```

4 2. Variables and Assignment Operators

4.1 Creating and Assigning Variables

Understanding variable assignment is crucial for writing effective R code.

4.1.1 Three Assignment Operators

```
# Left assignment with <- (RECOMMENDED)
x <- 10
patient_age <- 45
department_name <- "Cardiology"

# Left assignment with = (works but not preferred for variables)
y = 20
# Use = for function arguments: mean(x, na.rm = TRUE)

# Right assignment with -> (rarely used, but valid)
30 -> z

# All three created variables
print(paste("x =", x, "| y =", y, "| z =", z))

# Why <- is preferred:
# 1. Clearly distinguishes assignment from function arguments
# 2. Consistent with R conventions and style guides
# 3. Can be read as "gets" or "assign to"

# Example of clarity with <-
my_data <- read.csv("file.csv", header = TRUE) # Clear distinction
# vs
# my_data = read.csv("file.csv", header = TRUE) # Less clear
```


4.1.2 Chaining Assignments

```
# Multiple assignments in one line
a <- b <- c <- 100
print(paste("a =", a, "| b =", b, "| c =", c))

# Assignment with computation
result <- (x <- 5) + (y <- 10)
print(paste("x =", x, "| y =", y, "| result =", result))

# Practical example: assign and use
data_subset <- subset(employees, salary > (threshold <- 50000))
print(paste("Threshold used:", threshold))
print(data_subset)
```

4.1.3 Variable Naming Rules and Conventions

```
# VALID variable names
valid_name <- 1
valid.name <- 2          # Dots allowed (unlike most languages)
validName <- 3           # camelCase
valid_name_123 <- 4      # Numbers allowed (but not at start)
.hidden_var <- 5         # Starting with dot (hidden from ls())

# Common naming conventions
snake_case_variable <- "preferred in R"
camelCaseVariable <- "common in some R code"
dot.separated.name <- "traditional R style"
PascalCase <- "typically for functions/classes"

# INVALID variable names (will cause errors)
# 123invalid <- 5        # ERROR: Cannot start with number
# invalid-name <- 6      # ERROR: Hyphens not allowed (minus sign)
# _invalid <- 7          # ERROR: Cannot start with underscore
# my variable <- 8       # ERROR: Spaces not allowed

# Reserved words cannot be used as variable names
reserved_words <- c("if", "else", "repeat", "while", "function",
                   "for", "in", "next", "break", "TRUE", "FALSE",
```

```

"NULL", "Inf", "NaN", "NA")
print(reserved_words)

```

```

[1] "if"      "else"    "repeat"  "while"   "function" "for"
[7] "in"      "next"    "break"   "TRUE"    "FALSE"    "NULL"
[13] "Inf"     "NaN"     "NA"

```

```

# if <- 10 # ERROR: 'if' is reserved
# BUT you can use them with backticks (not recommended)
`if` <- 10
print(`if`)

```

```

[1] 10

```

4.1.4 Checking and Removing Variables

```

# Create some variables
var1 <- 100
var2 <- 200
var3 <- 300

# List all variables in environment
current_vars <- ls()
print(current_vars)

```

```

[1] "camelCaseVariable" "df" "dot.separated.name"
[4] "employees"        "if" "long_vec"
[7] "my_calculation"    "my_function" "mydata"
[10] "MyData"            "MYDATA" "PascalCase"
[13] "required_packages" "reserved_words" "result"
[16] "result1"           "result2" "result3"
[19] "short_vec"         "snake_case_variable" "valid_name"
[22] "valid_name_123"    "valid.name" "validName"
[25] "values"            "values_plus_10" "var1"
[28] "var2"              "var3" "variable"
[31] "Variable"          "VARIABLE" "vec1"
[34] "vec2"              "vector1" "vector2"
[37] "x"                 "y" "z"

```

```
# Check if variable exists
exists("var1") # TRUE
```

```
[1] TRUE
```

```
exists("var999") # FALSE
```

```
[1] FALSE
```

```
# Remove specific variables
rm(var3)
exists("var3") # FALSE
```

```
[1] FALSE
```

```
# Remove multiple variables
rm(var1, var2)

# Remove all variables (use with caution!)
# rm(list = ls())
```

4.2 Variable Scope and Environments

Understanding scope is crucial for writing functions and avoiding bugs.

4.2.1 Local vs Global Scope

```
# Global variable (available everywhere)
global_var <- 100

# Function with local variables
my_function <- function() {
  # Local variable (only exists inside function)
  local_var <- 200

  # Can read global variables
  print(paste("Inside function, global_var:", global_var))
}
```

```

    print(paste("Inside function, local_var:", local_var))

    # Return something
    return(local_var * 2)
}

# Call function
result <- my_function()

```

```

[1] "Inside function, global_var: 100"
[1] "Inside function, local_var: 200"

```

```

print(paste("Function returned:", result))

```

```

[1] "Function returned: 400"

```

```

# local_var doesn't exist outside function
print(paste("Outside function, global_var:", global_var))

```

```

[1] "Outside function, global_var: 100"

```

```

# print(local_var) # ERROR: object 'local_var' not found

```

4.2.2 Modifying Global Variables from Functions

```

# Global variable
counter <- 0

# Function that modifies local copy (default behavior)
increment_local <- function() {
  counter <- counter + 1 # Creates local 'counter'
  print(paste("Inside increment_local:", counter))
}

increment_local()

```

```

[1] "Inside increment_local: 1"

```

```
print(paste("Global counter after increment_local:", counter)) # Still 0!
```

```
[1] "Global counter after increment_local: 0"
```

```
# Function that modifies global variable (use <<- or assign)
increment_global <- function() {
  counter <<- counter + 1 # Modifies global 'counter'
  print(paste("Inside increment_global:", counter))
}

increment_global()
```

```
[1] "Inside increment_global: 1"
```

```
print(paste("Global counter after increment_global:", counter)) # Now 1!
```

```
[1] "Global counter after increment_global: 1"
```

```
increment_global()
```

```
[1] "Inside increment_global: 2"
```

```
print(paste("Global counter after second call:", counter)) # Now 2!
```

```
[1] "Global counter after second call: 2"
```

4.2.3 Understanding «- Super Assignment

super assignment operator <<- allows modification of variables in parent environments.

```
# <<- searches parent environments until it finds the variable
outer_function <- function() {
  x <- 10 # Local to outer_function

  inner_function <- function() {
    x <<- x + 5 # Modifies x in outer_function's environment
    print(paste("Inside inner_function, x:", x))
  }
}
```

```

    }

    print(paste("Before inner_function, x:", x))
    inner_function()
    print(paste("After inner_function, x:", x))
}

outer_function()

```

```

[1] "Before inner_function, x: 10"
[1] "Inside inner_function, x: 15"
[1] "After inner_function, x: 15"

```

```

# Use cases for <<-:
# 1. Counters and state in function factories
# 2. Caching/memoization
# 3. Building interactive applications
# WARNING: Use sparingly, can make code hard to understand

```

4.2.4 Function Factories and Closures

```

# Creating a function that returns a function
create_counter <- function() {
  count <- 0 # This variable persists across calls

  function() {
    count <<- count + 1
    return(count)
  }
}

# Create two independent counters
counter1 <- create_counter()
counter2 <- create_counter()

# Each maintains its own state
print(counter1()) # 1

```

```

[1] 1

```

```
print(counter1()) # 2
```

```
[1] 2
```

```
print(counter1()) # 3
```

```
[1] 3
```

```
print(counter2()) # 1
```

```
[1] 1
```

```
print(counter2()) # 2
```

```
[1] 2
```

```
# Practical example: create functions with specific parameters
create_multiplier <- function(factor) {
  function(x) {
    return(x * factor)
  }
}

times_2 <- create_multiplier(2)
times_10 <- create_multiplier(10)

print(times_2(5)) # 10
```

```
[1] 10
```

```
print(times_10(5)) # 50
```

```
[1] 50
```

4.2.5 Working with Environments Explicitly

the default environment is the global environment, but you can use the `new.env()` function to create and manipulate environments directly.

```
# Create new environment
my_env <- new.env()

# Assign variables to environment
my_env$var1 <- 100
my_env$var2 <- 200

# Access environment variables
print(my_env$var1)
```

```
[1] 100
```

```
# List variables in environment
ls(my_env)
```

```
[1] "var1" "var2"
```

```
# Environments are passed by reference (not copied)
modify_env <- function(env) {
  env$var1 <- 999
  env$new_var <- 777
}

modify_env(my_env)
print(my_env$var1)      # 999 (modified!)
```

```
[1] 999
```

```
print(my_env$new_var)  # 777 (added!)
```

```
[1] 777
```

```
# This is different from regular objects which are copied
```

4.3 Copy-on-Modify Behavior

R's copy-on-modify system is important for understanding performance and memory usage.

4.3.1 Default Copy-on-Modify

```
# Create original vector
original <- c(1, 2, 3, 4, 5)

# Assign to new variable (no copy yet!)
copy_var <- original

# Modify copy_var (NOW a copy is made)
copy_var[1] <- 999

# Original is unchanged
print(original)  # 1 2 3 4 5
```

```
[1] 1 2 3 4 5
```

```
print(copy_var)  # 999 2 3 4 5
```

```
[1] 999  2  3  4  5
```

```
# Same with data frames
original_df <- data.frame(x = 1:5, y = 6:10)
copy_df <- original_df
copy_df$x[1] <- 999

print(original_df$x)  # 1 2 3 4 5 (unchanged)
```

```
[1] 1 2 3 4 5
```

```
print(copy_df$x)      # 999 2 3 4 5 (modified)
```

```
[1] 999  2  3  4  5
```

4.3.2 Implications for Function Arguments

```
# Functions receive copies (so modifications don't affect original)
modify_vector <- function(vec) {
  vec[1] <- 999
  print(paste("Inside function:", paste(vec, collapse = " ")))
  return(vec)
}

my_vector <- c(1, 2, 3, 4, 5)
result <- modify_vector(my_vector)
```

```
[1] "Inside function: 999 2 3 4 5"
```

```
print(paste("Original vector:", paste(my_vector, collapse = " "))) # Unchanged
```

