



🔧 PROJECT LAUNCH

New Project

File > New Project > New Directory

New R script

File > New File > R Script

Where am I?

```
# Show current working directory
getwd()
# Set new working directory
setwd("C:/my-data-folder")
```

Install new packages

```
install.packages("readr")
library(readr)
```

PLOTS

Scatterplot

```
library(ggplot2)

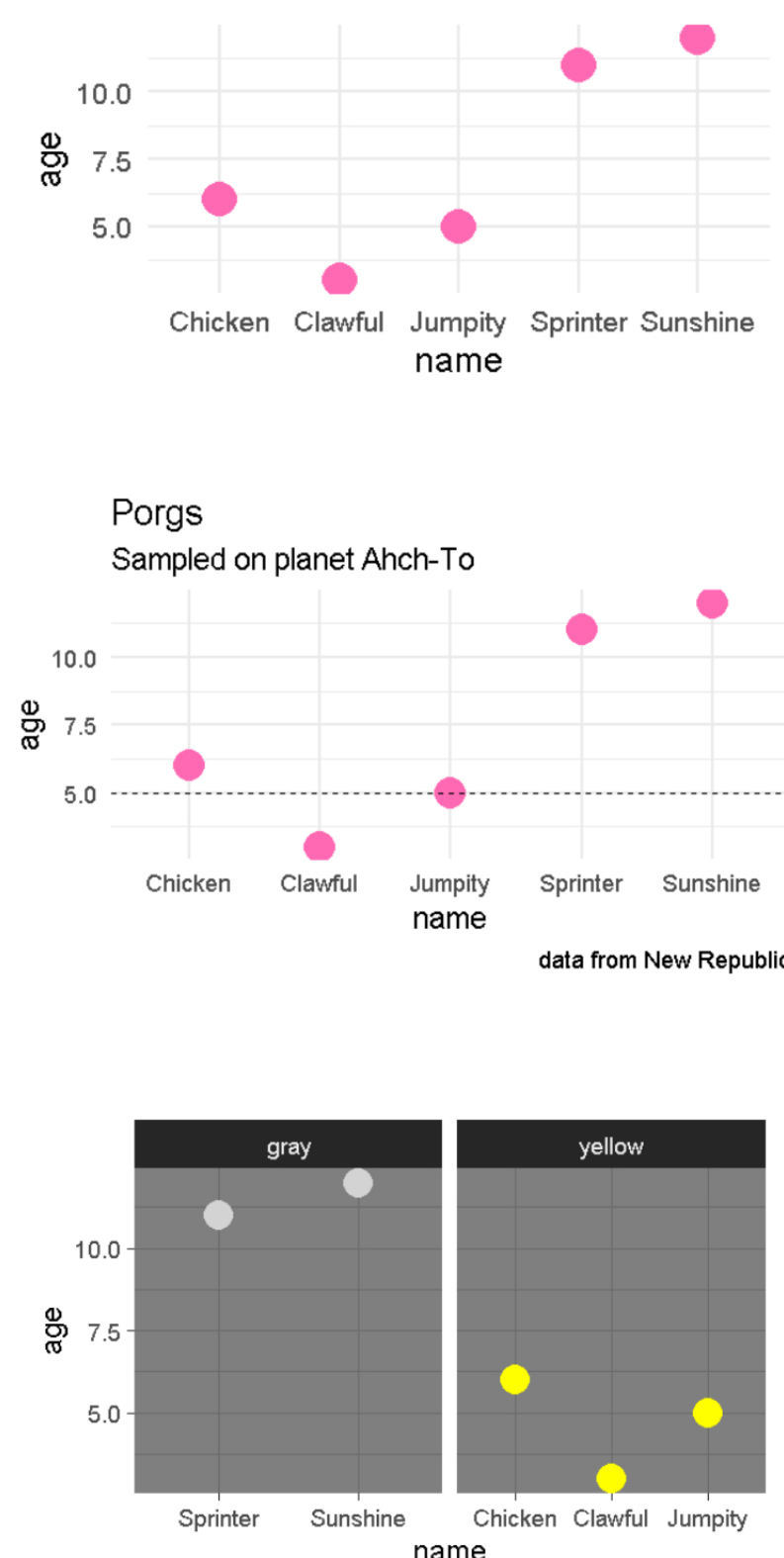
ggplot(porgs, aes(x = name, y = age)) +
  geom_point(size = 8, color = "hotpink")
```

Add titles & lines

```
ggplot(porgs, aes(x = name, y = age)) +
  geom_point(size = 8, color = "hotpink") +
  geom_hline(yintercept = 5,
            linetype = "dashed") +
  labs(title = "Porgs",
       subtitle = "Sampled on planet Ahch-To",
       caption = "data from New Republic")
```

Facet by group

```
ggplot(porgs, aes(x = name, y = age)) +
  geom_point(aes(color = color), size = 8) +
  facet_wrap(~color) +
  scale_color_manual(values = c("gray", "yellow")) +
  theme_dark()
```



STORE VALUES

```
# Use the Left-arrow
age <- 7.2

# Text goes in quotes
porg <- "Sunshine"

# Multiple values go inside c()
droids <- c("BB8", "R2D2", "C-3PO")

# Copy an object
my_droids <- droids

# Avoid numbers, spaces, & symbols
3-droids <- "error_invalid_name"
```

READ DATA

📄 Text files (.csv, .txt, .tab)

```
library(readr)
porgs <- read_csv("txt_file.csv")
```

📄 Excel files (.xlsx, .xls)

```
library(readxl)
porgs <- read_excel("Excel_file.xlsx")
```

CLEAN NAMES 🔧

```
# Simplify all column names
library(janitor)
porgs <- clean_names(porgs)

# Assign new names manually
library(dplyr)
# Put new name on left: new_name = oldName
rename(porgs, mass_kg = massKG)
```

FILTER

```
library(dplyr)
# Keep only Porgs with age greater than 3
filter(porgs, age > 3)
# Keep rows with name Jumpity
filter(porgs, name == "Jumpity")
# Keep Porgs named Jumpity -OR- Chicken
filter(porgs, name %in% c("Jumpity", "Chicken"))
```

SUMMARIZE

```
library(dplyr)
# Summarize the age for the entire table
summarize(porgs, avg_age = mean(age))
# Summarize the age for each color group
group_by(porgs, color) %>%
  summarize(avg_age = mean(age))
```

DESCRIBE DATA

```
library(dplyr)
nrow(porgs)
names(porgs)
summary(porgs)
glimpse(porgs)
class(porgs)
# View unique column values
distinct(porgs, age)
## 5 6 11 12 3
```

ADD COLUMNS

```
library(dplyr)

# Add home planet column
mutate(porgs,
       planet = "Ahch-To")
# Calculate new BMI column
mutate(porgs,
       bmi = mass / height)
```

COMPARISONS

Symbol	Comparison
>	greater than
>=	greater than or equal to
<	less than
<=	less than or equal to
==	equal to
!=	NOT equal to
%in%	is value in a list: X %in% c(1,3,5)
is.na(...)	is the value missing?