

Programming in Base R

```
library(tidyverse)
```

```
-- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
v dplyr      1.1.3      v readr      2.1.4
v forcats    1.0.0      v stringr    1.5.0
v ggplot2    3.5.1      v tibble     3.2.1
v lubridate  1.9.3      v tidyr      1.3.0
v purrr      1.0.2
-- Conflicts ----- tidyverse_conflicts() --
x dplyr::filter() masks stats::filter()
x dplyr::lag()     masks stats::lag()
i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become
```

```
library(palmerpenguins)
```

Task 1

Question 1

```
?read_csv
```

We cannot use `read_csv` specifically because our delimiter for the given data is a semicolon, not a comma. There are 2 other functions described in the help file called `read_csv2` and `read_delim` which allow for more control over the delimiter, and `read_csv2` specifically uses `;` as the delimiter which we will use.

```
data <- read_csv2("data/data.txt")
```

i Using "','" as decimal and "'.'" as grouping mark. Use `read_delim()` for more control.

Rows: 2 Columns: 3

-- Column specification -----

Delimiter: ";"

dbl (3): x, y, z

i Use `spec()` to retrieve the full column specification for this data.

i Specify the column types or set `show_col_types = FALSE` to quiet this message.

```
data
```

```
# A tibble: 2 x 3
```

	x	y	z
	<dbl>	<dbl>	<dbl>
1	1	2	3
2	5	3	8

Question 2

```
data2 <- read_delim("data/data2.txt", delim = '6', col_types = "fdc")
data2
```

```
# A tibble: 3 x 3
```

	x	y	z
	<fct>	<dbl>	<chr>
1	1	2	3
2	5	3	8
3	7	4	2

Task 2

Question 1

```
trailblazer <- read_csv("data/trailblazer.csv")
```

```

Rows: 9 Columns: 11
-- Column specification -----
Delimiter: ","
chr  (1): Player
dbl (10): Game1_Home, Game2_Home, Game3_Away, Game4_Home, Game5_Home, Game6_...

i Use `spec()` to retrieve the full column specification for this data.
i Specify the column types or set `show_col_types = FALSE` to quiet this message.

```

```
glimpse(trailblazer)
```

```

Rows: 9
Columns: 11
$ Player      <chr> "Damian Lillard", "CJ McCollum", "Norman Powell", "Robert ~
$ Game1_Home  <dbl> 20, 24, 14, 8, 20, 5, 11, 2, 7
$ Game2_Home  <dbl> 19, 28, 16, 6, 9, 5, 18, 8, 11
$ Game3_Away  <dbl> 12, 20, NA, 0, 4, 8, 12, 5, 5
$ Game4_Home  <dbl> 20, 25, NA, 3, 17, 10, 17, 8, 9
$ Game5_Home  <dbl> 25, 14, 12, 9, 14, 9, 5, 3, 8
$ Game6_Away  <dbl> 14, 25, 14, 6, 13, 6, 19, 8, 8
$ Game7_Away  <dbl> 20, 20, 22, 0, 7, 0, 17, 7, 4
$ Game8_Away  <dbl> 26, 21, 23, 6, 6, 7, 15, 0, 0
$ Game9_Home  <dbl> 4, 27, 25, 19, 10, 0, 16, 2, 7
$ Game10_Home <dbl> 25, 7, 13, 12, 15, 6, 10, 4, 8

```

Question 2

```

trailblazer_longer <- trailblazer |>
  pivot_longer(cols = 2:11,
               names_to = c('Game', 'Location'),
               names_sep = '_',
               values_to = 'Points')
head(trailblazer_longer, 5)

```

```

# A tibble: 5 x 4
  Player      Game Location Points
  <chr>      <chr> <chr>    <dbl>
1 Damian Lillard Game1 Home      20
2 Damian Lillard Game2 Home      19