```
[1] "Original vector: 1 2 3 4 5"
```

```
print(paste("Returned vector:", paste(result, collapse = " "))) # Modified
```

```
[1] "Returned vector: 999 2 3 4 5"
```

4.3.3 Using Environments for True Reference Behavior

When you need pass-by-reference behavior (like SAS datasets that get modified):

```
# Environment approach (pass by reference)
create_data_env <- function() {
  env <- new.env()
  env$data <- data.frame(id = 1:5, value = rnorm(5))
  return(env)
}

modify_data <- function(data_env, new_value) {
  data_env$data$value <- data_env$data$value + new_value
}

# Create data environment
my_data_env <- create_data_env()
print("Original:")
```

```
[1] "Original:"
```

```
print(my_data_env$data)
```

| | id | value |
|---|----|------------|
| 1 | 1 | 0.7563500 |
| 2 | 2 | -1.1486746 |
| 3 | 3 | 0.5043772 |
| 4 | 4 | 1.1743032 |
| 5 | 5 | -1.2969403 |

```
# Modify (changes original!)  
modify_data(my_data_env, 100)  
print("After modification:")
```

```
[1] "After modification:"
```

```
print(my_data_env$data) # Modified!
```

| | id | value |
|---|----|-----------|
| 1 | 1 | 100.75635 |
| 2 | 2 | 98.85133 |
| 3 | 3 | 100.50438 |
| 4 | 4 | 101.17430 |
| 5 | 5 | 98.70306 |

```
# This is similar to how SAS datasets work
```

5 3. Data Types and Structures

5.1 Atomic Data Types

R has six atomic (basic) types. Understanding these is fundamental to data manipulation.

5.1.1 Numeric Types

```
# Double (default numeric type in R)
num_double <- 42.5
class(num_double)
```

```
[1] "numeric"
```

```
typeof(num_double)
```

```
[1] "double"
```

```
# Integer (requires L suffix)
num_integer <- 42L
class(num_integer)
```

```
[1] "integer"
```

```
typeof(num_integer)
```

```
[1] "integer"
```

```
# Automatic conversion
x <- 10      # Double
y <- 10L     # Integer
z <- x + y   # Result is double
typeof(z)
```

```
[1] "double"
```

```
# Checking numeric types  
is.numeric(num_double) # TRUE
```

```
[1] TRUE
```

```
is.numeric(num_integer) # TRUE
```

```
[1] TRUE
```

```
is.integer(num_integer) # TRUE
```

```
[1] TRUE
```

```
is.integer(num_double) # FALSE
```

```
[1] FALSE
```

```
is.double(num_double) # TRUE
```

```
[1] TRUE
```

```
# Coercion  
as.integer(42.7) # 42 (truncates)
```

```
[1] 42
```

```
as.numeric("123") # 123
```

```
[1] 123
```

```
as.numeric("abc") # NA with warning
```

```
[1] NA
```

5.1.2 Character (String) Types

```

# Creating character variables
char1 <- "Hello"
char2 <- 'World' # Single or double quotes work
char3 <- "Can include 'quotes' inside"

# Multi-line strings
multi_line <- "This is line 1
This is line 2
This is line 3"

# String operations
paste("Hello", "World") # "Hello World"

```

```
[1] "Hello World"
```

```
paste0("No", "Space") # "NoSpace"
```

```
[1] "NoSpace"
```

```
paste("A", "B", "C", sep = "-") # "A-B-C"
```

```
[1] "A-B-C"
```

```
sprintf("Patient %d has BMI %.2f", 101, 24.5) # Formatted string
```

```
[1] "Patient 101 has BMI 24.50"
```

```

# String manipulation
toupper("hello") # "HELLO"

```

```
[1] "HELLO"
```

```
tolower("HELLO") # "hello"
```

```
[1] "hello"
```

```
nchar("Hello") # 5 (length)
```

```
[1] 5
```

```
substr("Hello World", 1, 5) # "Hello"
```

```
[1] "Hello"
```

```
gsub("o", "0", "Hello World") # "Hello W0rld" (replace)
```

```
[1] "Hello W0rld"
```

```
# Check and convert  
is.character(char1) # TRUE
```

```
[1] TRUE
```

```
as.character(123) # "123"
```

```
[1] "123"
```

5.1.3 Logical (Boolean) Types

```
# Creating logical values  
flag1 <- TRUE  
flag2 <- FALSE  
flag3 <- T      # Shorthand (but TRUE preferred)  
flag4 <- F      # Shorthand (but FALSE preferred)
```

```
# Logical operations  
TRUE & FALSE # AND: FALSE
```

```
[1] FALSE
```

```
TRUE | FALSE # OR: TRUE
```

```
[1] TRUE
```

```
!TRUE # NOT: FALSE
```

```
[1] FALSE
```

```
TRUE && FALSE # Short-circuit AND: FALSE (only evaluates what's needed)
```

```
[1] FALSE
```

```
TRUE || FALSE # Short-circuit OR: TRUE
```

```
[1] TRUE
```

```
# Comparisons return logical  
5 > 3 # TRUE
```

```
[1] TRUE
```

```
5 == 5 # TRUE
```

```
[1] TRUE
```

```
5 != 3 # TRUE
```

```
[1] TRUE
```

```
"a" %in% c("a", "b", "c") # TRUE
```

```
[1] TRUE
```

```
# Logical arithmetic (TRUE = 1, FALSE = 0)  
sum(c(TRUE, FALSE, TRUE, TRUE)) # 3
```

```
[1] 3
```



```
mean(c(TRUE, FALSE, TRUE, TRUE))    # 0.75
```

```
[1] 0.75
```

```
# Conversion  
as.logical(0)      # FALSE
```

```
[1] FALSE
```

```
as.logical(1)      # TRUE
```

```
[1] TRUE
```

```
as.logical("TRUE") # TRUE
```

```
[1] TRUE
```

```
as.logical("yes")  # NA
```

```
[1] NA
```

5.1.4 Special Values

```
# NA - Missing value (like . in SAS)  
na_value <- NA  
is.na(na_value) # TRUE
```

```
[1] TRUE
```

```
# Different types of NA  
na_real <- NA_real_      # Numeric NA  
na_int <- NA_integer_    # Integer NA  
na_char <- NA_character_ # Character NA  
na_logical <- NA        # Logical NA  
  
# NULL - Absence of value (different from NA)  
null_value <- NULL  
is.null(null_value) # TRUE
```

```
[1] TRUE
```

```
length(NULL)      # 0
```

```
[1] 0
```

```
length(NA)        # 1
```

```
[1] 1
```

```
# Inf - Infinity  
inf_value <- Inf  
1 / 0      # Inf
```

```
[1] Inf
```

```
-1 / 0      # -Inf
```

```
[1] -Inf
```

```
is.infinite(Inf)  # TRUE
```

```
[1] TRUE
```

```
is.finite(Inf)    # FALSE
```

```
[1] FALSE
```

```
# NaN - Not a Number  
nan_value <- NaN  
0 / 0      # NaN
```

```
[1] NaN
```

```
is.nan(nan_value) # TRUE
```

```
[1] TRUE
```

```
# Testing for special values
test_values <- c(1, NA, NaN, Inf, -Inf)
is.na(test_values)      # TRUE for NA and NaN
```

```
[1] FALSE TRUE TRUE FALSE FALSE
```

```
is.nan(test_values)     # TRUE only for NaN
```

```
[1] FALSE FALSE TRUE FALSE FALSE
```

```
is.infinite(test_values) # TRUE for Inf and -Inf
```

```
[1] FALSE FALSE FALSE TRUE TRUE
```

```
is.finite(test_values)  # TRUE only for 1
```

```
[1] TRUE FALSE FALSE FALSE FALSE
```

5.1.5 SAS to R Type Comparison

```
# Create comparison table
comparison <- data.frame(
  SAS_Type = c("Numeric", "Character", "Logical (1/0)", ". (missing)", "Date (numeric)"),
  R_Type = c("numeric/integer", "character", "logical (TRUE/FALSE)", "NA", "Date class"),
  Example = c("42.5 or 42L", '"text"', "TRUE/FALSE", "NA", 'as.Date("2025-12-18")'),
  stringsAsFactors = FALSE
)

print(comparison)
```

| | SAS_Type | R_Type | Example |
|---|----------------|----------------------|-----------------------|
| 1 | Numeric | numeric/integer | 42.5 or 42L |
| 2 | Character | character | "text" |
| 3 | Logical (1/0) | logical (TRUE/FALSE) | TRUE/FALSE |
| 4 | . (missing) | NA | NA |
| 5 | Date (numeric) | Date class | as.Date("2025-12-18") |

```
# Date example
sas_date <- 19709 # SAS date value for 2025-12-18
r_date <- as.Date("2025-12-18")
print(r_date)
```

```
[1] "2025-12-18"
```

```
print(class(r_date))
```

```
[1] "Date"
```

```
# Convert between systems
r_date_from_sas <- as.Date(sas_date, origin = "1960-01-01")
print(r_date_from_sas)
```

```
[1] "2013-12-17"
```

5.2 Vector Data Structures

Vectors are the fundamental data structure in R - even single values are vectors of length 1.

5.2.1 Creating Vectors

