



Introduction to Statistics in R

Workshop 1 – September 16, 2025



Data exploration and visualization in R

Hands-on tutorials to help build skills required in Precision Child Health, data science and bioinformatics for SickKids staff and trainees.

Topics include:

- Data exploration in R
- Preparing data for analysis
- Data visualization using ggplot

No prior knowledge of programming is required. We encourage you to bring your laptop to practice data analysis steps under our guidance.

To register please visit: http://ccm20250916.eventbrite.com/

Hands-on
Data Science
Tutorials
for Biologists

Tuesday Sep 16, 2025 12:00 – 1:00 pm

Multi-media Room PGCRL 3rd floor or online (hybrid event)



CCM Overview

Note that prior knowledge of R is recommended to get the most out of this workshop series.

Workshop 1	Data exploration and visualization		
Workshop 2	Statistical tests and models in R		

Check out other workshop series hosted by CCM at https://ccm.sickkids.ca/bioinformatics-training/

Supported by:

Centre for Computational Medicine (ccm.sickkids.ca)
Digital Research Alliance of Canada (alliancecan.ca)
Compute Ontario (computeontario.ca)

Table of contents

01

Data exploration and cleaning

Overview of the Framingham heart dataset Data exploration Basic data frame manipulation

02

Data visualization

Plotting with base R vs ggplot2.

Distributions

Bar charts

Scatterplots

Panel figures

01

Data exploration and cleaning

Framingham Heart Study

One of the longest prospective epidemiological studies of cardiovascular disease and its risk factors.

Began in 1948 in Framingham, MA.

- Initially enrolled 5209 men and women aged 29-62 years old.
- Followed them over time, with assessments every 2 years.
- Data collection for the original cohort ended in 2014.
- Over time, the researchers incorporated offspring and their spouses into the study.
- Today, they have data from three generations of participants as well as more ethnically diverse cohorts.

Framingham Heart Study

Examinations included:

- Detailed medical history
- Physical exams
- Lab tests
- Lifestyle and habits
- Noninvasive imaging

Our dataset: subset of the FHS from Kaggle

• 4000+ records and 16 variables.

(https://www.kaggle.com/datasets/captainozlem/framingham-chd-preprocessed-data)

Variable	Description	Class/type	
male	Sex (male or female)	Binary	
age	Age	Continuous	
education	Education (0-11 years, HS or GED, some uni, uni grad+)	Ordinal/categorical	
currentSmoker	Whether the patient is currently a smoker	Binary	
cigsPerDay	Number of cigarettes smoked per day, on average	Continuous	
BPMeds	Whether the patient is on blood pressure medication	Binary	
prevalentStroke	Whether the patient previously had a stroke	Binary	
prevalentHyp	Whether the patient is hypertensive	Binary	
diabetes	Whether the patient has diabetes	Binary	
totChol	Total cholesterol level	Continuous	
sysBP	Systolic blood pressure	Continuous	
diaBP	Diastolic blood pressure	Continuous	
BMI	Body mass index	Continuous	
heartRate	Heart rate	Continuous	
glucose	Glucose level	Continuous	
TenYearCHD	10-year risk of coronary heart disease	Binary	

Some common and/or important data cleaning steps include making sure that:

- 1. All variables are in the right format and/or correctly classified.
- 2. Duplicate or missing data are dealt with.
- 3. Variables of interest have been transformed appropriately.
- 4. Outliers are handled.
- 5. Data are validated.

1. Look at the data and its structure – are all the variables in the right format? str()

```
2. Tabulate data
```

```
table(df$column) or table(df$column1, df$column2, ...)
```

3. Modify variables

```
mutate(column = function(column))
mutate(df, across(columns, function))
```

4. Transform variables – center, scale, log-transform

```
log(x, base = exp(1))
scale(x, center = TRUE, scale = TRUE)
```

```
5. Subset dataframes
filter() rows according to some condition
select() columns to keep
subset(df, subset = row_vals == "abc", cols = X)
6. Combine vectors and dataframes
cbind() to join by columns
rbind() to join by rows
merge() to join dataframes
```

7. Reshape dataframes

pivot_wider() each row is a participant/patient

pivot_longer() each row is an observation, with multiple rows per participant

8. Handy functions, pipes, and operators

ifelse() for conditional element selection
case_when() as a vectorized version of multiple ifelse() statements, uses formula
notation (more on that later)

%>% passes the result of the function on the left as input to the next function %in% to compare vectors

Create a summary table

Use summarize() to create a dataframe with summaries of desired variables. For example, what if you're only interested in the educational attainment and smoking habits of people who go on to develop heart disease?

•	TenYearCHD	† education_class †	n_participants ‡	age ‡	currentSmoker ‡	prop_smokers ‡	cigsPerDay ‡
1	0	< High School	1397	51	645	0.46	8.39
2	0	College Graduate	403	47	204	0.51	9.09
3	0	High School Graduate	1106	47	598	0.54	9.67
4	0	Some College	599	48	275	0.46	7.70
5	1	< High School	323	56	160	0.50	9.93
6	1	College Graduate	70	53	36	0.51	11.97
7	1	High School Graduate	147	52	81	0.55	11.14
8	1	Some College	88	53	46	0.52	10.74