```

3	Damian Lillard	Game3	Away	12
4	Damian Lillard	Game4	Home	20
5	Damian Lillard	Game5	Home	25

Question 3

```
trailblazer_longer |>
  pivot_wider(names_from = Location, values_from = Points) |>
  group_by(Player) |>
  summarise(mean_home = mean(Home, na.rm = T),
            mean_away = mean(Away, na.rm = T),
            diff = mean_home - mean_away) |>
  arrange(desc(diff))
```

```
# A tibble: 9 x 4
  Player      mean_home mean_away   diff
  <chr>      <dbl>    <dbl> <dbl>
1 Jusuf Nurkic      14.2      7.5  6.67
2 Robert Covington   9.5       3   6.5
3 Nassir Little      8.33     4.25 4.08
4 Damian Lillard     18.8     18  0.833
5 Cody Zeller        5.83     5.25 0.583
6 Larry Nance Jr     4.5       5  -0.5
7 CJ McCollum       20.8     21.5 -0.667
8 Anfernee Simons    12.8     15.8 -2.92
9 Norman Powell      16      19.7 -3.67
```

Jusuf Nurkic, Robert Covington and Nassir Little were the top three players who scored more points at home than away throughout the first 10 games of the season.

Task 3

Question 1

```
?penguins
penguins |>
  select(species, island, bill_length_mm) |>
  pivot_wider(
    names_from = island, values_from = bill_length_mm
```

```
)
```

Warning: Values from `bill_length_mm` are not uniquely identified; output will contain list-cols.

```
* Use `values_fn = list` to suppress this warning.
* Use `values_fn = {summary_fun}` to summarise duplicates.
* Use the following dplyr code to identify duplicates.
{data} %>%
  dplyr::group_by(species, island) %>%
  dplyr::summarise(n = dplyr::n(), .groups = "drop") %>%
  dplyr::filter(n > 1L)
```

```
# A tibble: 3 x 4
  species   Torgersen   Biscoe     Dream
  <fct>     <list>      <list>     <list>
1 Adelie   <dbl [52]> <dbl [44]> <dbl [56]>
2 Gentoo   <NULL>     <dbl [124]> <NULL>
3 Chinstrap <NULL>     <NULL>     <dbl [68]>
```

We did not group by the species and island, so there are errors when we run this code since there are multiple values for `bill_length_mm`. means that there are no penguins for that combination of species and island. `<dbl [52]>` means that for the Adelie species on Torgersen, there are 52 dbl values, so there are 52 different values of `bill_length_mm` for that group. means that the column is a list since it is comprised of many dbls or null values.

Question 2

```
penguins |>
  group_by(species, island) |>
  summarise(count = n()) |>
  pivot_wider(names_from = island, values_from = count, values_fill = 0)
```

``summarise()`` has grouped output by 'species'. You can override using the ``.groups`` argument.

```
# A tibble: 3 x 4
# Groups:   species [3]
  species   Biscoe Dream Torgersen
  <fct>     <int> <int>    <int>
1 Adelie     44    56     52
2 Gentoo    124    NA     NA
3 Chinstrap  68    NA     NA
```

1 Adelie	44	56	52
2 Chinstrap	0	68	0
3 Gentoo	124	0	0

Task 4

Question 1

```
penguins_nona <- penguins |>
  mutate(bill_length_mm = case_when(
    (species == 'Adelie') & is.na(bill_length_mm) ~ 26,
    (species == 'Gentoo') & is.na(bill_length_mm) ~ 30,
    TRUE ~ bill_length_mm
  ))

head(penguins_nona, 10)
```

A tibble: 10 x 8

	species	island	bill_length_mm	bill_depth_mm	flipper_length_mm	body_mass_g
	<fct>	<fct>	<dbl>	<dbl>	<int>	<int>
1	Adelie	Torgersen	39.1	18.7	181	3750
2	Adelie	Torgersen	39.5	17.4	186	3800
3	Adelie	Torgersen	40.3	18	195	3250
4	Adelie	Torgersen	26	NA	NA	NA
5	Adelie	Torgersen	36.7	19.3	193	3450
6	Adelie	Torgersen	39.3	20.6	190	3650
7	Adelie	Torgersen	38.9	17.8	181	3625
8	Adelie	Torgersen	39.2	19.6	195	4675
9	Adelie	Torgersen	34.1	18.1	193	3475
10	Adelie	Torgersen	42	20.2	190	4250

i 2 more variables: sex <fct>, year <int>