```
# Combine function c()
numeric_vec <- c(1, 2, 3, 4, 5)
char_vec <- c("apple", "banana", "cherry", "date")
logical_vec <- c(TRUE, FALSE, TRUE, TRUE, FALSE)

# Sequences
seq1 <- 1:10 # Simple sequence
seq2 <- 10:1 # Descending
seq3 <- seq(0, 100, by = 10) # With step
seq4 <- seq(0, 1, length.out = 11) # Specific length
seq5 <- rep(5, times = 3) # Repeat value
seq6 <- rep(c(1, 2), times = 3) # Repeat vector
seq7 <- rep(c(1, 2), each = 3) # Repeat each element

print(seq1)
```

```
[1] 1 2 3 4 5 6 7 8 9 10
```

```
print(seq3)
```

```
[1] 0 10 20 30 40 50 60 70 80 90 100
```

```
print(seq6) # 1 2 1 2 1 2
```

```
[1] 1 2 1 2 1 2
```

```
print(seq7) # 1 1 1 2 2 2
```

```
[1] 1 1 1 2 2 2
```

5.2.2 Vector Properties and Operations

```
# Vector properties  
vec <- c(10, 20, 30, 40, 50)  
  
length(vec) # 5
```

```
[1] 5
```

```
class(vec) # "numeric"
```

```
[1] "numeric"
```

```
typeof(vec) # "double"
```

```
[1] "double"
```

```
str(vec) # Structure
```

```
num [1:5] 10 20 30 40 50
```

```
# Vectorized operations (applied element-wise)
vec + 10          # 20 30 40 50 60
```

```
[1] 20 30 40 50 60
```

```
vec * 2          # 20 40 60 80 100
```

```
[1] 20 40 60 80 100
```

```
vec^2           # 100 400 900 1600 2500
```

```
[1] 100 400 900 1600 2500
```

```
sqrt(vec)       # Square root of each element
```

```
[1] 3.162278 4.472136 5.477226 6.324555 7.071068
```

```
log(vec)        # Natural log of each element
```

```
[1] 2.302585 2.995732 3.401197 3.688879 3.912023
```

```
# Operations between vectors (element-wise)
vec1 <- c(1, 2, 3, 4, 5)
vec2 <- c(10, 20, 30, 40, 50)
vec1 + vec2      # 11 22 33 44 55
```

```
[1] 11 22 33 44 55
```

```
vec1 * vec2     # 10 40 90 160 250
```

```
[1] 10 40 90 160 250
```

5.2.3 Named Vectors

```
# Create named vector
scores <- c(Math = 95, English = 88, Science = 92, History = 85)
print(scores)
```

```
Math English Science History
95      88      92      85
```

```
# Access by name
scores["Math"]
```

```
Math
95
```

```
scores[c("Math", "Science")]
```

```
Math Science
95      92
```

```
# Get names
names(scores)
```

```
[1] "Math"      "English" "Science" "History"
```

```
# Add/modify names
grades <- c(95, 88, 92, 85)
names(grades) <- c("Math", "English", "Science", "History")
print(grades)
```

```
Math English Science History
95      88      92      85
```

```
# Named vectors are useful for lookup tables
month_days <- c(Jan = 31, Feb = 28, Mar = 31, Apr = 30, May = 31, Jun = 30,
                Jul = 31, Aug = 31, Sep = 30, Oct = 31, Nov = 30, Dec = 31)
month_days["Feb"]
```

```
Feb
28
```

5.2.4 Vector Indexing and Subsetting

which function is very useful for logical indexing.

```
# Create vector for examples
vec <- c(10, 20, 30, 40, 50, 60, 70, 80, 90, 100)

# Positive indexing (1-based, not 0-based like Python!)
vec[1]           # First element: 10
```

```
[1] 10
```

```
vec[5]           # Fifth element: 50
```

```
[1] 50
```

```
vec[c(1, 3, 5)]  # Multiple elements: 10 30 50
```

```
[1] 10 30 50
```

```
vec[1:5]         # Range: 10 20 30 40 50
```

```
[1] 10 20 30 40 50
```

```
# Negative indexing (exclusion)
vec[-1]          # All except first
```

```
[1] 20 30 40 50 60 70 80 90 100
```

```
vec[-c(1, 2)]    # All except first two
```

```
[1] 30 40 50 60 70 80 90 100
```

```
vec[-(1:5)]      # All except first five
```

```
[1] 60 70 80 90 100
```



```
# Logical indexing (very powerful!)
vec[vec > 50]      # Elements greater than 50
```

```
[1] 60 70 80 90 100
```

```
vec[vec %% 2 == 0] # Even elements
```

```
[1] 10 20 30 40 50 60 70 80 90 100
```

```
vec[vec >= 30 & vec <= 70] # Between 30 and 70
```

```
[1] 30 40 50 60 70
```

```
# Which function (returns indices)
which(vec > 50)      # Indices where condition is TRUE
```

```
[1] 6 7 8 9 10
```

```
vec[which.max(vec)] # Maximum value
```

```
[1] 100
```

```
vec[which.min(vec)] # Minimum value
```

```
[1] 10
```

5.2.5 Vector Type Coercion

```
# Vectors must be homogeneous (all same type)
# R coerces to most flexible type: logical < integer < double < character

mixed1 <- c(1, 2, "three")      # All become character
print(mixed1)
```

```
[1] "1"      "2"      "three"
```

```
class(mixed1)
```

```
[1] "character"
```

```
mixed2 <- c(TRUE, FALSE, 1, 2) # All become numeric  
print(mixed2)
```

```
[1] 1 0 1 2
```

```
class(mixed2)
```

```
[1] "numeric"
```

```
mixed3 <- c(TRUE, FALSE, "yes") # All become character  
print(mixed3)
```

```
[1] "TRUE" "FALSE" "yes"
```

```
class(mixed3)
```

```
[1] "character"
```

```
# Explicit coercion  
char_nums <- c("1", "2", "3", "4")  
as.numeric(char_nums) # Convert to numeric
```

```
[1] 1 2 3 4
```

```
logical_vals <- c(1, 0, 1, 1, 0)  
as.logical(logical_vals) # Convert to logical
```

```
[1] TRUE FALSE TRUE TRUE FALSE
```

5.3 Lists - Flexible Containers

Lists can contain elements of different types and structures, including other lists.

5.3.1 Creating Lists

```
# Basic list creation
my_list <- list(
  numbers = c(1, 2, 3, 4, 5),
  text = "Hello World",
  flag = TRUE,
  matrix_data = matrix(1:6, nrow = 2),
  nested_list = list(a = 10, b = 20)
)

print(my_list)
```

```
$numbers
[1] 1 2 3 4 5
```

```
$text
[1] "Hello World"
```

```
$flag
[1] TRUE
```

```
$matrix_data
      [,1] [,2] [,3]
[1,]    1    3    5
[2,]    2    4    6
```

```
$nested_list
$nested_list$a
[1] 10
```

```
$nested_list$b
[1] 20
```

```
str(my_list) # Structure is very helpful for lists
```

```
List of 5
 $ numbers      : num [1:5] 1 2 3 4 5
 $ text         : chr "Hello World"
 $ flag         : logi TRUE
```

```
$ matrix_data: int [1:2, 1:3] 1 2 3 4 5 6
$ nested_list:List of 2
..$ a: num 10
..$ b: num 20
```

5.3.2 Accessing List Elements

```
# Three ways to access list elements
```

```
# 1. Using $ (by name)
my_list$numbers
```

```
[1] 1 2 3 4 5
```

```
my_list$text
```

```
[1] "Hello World"
```

```
# 2. Using [[ ]] (by name or index) - extracts element
my_list[["numbers"]]
```

```
[1] 1 2 3 4 5
```

```
my_list[[1]]
```

```
[1] 1 2 3 4 5
```

```
my_list[["nested_list"]]$a
```

```
[1] 10
```

```
# 3. Using [ ] - returns a list
my_list["numbers"] # Returns list with one element
```

```
$numbers
[1] 1 2 3 4 5
```

```
my_list[1]          # Same
```

```
$numbers  
[1] 1 2 3 4 5
```

```
class(my_list[[1]]) # "numeric"
```

```
[1] "numeric"
```

```
class(my_list[1])   # "list"
```

```
[1] "list"
```

```
# Accessing nested elements  
my_list$nested_list$a
```

```
[1] 10
```

```
my_list[["nested_list"]][["a"]]
```

```
[1] 10
```

```
my_list[[5]][[1]]
```

```
[1] 10
```

5.3.3 Modifying Lists

```
# Add new elements  
my_list$new_element <- c(100, 200, 300)  
my_list[["another_element"]] <- "New value"  
  
# Modify existing elements  
my_list$text <- "Modified text"
```

```
# Remove elements
my_list$new_element <- NULL

# Append lists
list1 <- list(a = 1, b = 2)
list2 <- list(c = 3, d = 4)
combined <- c(list1, list2)
print(combined)
```

```
$a
[1] 1
```

```
$b
[1] 2
```

```
$c
[1] 3
```

```
$d
[1] 4
```

```
# Get list names
names(my_list)
```

```
[1] "numbers"      "text"          "flag"          "matrix_data"
[5] "nested_list"  "another_element"
```

```
# Unnamed lists
unnamed_list <- list(10, "text", TRUE)
print(unnamed_list)
```

```
[[1]]
[1] 10
```

```
[[2]]
[1] "text"
```

```
[[3]]
[1] TRUE
```

```
unnamed_list[[2]] # Access by index only
```

```
[1] "text"
```

5.3.4 Practical List Applications

```
# Storing analysis results
perform_analysis <- function(data) {
  results <- list(
    summary_stats = summary(data),
    mean_value = mean(data, na.rm = TRUE),
    sd_value = sd(data, na.rm = TRUE),
    n_missing = sum(is.na(data)),
    n_total = length(data)
  )
  return(results)
}

sample_data <- c(12, 15, 18, NA, 22, 25, 28, 31)
analysis_results <- perform_analysis(sample_data)
print(analysis_results)
```

```
$summary_stats
  Min. 1st Qu.  Median    Mean 3rd Qu.    Max.   NA's
12.00  16.50   22.00   21.57  26.50   31.00     1
```

```
$mean_value
[1] 21.57143
```

```
$sd_value
[1] 6.948792
```

```
$n_missing
[1] 1
```

```
$n_total
[1] 8
```

```
# Accessing results
print(paste("Mean:", analysis_results$mean_value))
```

```
[1] "Mean: 21.5714285714286"
```

```
print(paste("Missing values:", analysis_results$n_missing))
```

```
[1] "Missing values: 1"
```

5.4 Data Frames - R's Version of SAS Datasets

Data frames are the primary structure for tabular data in R.

5.4.1 Creating Data Frames

```
# Method 1: From vectors
employees_df <- data.frame(
  employee_id = 1:6,
  name = c("John", "Jane", "Bob", "Alice", "Charlie", "Diana"),
  department = c("Sales", "IT", "IT", "Sales", "HR", "Sales"),
  salary = c(50000, 75000, 68000, 52000, 48000, 55000),
  years_employed = c(2, 5, 3, 1, 4, 3),
  full_time = c(TRUE, TRUE, FALSE, TRUE, TRUE, FALSE),
  stringsAsFactors = FALSE # Keep strings as character (important!)
)

print(employees_df)
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|---------|------------|--------|----------------|-----------|
| 1 | 1 | John | Sales | 50000 | 2 | TRUE |
| 2 | 2 | Jane | IT | 75000 | 5 | TRUE |
| 3 | 3 | Bob | IT | 68000 | 3 | FALSE |
| 4 | 4 | Alice | Sales | 52000 | 1 | TRUE |
| 5 | 5 | Charlie | HR | 48000 | 4 | TRUE |
| 6 | 6 | Diana | Sales | 55000 | 3 | FALSE |


```
# Method 2: From lists
emp_list <- list(
  employee_id = 1:3,
  name = c("John", "Jane", "Bob"),
  salary = c(50000, 75000, 68000)
)
df_from_list <- as.data.frame(emp_list)
print(df_from_list)
```

```
   employee_id name salary
1            1  John 50000
2            2  Jane 75000
3            3   Bob 68000
```

```
# Method 3: Read from file (most common)
# employees_df <- read.csv("employees.csv")
# employees_df <- read.table("employees.txt", header = TRUE)
```

5.4.2 Data Frame Properties

```
# Dimensions
nrow(employees_df)      # Number of rows
```

```
[1] 6
```

```
ncol(employees_df)      # Number of columns
```

```
[1] 6
```

```
dim(employees_df)       # Both: rows, columns
```

```
[1] 6 6
```

```
# Names
names(employees_df)     # Column names
```

```
[1] "employee_id"      "name"              "department"        "salary"
[5] "years_employed"  "full_time"
```

```
colnames(employees_df) # Same
```

```
[1] "employee_id"      "name"              "department"        "salary"
[5] "years_employed"  "full_time"
```

```
rownames(employees_df) # Row names (usually just numbers)
```

```
[1] "1" "2" "3" "4" "5" "6"
```

```
# Structure and summary
```

```
str(employees_df) # Structure (like PROC CONTENTS)
```

```
'data.frame':  6 obs. of  6 variables:
 $ employee_id   : int  1 2 3 4 5 6
 $ name          : chr  "John" "Jane" "Bob" "Alice" ...
 $ department    : chr  "Sales" "IT" "IT" "Sales" ...
 $ salary        : num  50000 75000 68000 52000 48000 55000
 $ years_employed: num  2 5 3 1 4 3
 $ full_time     : logi  TRUE TRUE FALSE TRUE TRUE FALSE
```

```
summary(employees_df) # Summary statistics
```

| employee_id | name | department | salary |
|----------------|------------------|------------------|---------------|
| Min. :1.00 | Length:6 | Length:6 | Min. :48000 |
| 1st Qu.:2.25 | Class :character | Class :character | 1st Qu.:50500 |
| Median :3.50 | Mode :character | Mode :character | Median :53500 |
| Mean :3.50 | | | Mean :58000 |
| 3rd Qu.:4.75 | | | 3rd Qu.:64750 |
| Max. :6.00 | | | Max. :75000 |
| years_employed | full_time | | |
| Min. :1.00 | Mode :logical | | |
| 1st Qu.:2.25 | FALSE:2 | | |
| Median :3.00 | TRUE :4 | | |
| Mean :3.00 | | | |
| 3rd Qu.:3.75 | | | |
| Max. :5.00 | | | |

```
head(employees_df, 3) # First 3 rows
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|------|------------|--------|----------------|-----------|
| 1 | 1 | John | Sales | 50000 | 2 | TRUE |
| 2 | 2 | Jane | IT | 75000 | 5 | TRUE |
| 3 | 3 | Bob | IT | 68000 | 3 | FALSE |

```
tail(employees_df, 2) # Last 2 rows
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|---------|------------|--------|----------------|-----------|
| 5 | 5 | Charlie | HR | 48000 | 4 | TRUE |
| 6 | 6 | Diana | Sales | 55000 | 3 | FALSE |

```
glimpse(employees_df) # tidyverse version (if loaded)
```

```
Rows: 6
Columns: 6
$ employee_id    <int> 1, 2, 3, 4, 5, 6
$ name           <chr> "John", "Jane", "Bob", "Alice", "Charlie", "Diana"
$ department     <chr> "Sales", "IT", "IT", "Sales", "HR", "Sales"
$ salary         <dbl> 50000, 75000, 68000, 52000, 48000, 55000
$ years_employed <dbl> 2, 5, 3, 1, 4, 3
$ full_time      <lgl> TRUE, TRUE, FALSE, TRUE, TRUE, FALSE
```

5.4.3 Accessing Data Frame Elements

```
# Access columns
employees_df$name # By name with $
```

```
[1] "John"    "Jane"    "Bob"     "Alice"   "Charlie" "Diana"
```

```
employees_df[["name"]] # By name with [[]]
```

```
[1] "John"    "Jane"    "Bob"     "Alice"   "Charlie" "Diana"
```

```
employees_df[, "name"] # By name with [,]
```

```
[1] "John" "Jane" "Bob" "Alice" "Charlie" "Diana"
```

```
employees_df[, 2] # By position
```

```
[1] "John" "Jane" "Bob" "Alice" "Charlie" "Diana"
```

```
# Access rows  
employees_df[1, ] # First row
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|------|------------|--------|----------------|-----------|
| 1 | 1 | John | Sales | 50000 | 2 | TRUE |

```
employees_df[1:3, ] # First three rows
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|------|------------|--------|----------------|-----------|
| 1 | 1 | John | Sales | 50000 | 2 | TRUE |
| 2 | 2 | Jane | IT | 75000 | 5 | TRUE |
| 3 | 3 | Bob | IT | 68000 | 3 | FALSE |

```
# Access specific elements  
employees_df[1, 2] # Row 1, column 2
```

```
[1] "John"
```

```
employees_df[1, "name"] # Row 1, column "name"
```

```
[1] "John"
```

```
employees_df$name[1] # Element 1 of column "name"
```

```
[1] "John"
```

```
# Multiple columns
employees_df[, c("name", "salary")]
```

| | name | salary |
|---|---------|--------|
| 1 | John | 50000 |
| 2 | Jane | 75000 |
| 3 | Bob | 68000 |
| 4 | Alice | 52000 |
| 5 | Charlie | 48000 |
| 6 | Diana | 55000 |

```
employees_df[, c(2, 4)]
```

| | name | salary |
|---|---------|--------|
| 1 | John | 50000 |
| 2 | Jane | 75000 |
| 3 | Bob | 68000 |
| 4 | Alice | 52000 |
| 5 | Charlie | 48000 |
| 6 | Diana | 55000 |

```
# Subset with conditions
employees_df[employees_df$salary > 50000, ]
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|-------|------------|--------|----------------|-----------|
| 2 | 2 | Jane | IT | 75000 | 5 | TRUE |
| 3 | 3 | Bob | IT | 68000 | 3 | FALSE |
| 4 | 4 | Alice | Sales | 52000 | 1 | TRUE |
| 6 | 6 | Diana | Sales | 55000 | 3 | FALSE |

```
employees_df[employees_df$department == "Sales", ]
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|-------|------------|--------|----------------|-----------|
| 1 | 1 | John | Sales | 50000 | 2 | TRUE |
| 4 | 4 | Alice | Sales | 52000 | 1 | TRUE |
| 6 | 6 | Diana | Sales | 55000 | 3 | FALSE |

```
employees_df[employees_df$salary > 50000 & employees_df$full_time, ]
```

| | employee_id | name | department | salary | years_employed | full_time |
|---|-------------|-------|------------|--------|----------------|-----------|
| 2 | 2 | Jane | IT | 75000 | 5 | TRUE |
| 4 | 4 | Alice | Sales | 52000 | 1 | TRUE |

5.4.4 Modifying Data Frames

```
# Add new column
employees_df$bonus <- employees_df$salary * 0.10

# Modify existing column
employees_df$salary <- employees_df$salary * 1.05 # 5% raise

# Add calculated column
employees_df$total_comp <- employees_df$salary + employees_df$bonus

# Delete column
employees_df$bonus <- NULL

# Rename columns
names(employees_df)[names(employees_df) == "name"] <- "employee_name"

# Add rows
new_employee <- data.frame(
  employee_id = 7,
  employee_name = "Eve",
  department = "IT",
  salary = 70000,
  years_employed = 2,
  full_time = TRUE,
  total_comp = 77000
)
employees_df <- rbind(employees_df, new_employee)

# Reset column name back
names(employees_df)[names(employees_df) == "employee_name"] <- "name"
```

5.4.5 SAS DATA Step vs R Data Frame Operations

: :: {.panel-tabset}

5.5 SAS

```
/* SAS DATA step */
DATA analysis;
  SET employees;
  WHERE salary > 50000;

  /* Create variables */
  bonus = salary * 0.10;
  total_comp = salary + bonus;

  /* Conditional logic */
  IF department = 'Sales' THEN sales_flag = 1;
  ELSE sales_flag = 0;

  /* Keep specific variables */
  KEEP employee_id name salary bonus total_comp;
RUN;
```

5.6 R (Base)

```
# Base R approach
analysis <- employees[employees$salary > 50000, ]
analysis$bonus <- analysis$salary * 0.10
analysis$total_comp <- analysis$salary + analysis$bonus
analysis$sales_flag <- ifelse(analysis$department == "Sales", 1, 0)
analysis <- analysis[, c("employee_id", "name", "salary", "bonus", "total_comp")]
```

5.7 R (tidyverse)

```
# tidyverse approach (more readable)
analysis <- employees_df %>%
  filter(salary > 50000) %>%
  mutate(
    bonus = salary * 0.10,
    total_comp = salary + bonus,
    sales_flag = if_else(department == "Sales", 1, 0)
  ) %>%
  select(employee_id, name, salary, bonus, total_comp)

print(analysis)
```

...

5.8 Matrices and Arrays

Matrices (2D) and arrays (N-dimensional) are homogeneous data structures.

5.8.1 Creating and Using Matrices

```
# Create matrix
mat <- matrix(1:12, nrow = 3, ncol = 4)
print(mat)
```

```
      [,1] [,2] [,3] [,4]
[1,]    1    4    7   10
[2,]    2    5    8   11
[3,]    3    6    9   12
```

```
# Matrix with specific fill pattern
mat_byrow <- matrix(1:12, nrow = 3, ncol = 4, byrow = TRUE)
print(mat_byrow)
```

```
      [,1] [,2] [,3] [,4]
[1,]    1    2    3    4
[2,]    5    6    7    8
[3,]    9   10   11   12
```



```
# Named dimensions
mat_named <- matrix(1:12, nrow = 3, ncol = 4,
                    dimnames = list(c("Row1", "Row2", "Row3"),
                                    c("Col1", "Col2", "Col3", "Col4")))
print(mat_named)
```

```
      Col1 Col2 Col3 Col4
Row1     1     4     7    10
Row2     2     5     8    11
Row3     3     6     9    12
```

```
# Matrix properties
dim(mat)          # Dimensions
```

```
[1] 3 4
```

```
nrow(mat)         # Number of rows
```

```
[1] 3
```

```
ncol(mat)         # Number of columns
```

```
[1] 4
```

```
length(mat)       # Total elements
```

```
[1] 12
```

```
# Matrix indexing
mat[1, 2]         # Element at row 1, column 2
```

```
[1] 4
```

```
mat[1, ]         # First row
```

```
[1] 1 4 7 10
```

```
mat[, 2]          # Second column
```

```
[1] 4 5 6
```

```
mat[1: 2, 2: 3]   # Submatrix
```

```
      [,1] [,2]  
[1,]    4    7  
[2,]    5    8
```

5.8.2 Matrix Operations

```
# Create matrices for operations  
A <- matrix(c(1, 2, 3, 4), nrow = 2)  
B <- matrix(c(5, 6, 7, 8), nrow = 2)  
  
# Element-wise operations  
A + B          # Element-wise addition
```

```
      [,1] [,2]  
[1,]    6   10  
[2,]    8   12
```

```
A * B          # Element-wise multiplication
```

```
      [,1] [,2]  
[1,]    5   21  
[2,]   12   32
```

```
A / B          # Element-wise division
```

```
      [,1]      [,2]  
[1,] 0.2000000 0.4285714  
[2,] 0.3333333 0.5000000
```

```
# Matrix multiplication
A %*% B      # Matrix product
```

```
      [,1] [,2]
[1,]   23   31
[2,]   34   46
```

```
# Transpose
t(A)
```

```
      [,1] [,2]
[1,]     1     2
[2,]     3     4
```

```
# Inverse (if square and invertible)
solve(A)
```

```
      [,1] [,2]
[1,]   -2  1.5
[2,]     1 -0.5
```

```
# Determinant
det(A)
```

```
[1] -2
```

```
# Eigenvalues and eigenvectors
eigen(A)
```

```
eigen() decomposition
$values
[1]  5.3722813 -0.3722813
```

```
$vectors
      [,1]      [,2]
[1,] -0.5657675 -0.9093767
[2,] -0.8245648  0.4159736
```

5.8.3 Arrays (Multi-dimensional)

```
# Create 3D array
arr <- array(1:24, dim = c(3, 4, 2))
print(arr)
```

, , 1

| | [,1] | [,2] | [,3] | [,4] |
|------|------|------|------|------|
| [1,] | 1 | 4 | 7 | 10 |
| [2,] | 2 | 5 | 8 | 11 |
| [3,] | 3 | 6 | 9 | 12 |

, , 2

| | [,1] | [,2] | [,3] | [,4] |
|------|------|------|------|------|
| [1,] | 13 | 16 | 19 | 22 |
| [2,] | 14 | 17 | 20 | 23 |
| [3,] | 15 | 18 | 21 | 24 |

```
# Access elements
arr[1, 2, 1]      # Single element
```

[1] 4

```
arr[, , 1]        # First "slice"
```

| | [,1] | [,2] | [,3] | [,4] |
|------|------|------|------|------|
| [1,] | 1 | 4 | 7 | 10 |
| [2,] | 2 | 5 | 8 | 11 |
| [3,] | 3 | 6 | 9 | 12 |

```
arr[1, , ]        # First row across all slices
```

| | [,1] | [,2] |
|------|------|------|
| [1,] | 1 | 13 |
| [2,] | 4 | 16 |
| [3,] | 7 | 19 |
| [4,] | 10 | 22 |

```
# Array properties
dim(arr)
```

```
[1] 3 4 2
```

```
length(arr)
```

```
[1] 24
```

5.9 Tibbles - Modern Data Frames

Tibbles are enhanced data frames from the tidyverse with better printing and behavior.

5.9.1 Creating Tibbles

```
library(tibble)

# Create tibble
employees_tbl <- tibble(
  employee_id = 1:6,
  name = c("John", "Jane", "Bob", "Alice", "Charlie", "Diana"),
  department = c("Sales", "IT", "IT", "Sales", "HR", "Sales"),
  salary = c(50000, 75000, 68000, 52000, 48000, 55000),
  start_date = as.Date(c("2023-01-15", "2020-06-01", "2022-03-10",
                        "2024-02-20", "2021-09-05", "2022-11-12"))
)

print(employees_tbl) # Better printing than data.frame
```

```
# A tibble: 6 x 5
  employee_id name      department salary start_date
    <int> <chr>      <chr>      <dbl> <date>
1         1 John      Sales      50000 2023-01-15
2         2 Jane      IT         75000 2020-06-01
3         3 Bob       IT         68000 2022-03-10
4         4 Alice     Sales      52000 2024-02-20
5         5 Charlie   HR         48000 2021-09-05
6         6 Diana     Sales      55000 2022-11-12
```

```
# Tribble: row-wise tibble creation
employees_tribble <- tribble(
  ~employee_id, ~name,      ~department, ~salary,
  1,            "John",     "Sales",    50000,
  2,            "Jane",     "IT",      75000,
  3,            "Bob",      "IT",      68000
)
print(employees_tribble)
```

```
# A tibble: 3 x 4
  employee_id name  department salary
      <dbl> <chr> <chr>      <dbl>
1           1 John  Sales      50000
2           2 Jane  IT        75000
3           3 Bob   IT        68000
```

5.9.2 Tibble vs Data Frame Differences

```
# Difference 1: Printing
# Data frames print everything, tibbles show first 10 rows
df_large <- data.frame(x = 1:100, y = 101:200)
tbl_large <- tibble(x = 1:100, y = 101:200)
# print(df_large)    # Would print all 100 rows
print(tbl_large)     # Prints first 10 rows nicely
```

```
# A tibble: 100 x 2
      x     y
  <int> <int>
1     1  101
2     2  102
3     3  103
4     4  104
5     5  105
6     6  106
7     7  107
8     8  108
9     9  109
10    10  110
# i 90 more rows
```

```
# Difference 2: Subsetting
df <- data.frame(x = 1:3, y = 4:6)
tbl <- tibble(x = 1:3, y = 4:6)

df[, "x"]          # Returns vector
```

```
[1] 1 2 3
```

```
tbl[, "x"]          # Returns tibble
```

```
# A tibble: 3 x 1
      x
  <int>
1     1
2     2
3     3
```

```
# Difference 3: Character vectors
df_char <- data.frame(name = c("John", "Jane"))
tbl_char <- tibble(name = c("John", "Jane"))

str(df_char)        # Might convert to factor (depends on R version)
```

```
'data.frame':  2 obs. of  1 variable:
 $ name: chr  "John" "Jane"
```

```
str(tbl_char)       # Always character
```

```
tibble [2 x 1] (S3: tbl_df/tbl/data.frame)
 $ name: chr [1:2] "John" "Jane"
```

```
# Difference 4: Column names
# Tibbles allow non-syntactic names
tbl_special <- tibble(
  `Column with spaces` = 1:3,
  `2024` = 4:6
)
print(tbl_special)
```

```
# A tibble: 3 x 2
  `Column with spaces` `2024`
    <int>    <int>
1         1         4
2         2         5
3         3         6
```

5.9.3 Converting Between Data Frames and Tibbles

```
# Convert data frame to tibble
df <- data.frame(x = 1:3, y = 4:6)
tbl <- as_tibble(df)

# Convert tibble to data frame
tbl <- tibble(x = 1:3, y = 4:6)
df <- as.data.frame(tbl)

# Check type
is.data.frame(tbl) # TRUE (tibbles are also data frames)
```

```
[1] TRUE
```

```
is_tibble(df) # FALSE
```

```
[1] FALSE
```

```
is_tibble(tbl) # TRUE
```

```
[1] TRUE
```

5.10 Data. table - High Performance Option

Data.table is optimized for speed and memory efficiency with large datasets.

5.10.1 Creating Data.tables


```
library(data.table)

# Create data. table
employees_dt <- data.table(
  employee_id = 1:1000,
  department = sample(c("Sales", "IT", "HR", "Finance"), 1000, replace = TRUE),
  salary = round(rnorm(1000, mean = 60000, sd = 15000), 0),
  years = sample(1:20, 1000, replace = TRUE)
)

print(head(employees_dt))
```

| | employee_id | department | salary | years |
|----|-------------|------------|--------|-------|
| | <int> | <char> | <num> | <int> |
| 1: | 1 | HR | 51842 | 17 |
| 2: | 2 | HR | 66407 | 13 |
| 3: | 3 | IT | 71294 | 11 |
| 4: | 4 | HR | 81347 | 20 |
| 5: | 5 | Finance | 45480 | 6 |
| 6: | 6 | Sales | 74822 | 20 |

```
# Convert from data frame
df <- data.frame(x = 1:3, y = 4:6)
dt <- as.data.table(df)
```

5.10.2 Data.table Syntax: DT[i, j, by]

The data.table syntax follows the pattern: DT[where, select, group by]

```
# i: Filter rows (WHERE clause)
employees_dt[salary > 70000]
```

| | employee_id | department | salary | years |
|----|-------------|------------|--------|-------|
| | <int> | <char> | <num> | <int> |
| 1: | 3 | IT | 71294 | 11 |
| 2: | 4 | HR | 81347 | 20 |
| 3: | 6 | Sales | 74822 | 20 |
| 4: | 10 | Finance | 70873 | 13 |
| 5: | 12 | HR | 80826 | 1 |

```

---
234:      977      IT  80616      8
235:      978      HR  70166     14
236:      989  Finance 105046     14
237:      996      IT  93737      2
238:      997     Sales  75090      1

```

```
employees_dt[department == "IT"]
```

```

      employee_id department salary years
      <int>      <char>  <num> <int>
1:           3         IT  71294     11
2:           9         IT  51208     16
3:          16         IT  56581     15
4:          19         IT  35140      7
5:          20         IT  64713      9
---
241:         977         IT  80616      8
242:         994         IT  54681      9
243:         996         IT  93737      2
244:         999         IT  49916     16
245:        1000         IT  68992      9

```

```

# j: Select/compute columns (SELECT clause)
employees_dt[, .(employee_id, salary)]

```

```

      employee_id salary
      <int>  <num>
1:           1  51842
2:           2  66407
3:           3  71294
4:           4  81347
5:           5  45480
---
996:         996  93737
997:         997  75090
998:         998  56055
999:         999  49916
1000:        1000  68992

```

```
employees_dt[, mean_salary := mean(salary)]
```

```
# by: Group by (GROUP BY clause)
```

```
employees_dt[, .(avg_salary = mean(salary)), by = department]
```

| | department | avg_salary |
|----|------------|------------|
| | <char> | <num> |
| 1: | HR | 59172.37 |
| 2: | IT | 60116.66 |
| 3: | Finance | 56665.81 |
| 4: | Sales | 58846.69 |

```
employees_dt[, .(count = .N), by = department] # . N is row count
```

| | department | count |
|----|------------|-------|
| | <char> | <int> |
| 1: | HR | 251 |
| 2: | IT | 245 |
| 3: | Finance | 249 |
| 4: | Sales | 255 |

```
# Combining i, j, by
```

```
employees_dt[salary > 50000,  
              .(avg_salary = mean(salary), count = .N),  
              by = department]
```

| | department | avg_salary | count |
|----|------------|------------|-------|
| | <char> | <num> | <int> |
| 1: | HR | 66055.12 | 184 |
| 2: | IT | 66951.79 | 184 |
| 3: | Sales | 65162.32 | 186 |
| 4: | Finance | 64984.46 | 164 |

```
# Multiple group by
```

```
employees_dt[, .(avg_salary = mean(salary)),  
              by = .(department, years_bucket = cut(years, breaks = c(0, 5, 10, 20))))]
```

| | department | years_bucket | avg_salary |
|--|------------|--------------|------------|
| | <char> | <fctr> | <num> |

| | | | |
|-----|---------|---------|----------|
| 1: | HR | (10,20] | 58712.06 |
| 2: | IT | (10,20] | 60364.82 |
| 3: | Finance | (5,10] | 56291.61 |
| 4: | Sales | (10,20] | 58037.80 |
| 5: | Finance | (10,20] | 56846.53 |
| 6: | HR | (0,5] | 59166.40 |
| 7: | IT | (5,10] | 59374.37 |
| 8: | Sales | (0,5] | 59515.41 |
| 9: | Finance | (0,5] | 56726.71 |
| 10: | Sales | (5,10] | 59447.41 |
| 11: | IT | (0,5] | 60374.79 |
| 12: | HR | (5,10] | 60264.12 |

5.10.3 Data.table Performance Advantages

```
# Create large datasets for comparison
n <- 1000000

# Data frame approach
df_large <- data.frame(
  id = 1:n,
  group = sample(letters[1:100], n, replace = TRUE),
  value = rnorm(n)
)

# Data.table approach
dt_large <- data.table(
  id = 1:n,
  group = sample(letters[1:100], n, replace = TRUE),
  value = rnorm(n)
)

# Speed comparison: group aggregation
system.time({
  result_df <- aggregate(value ~ group, data = df_large, FUN = mean)
})
```

| | user | system | elapsed |
|--|-------|--------|---------|
| | 0.092 | 0.014 | 0.106 |

```
system.time({
  result_dt <- dt_large[, .(mean_value = mean(value)), by = group]
})
```

```
      user  system elapsed
0.081    0.003    0.044
```

```
# data.table is typically much faster for large data operations
```

5.11 Factors - Categorical Variables

Factors represent categorical data, similar to SAS formats/value labels.

5.11.1 Creating Factors

```
# Create factor from character vector
gender_char <- c("M", "F", "F", "M", "M", "F")
gender_factor <- factor(gender_char)
print(gender_factor)
```

```
[1] M F F M M F
Levels: F M
```

```
# Levels are shown
levels(gender_factor)
```

```
[1] "F" "M"
```

```
# With explicit levels and labels
gender_factor2 <- factor(
  c("M", "F", "F", "M", "M", "F"),
  levels = c("M", "F"),
  labels = c("Male", "Female")
)
print(gender_factor2)
```

```
[1] Male   Female Female Male   Male   Female
Levels: Male Female
```

```
# Ordered factors
education <- factor(
  c("HS", "BS", "MS", "PhD", "BS", "HS"),
  levels = c("HS", "BS", "MS", "PhD"),
  ordered = TRUE
)
print(education)
```

```
[1] HS   BS   MS   PhD BS   HS
Levels: HS < BS < MS < PhD
```

```
print(education[2] < education[3]) # TRUE: BS < MS
```

```
[1] TRUE
```

5.11.2 Working with Factors

```
# Factor properties
levels(gender_factor)
```

```
[1] "F" "M"
```

```
nlevels(gender_factor)
```

```
[1] 2
```

```
is.factor(gender_factor)
```

```
[1] TRUE
```

```
is.ordered(education)
```

```
[1] TRUE
```

```
# Convert factor to character/numeric
as.character(gender_factor)
```

```
[1] "M" "F" "F" "M" "M" "F"
```

```
as.numeric(gender_factor) # Returns level codes (1, 2, ...)
```

```
[1] 2 1 1 2 2 1
```

```
# Reorder levels
gender_reordered <- factor(gender_factor, levels = c("F", "M"))
levels(gender_reordered)
```

```
[1] "F" "M"
```

```
# Add levels
gender_expanded <- factor(gender_factor, levels = c("M", "F", "O"))
levels(gender_expanded)
```

```
[1] "M" "F" "O"
```

```
# Relabel levels
levels(gender_factor) <- c("Male", "Female")
print(gender_factor)
```

```
[1] Female Male   Male   Female Female Male
Levels: Male Female
```

5.11.3 Practical Uses for Factors

```
# Create survey data
survey <- data.frame(
  id = 1:10,
  satisfaction = factor(
    c("High", "Low", "Medium", "High", "Low", "High", "Medium", "High", "Low", "Medium"),
    levels = c("Low", "Medium", "High"),
```

```

        ordered = TRUE
    ),
    region = factor(c("North", "South", "North", "West", "East",
                      "North", "South", "East", "West", "North"))
)

# Frequency tables (like PROC FREQ)
table(survey$satisfaction)

```

```

Low Medium    High
  3      3      4

```

```
table(survey$region)
```

```

East North South West
  2      4      2      2

```

```
table(survey$satisfaction, survey$region) # Cross-tabulation
```

```

      East North South West
Low      1      0      1      1
Medium   0      2      1      0
High     1      2      0      1

```

```

# Proportions
prop.table(table(survey$satisfaction))

```

```

Low Medium    High
0.3    0.3    0.4

```

```

# Factors preserve level order in plots and analyses
# (Very useful for ordered categories like satisfaction, education, etc.)

```

5.11.4 Factor Pitfalls and Solutions


```
# Pitfall 1: Factors look like characters but aren't
factor_var <- factor(c("10", "20", "30"))
# as.numeric(factor_var) # Returns 1, 2, 3 (NOT 10, 20, 30!)
as.numeric(as.character(factor_var)) # Returns 10, 20, 30 (correct)
```

```
[1] 10 20 30
```

```
# Pitfall 2: Can't add values not in levels
colors <- factor(c("red", "blue", "green"), levels = c("red", "blue", "green"))
# colors[4] <- "yellow" # Would produce NA (not an error!)

# Solution: Add level first
levels(colors) <- c(levels(colors), "yellow")
colors[4] <- "yellow"
print(colors)
```

```
[1] red    blue   green  yellow
Levels: red blue green yellow
```

```
# Pitfall 3: stringsAsFactors in data.frame (older R versions)
# In R < 4.0.0, characters were automatically converted to factors
# Always use stringsAsFactors = FALSE or upgrade to R >= 4.0.0
df_safe <- data.frame(
  name = c("John", "Jane"),
  stringsAsFactors = FALSE
)
str(df_safe) # character, not factor
```

```
'data.frame':  2 obs. of  1 variable:
 $ name: chr  "John" "Jane"
```

6 4. Functions and Help System

6.1 Using Built-in Functions

R has thousands of built-in functions for data analysis.

6.1.1 Statistical Functions

```
# Create sample data
numbers <- c(12, 18, 23, 28, 15, NA, 34, 29, 19, 25)

# Measures of central tendency
mean(numbers, na.rm = TRUE)      # Mean
```

```
[1] 22.55556
```

```
median(numbers, na.rm = TRUE)    # Median
```

```
[1] 23
```

```
# mode (no built-in mode function)

# Measures of dispersion
sd(numbers, na.rm = TRUE)        # Standard deviation
```

```
[1] 7.16085
```

```
var(numbers, na.rm = TRUE)      # Variance
```

```
[1] 51.27778
```

```
IQR(numbers, na.rm = TRUE)      # Interquartile range
```

```
[1] 10
```

```
range(numbers, na.rm = TRUE)    # Min and max
```

```
[1] 12 34
```

```
mad(numbers, na.rm = TRUE)      # Median absolute deviation
```

```
[1] 7.413
```

```
# Summary statistics
```

```
min(numbers, na.rm = TRUE)
```

```
[1] 12
```

```
max(numbers, na.rm = TRUE)
```

```
[1] 34
```

```
sum(numbers, na.rm = TRUE)
```

```
[1] 203
```

```
prod(numbers, na.rm = TRUE)     # Product
```

```
[1] 977240376000
```

```
length(numbers)                 # Count all
```

```
[1] 10
```

```
sum(! is.na(numbers))          # Count non-missing
```

```
[1] 9
```

```
# Quantiles  
quantile(numbers, na.rm = TRUE)  # Default: 0%, 25%, 50%, 75%, 100%
```

```
0%  25%  50%  75% 100%  
12   18   23   28  34
```

```
quantile(numbers, probs = c(0.1, 0.9), na.rm = TRUE) # 10th and 90th percentiles
```

```
10%  90%  
14.4 30.0
```

```
# Summary function (multiple stats at once)  
summary(numbers)
```

```
      Min. 1st Qu.  Median    Mean 3rd Qu.    Max.     NA's  
12.00   18.00   23.00   22.56   28.00   34.00         1
```

6.1.2 Mathematical Functions

```
# Arithmetic  
abs(-5)          # Absolute value
```

```
[1] 5
```

```
sqrt(16)         # Square root
```

```
[1] 4
```

```
exp(2)           # Exponential
```

```
[1] 7.389056
```

```
log(100) # Natural logarithm
```

```
[1] 4.60517
```

```
log10(100) # Base-10 logarithm
```

```
[1] 2
```

```
log(8, base = 2) # Custom base logarithm
```

```
[1] 3
```

```
# Rounding  
round(3.14159, digits = 2) # 3.14
```

```
[1] 3.14
```

```
floor(3.9) # 3 (round down)
```

```
[1] 3
```

```
ceiling(3.1) # 4 (round up)
```

```
[1] 4
```

```
trunc(3.9) # 3 (truncate decimal)
```

```
[1] 3
```

```
signif(123456, digits = 3) # 123000 (significant figures)
```

```
[1] 123000
```

```
# Trigonometry
```

```
sin(pi/2) # 1
```

```
[1] 1
```

```
cos(0) # 1
```

```
[1] 1
```

```
tan(pi/4) # 1
```

```
[1] 1
```

```
asin(1) # pi/2 (arcsin)
```

```
[1] 1.570796
```

```
# Powers and roots
```

```
2^10 # 1024
```

```
[1] 1024
```

```
10^3 # 1000
```

```
[1] 1000
```

```
27^(1/3) # 3 (cube root)
```

```
[1] 3
```

6.1.3 String Functions

```
# Basic string operations  
text <- " Hello World "
```

```
nchar(text)                                # Length (with spaces)
```

```
[1] 15
```

```
toupper(text)                             # " HELLO WORLD "
```

```
[1] " HELLO WORLD "
```

```
tolower(text)                             # " hello world "
```

```
[1] " hello world "
```

```
trimws(text)                             # Remove leading/trailing whitespace
```

```
[1] "Hello World"
```

```
# Substring operations  
substr("Hello World", 1, 5)               # "Hello"
```

```
[1] "Hello"
```

```
substring("Hello World", 7)              # "World"
```

```
[1] "World"
```

```
# Search and replace  
grepl("World", text)                     # TRUE (pattern found)
```

```
[1] TRUE
```

```
grep("World", c("Hello", "World", "Goodbye")) # 2 (position)
```

```
[1] 2
```

```
sub("World", "Universe", text) # Replace first occurrence
```

```
[1] " Hello Universe "
```

```
gsub("o", "0", text) # Replace all occurrences
```

```
[1] " Hello W0rld "
```

```
# Split strings
```

```
strsplit("apple,banana,cherry", ",") # Split by delimiter
```

```
[[1]]
```

```
[1] "apple" "banana" "cherry"
```

```
# Paste strings
```

```
paste("Hello", "World") # "Hello World" (with space)
```

```
[1] "Hello World"
```

```
paste0("Hello", "World") # "HelloWorld" (no space)
```

```
[1] "HelloWorld"
```

```
paste(c("A", "B", "C"), collapse = "-") # "A-B-C"
```

```
[1] "A-B-C"
```

```
# Formatted strings (like sprintf in C)
```

```
sprintf("Patient %d: BMI = %.2f", 12345, 24.567)
```

```
[1] "Patient 12345: BMI = 24.57"
```

6.1.4 Type Checking and Conversion Functions


```
# Type checking (is.* functions)
x <- 42
is.numeric(x)
```

```
[1] TRUE
```

```
is.integer(x)
```

```
[1] FALSE
```

```
is.character(x)
```

```
[1] FALSE
```

```
is.logical(x)
```

```
[1] FALSE
```

```
is.factor(x)
```

```
[1] FALSE
```

```
is.data.frame(x)
```

```
[1] FALSE
```

```
is.list(x)
```

```
[1] FALSE
```

```
is.matrix(x)
```

```
[1] FALSE
```

```
is.na(x)
```

```
[1] FALSE
```

```
is.null(x)
```

```
[1] FALSE
```

```
# Type conversion (as.* functions)  
as.character(42)          # "42"
```

```
[1] "42"
```

```
as.numeric("42")          # 42
```

```
[1] 42
```

```
as.integer(42.7)          # 42 (truncates)
```

```
[1] 42
```

```
as.logical(1)             # TRUE
```

```
[1] TRUE
```

```
as.factor(c("A", "B", "C"))
```

```
[1] A B C  
Levels: A B C
```

```
as.data.frame(matrix(1:6, nrow = 2))
```

```
  V1 V2 V3  
1  1  3  5  
2  2  4  6
```

```
as.list(c(1, 2, 3))
```

```
[[1]]  
[1] 1
```

```
[[2]]  
[1] 2
```

```
[[3]]  
[1] 3
```

```
as.matrix(data.frame(x = 1:3, y = 4:6))
```

```
      x y  
[1,] 1 4  
[2,] 2 5  
[3,] 3 6
```

6.2 SAS PROC to R Function Mapping

Here's a comprehensive mapping of common SAS procedures to R equivalents:

```
# Create mapping table  
sas_to_r <- data.frame(  
  SAS_Proc = c(  
    "PROC MEANS",  
    "PROC FREQ",  
    "PROC PRINT",  
    "PROC SORT",  
    "PROC CONTENTS",  
    "PROC SQL",  
    "PROC UNIVARIATE",  
    "PROC CORR",  
    "PROC REG",  
    "PROC LOGISTIC",  
    "PROC TABULATE",  
    "PROC TRANSPOSE",  
    "PROC IMPORT",  
    "PROC EXPORT"
```

```

),
R_Function = c(
  "summary(), mean(), sd()",
  "table(), prop.table()",
  "print(), head(), View()",
  "sort(), order(), arrange()",
  "str(), glimpse(), attributes()",
  "sqldf(), dplyr verbs",
  "summary(), psych::describe()",
  "cor(), cor.test()",
  "lm(), summary()",
  "glm(family='binomial')",
  "ftable(), aggregate()",
  "t(), pivot_longer()/pivot_wider()",
  "read.csv(), read_excel()",
  "write.csv(), write_xlsx()"
),
Package = c(
  "base",
  "base",
  "base",
  "base/dplyr",
  "base/dplyr",
  "sqldf/dplyr",
  "base/psych",
  "base",
  "stats",
  "stats",
  "base/tidyr",
  "base/tidyr",
  "base/readxl",
  "base/writexl"
),
stringsAsFactors = FALSE
)

print(sas_to_r)

```

| | SAS_Proc | R_Function | Package |
|---|------------|-------------------------|---------|
| 1 | PROC MEANS | summary(), mean(), sd() | base |
| 2 | PROC FREQ | table(), prop.table() | base |
| 3 | PROC PRINT | print(), head(), View() | base |

| | | | |
|----|-----------------|-----------------------------------|--------------|
| 4 | PROC SORT | sort(), order(), arrange() | base/dplyr |
| 5 | PROC CONTENTS | str(), glimpse(), attributes() | base/dplyr |
| 6 | PROC SQL | sqldf(), dplyr verbs | sqldf/dplyr |
| 7 | PROC UNIVARIATE | summary(), psych::describe() | base/psych |
| 8 | PROC CORR | cor(), cor.test() | base |
| 9 | PROC REG | lm(), summary() | stats |
| 10 | PROC LOGISTIC | glm(family='binomial') | stats |
| 11 | PROC TABULATE | ftable(), aggregate() | base/tidyr |
| 12 | PROC TRANSPOSE | t(), pivot_longer()/pivot_wider() | base/tidyr |
| 13 | PROC IMPORT | read.csv(), read_excel() | base/readxl |
| 14 | PROC EXPORT | write.csv(), write_xlsx() | base/writexl |

6.2.1 Practical Examples: SAS to R

```
: :: {.panel-tabset}
```

6.3 PROC MEANS

```
# Create data
data <- data.frame(
  group = rep(c("A", "B", "C"), each = 10),
  value = c(rnorm(10, 100, 15), rnorm(10, 110, 15), rnorm(10, 95, 15))
)

# SAS: PROC MEANS DATA=data MEAN STD MIN MAX; VAR value; CLASS group; RUN;

# R (base)
aggregate(value ~ group, data = data,
  FUN = function(x) c(mean = mean(x), sd = sd(x),
    min = min(x), max = max(x)))
```

| | group | value.mean | value.sd | value.min | value.max |
|---|-------|------------|----------|-----------|-----------|
| 1 | A | 94.89372 | 15.21235 | 62.79850 | 113.28803 |
| 2 | B | 110.89518 | 14.17912 | 86.46798 | 127.66605 |
| 3 | C | 100.08641 | 15.35787 | 81.35258 | 123.05135 |

```
# R (dplyr)
data %>%
  group_by(group) %>%
```

```

summarise(
  n = n(),
  mean = mean(value),
  sd = sd(value),
  min = min(value),
  max = max(value)
)

```

```

# A tibble: 3 x 6
  group    n mean   sd   min   max
<chr> <int> <dbl> <dbl> <dbl> <dbl>
1 A         10  94.9  15.2  62.8  113.
2 B         10  111.   14.2  86.5  128.
3 C         10  100.   15.4  81.4  123.

```

6.4 PROC FREQ

```

# Create data
survey_data <- data.frame(
  gender = sample(c("M", "F"), 100, replace = TRUE),
  satisfaction = sample(c("Low", "Medium", "High"), 100, replace = TRUE)
)

# SAS: PROC FREQ DATA=survey_data; TABLES gender satisfaction gender*satisfaction; RUN;

# R: One-way frequency
table(survey_data$gender)

```

```

  F  M
50 50

```

```
prop.table(table(survey_data$gender))
```

```

  F  M
0.5 0.5

```

```
# Two-way cross-tabulation
table(survey_data$gender, survey_data$satisfaction)
```

| | High | Low | Medium |
|---|------|-----|--------|
| F | 16 | 17 | 17 |
| M | 13 | 18 | 19 |

```
prop.table(table(survey_data$gender, survey_data$satisfaction))
```

| | High | Low | Medium |
|---|------|------|--------|
| F | 0.16 | 0.17 | 0.17 |
| M | 0.13 | 0.18 | 0.19 |

```
# With dplyr
survey_data %>%
  count(gender, satisfaction) %>%
  mutate(proportion = n / sum(n))
```

| | gender | satisfaction | n | proportion |
|---|--------|--------------|----|------------|
| 1 | F | High | 16 | 0.16 |
| 2 | F | Low | 17 | 0.17 |
| 3 | F | Medium | 17 | 0.17 |
| 4 | M | High | 13 | 0.13 |
| 5 | M | Low | 18 | 0.18 |
| 6 | M | Medium | 19 | 0.19 |

6.5 PROC SORT

```
# Create data
unsorted <- data.frame(
  id = c(3, 1, 4, 2, 5),
  name = c("Charlie", "Alice", "David", "Bob", "Eve"),
  score = c(85, 92, 78, 88, 95)
)
```

```
# SAS: PROC SORT DATA=unsorted OUT=sorted; BY score; RUN;

# R (base)
sorted_base <- unsorted[order(unsorted$score), ]
sorted_desc <- unsorted[order(-unsorted$score), ] # Descending

# R (dplyr)
sorted_dplyr <- unsorted %>% arrange(score)
sorted_desc_dplyr <- unsorted %>% arrange(desc(score))

print(sorted_dplyr)
```

| | id | name | score |
|---|----|---------|-------|
| 1 | 4 | David | 78 |
| 2 | 3 | Charlie | 85 |
| 3 | 2 | Bob | 88 |
| 4 | 1 | Alice | 92 |
| 5 | 5 | Eve | 95 |

```
:::
```

6.6 Getting Help in R

R has comprehensive built-in documentation and help systems.

6.6.1 Basic Help Commands

```
# Help on specific function
? mean
help(mean)

# Search help for keyword
?? regression
help.search("regression")

# Find functions with pattern in name
apropos("mean")
```



```
# Examples from help page
example(mean)
example(plot)

# See function arguments
args(mean)
args(lm)

# See function source code
mean.default
lm # For most functions, just type the name
```

6.6.2 Exploring Packages and Functions

```
# List all installed packages
installed.packages()

# List functions in a package
ls("package:dplyr")

# Help on a package
help(package = "dplyr")

# Vignettes (package tutorials)
vignette()           # List all vignettes
vignette("dplyr")    # Specific vignette
browseVignettes("dplyr") # Browse all package vignettes
```

6.6.3 Online Resources and Cheat Sheets

```
# R documentation website
# https://www.rdocumentation.org/

# RStudio cheat sheets
# https://posit.co/resources/cheatsheets/

# Stack Overflow for R questions
# https://stackoverflow.com/questions/tagged/r
```

```
# R-bloggers for tutorials
# https://www.r-bloggers.com/
```

6.7 Writing Custom Functions

Creating reusable functions is essential for efficient R programming.

6.7.1 Basic Function Definition

```
# Simple function
calculate_bmi <- function(weight_kg, height_m) {
  bmi <- weight_kg / (height_m^2)
  return(bmi)
}

# Test function
my_bmi <- calculate_bmi(70, 1.75)
print(paste("BMI:", round(my_bmi, 2)))
```

```
[1] "BMI: 22.86"
```

```
# Calculate for multiple people
weights <- c(70, 85, 60)
heights <- c(1.75, 1.80, 1.65)
bmis <- mapply(calculate_bmi, weights, heights)
print(round(bmis, 2))
```

```
[1] 22.86 26.23 22.04
```

6.7.2 Function with Default Arguments

```
# Function with default parameters
greet_user <- function(name, greeting = "Hello") {
  message <- paste(greeting, name)
  return(message)
}
```

```
}  
# Test function  
print(greet_user("Alice"))           # Uses default greeting
```

```
[1] "Hello Alice"
```

```
print(greet_user("Bob", greeting = "Hi")) # Custom greeting
```

```
[1] "Hi Bob"
```

6.7.3 Function with Variable Number of Arguments

```
# Function that accepts variable number of arguments  
sum_numbers <- function(...) {  
  numbers <- c(...)  
  total <- sum(numbers, na.rm = TRUE)  
  return(total)  
}  
# Test function  
print(sum_numbers(1, 2, 3, 4, 5))
```

```
[1] 15
```

```
print(sum_numbers(10, 20, NA, 30))
```

```
[1] 60
```

7 datatype and structure exercise

8 Exercise 1

Install and load the following packages

`{tidyverse}` `{admiral}` `{dplyr}` `{tidyr}` `{admiral.test}`

```
#installing the packages
install.packages(c("tidyverse", "admiral", "dplyr", "tidyr"))

library(tidyverse)
library(admiral)
library(admiral.test)
library(dplyr)
library(tidyr)
```

9 Exercise 2

Import `adsl.sas7bdat` as `adsl`

```
library(haven)  
adsl <- read_sas("data/adsl.sas7bdat")
```

Part II

data manipulation

10 Introduction

This is a book created from markdown and executable code.

See Knuth (1984) for additional discussion of literate programming.

```
1 + 1
```

```
[1] 2
```


11 Summary

In summary, this book has no content whatsoever.

1 + 1

[1] 2

References

Knuth, Donald E. 1984. “Literate Programming.” *Comput. J.* 27 (2): 97–111. <https://doi.org/10.1093/comjnl/27.2.97>.