



Manipulação e Apresentação de Dados

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```
modelo.aov <- aov(Sepal.Length ~ Species, data=iris)
modelo.lm <- lm(Sepal.Width ~ Petal.Width, data=iris)
modelo.wt <- wilcox.test(iris$Petal.Length, g=iris$Species)</pre>
```

##

1



```
library(broom)
tidy(modelo.aov)
## # A tibble: 2 x 6
## term df sumsq meansq statistic p.value
## <chr> <dbl> <dbl> <dbl> <dbl>
                                       <dbl>
## 1 Species 2 63.2 31.6 119. 1.67e-31
## 2 Residuals 147 39.0 0.265 NA NA
tidy(modelo.lm)
## # A tibble: 2 x 5
## term estimate std.error statistic p.value
## <chr> <dbl> <dbl> <dbl>
                                        <dbl>
## 1 (Intercept) 3.31 0.0621 53.3 1.84e-98
## 2 Petal.Width -0.209 0.0437 -4.79 4.07e- 6
tidy(modelo.wt)
## # A tibble: 1 x 4
```

11325 2.25e-26 Wilcoxon signed rank test with continuity corr~ two.sid

statistic p.value method

<dbl> <dbl> <chr>

5-

alterna

<chr>



```
glance(modelo.aov)
## # A tibble: 1 x 6
##
    logLik AIC BIC deviance nobs r.squared
## <dbl> <dbl> <dbl> <int> <dbl> <int>
## 1 -112, 231, 243, 39,0 150
                                      0.619
glance(modelo.lm)
## # A tibble: 1 x 12
##
    r.squared adj.r.squared sigma statistic p.value df logLik AIC
                                                                   BI
        <dbl>
                    <dbl> <dbl> <dbl>
                                             <dbl> <dbl> <dbl> <dbl> <dbl> <dbl
##
## 1
       0.134
                    0.128 0.407 22.9 0.00000407 1 -77.0 160. 169
## # i 3 more variables: deviance <dbl>, df.residual <int>, nobs <int>
glance(modelo.wt)
## # A tibble: 1 x 4
##
    statistic p.value method
                                                                 alterna
##
        <dbl> <dbl> <chr>
                                                                 <chr>
        11325 2.25e-26 Wilcoxon signed rank test with continuity corr~ two.sid
## 1
```



augment(modelo.lm, interval="confidence")

```
A tibble: 150 x 10
##
      Sepal.Width Petal.Width .fitted .lower .upper
                                                         .resid
                                                                   .hat
                                                                        .sig
             <dbl>
                         <dbl>
                                  <dbl>
                                          <dbl>
                                                 <dbl>
                                                         <dbl>
                                                                 <dbl>
##
                                                                         <db
##
    1
               3.5
                            0.2
                                   3.27
                                          3.16
                                                  3.38
                                                         0.233
                                                                0.0182
                                                                         0.4
    2
               3
##
                            0.2
                                   3.27
                                          3.16
                                                  3.38 - 0.267
                                                                0.0182
                                                                         0.4
    3
                                   3.27
                                                                         0.4
##
               3.2
                            0.2
                                          3.16
                                                  3.38 -0.0666 0.0182
##
    4
               3.1
                            0.2
                                   3.27
                                          3.16
                                                  3.38 -0.167
                                                                0.0182
                                                                         0.4
##
    5
               3.6
                            0.2
                                   3.27
                                           3.16
                                                  3.38
                                                         0.333
                                                                0.0182
                                                                         0.4
##
    6
               3.9
                            0.4
                                   3.22
                                           3.13
                                                  3.32
                                                         0.675
                                                                0.0140
                                                                         0.4
##
    7
               3.4
                            0.3
                                   3.25
                                          3.14
                                                  3.35
                                                         0.154
                                                                0.0160
                                                                         0.4
                           0.2
                                                                         0.4
##
    8
               3.4
                                   3.27
                                          3.16
                                                  3.38
                                                         0.133
                                                                0.0182
##
    9
               2.9
                            0.2
                                   3.27
                                          3.16
                                                  3.38 -0.367
                                                                0.0182
                                                                         0.4
## 10
               3.1
                            0.1
                                   3.29
                                           3.17
                                                  3.40 - 0.187
                                                                0.0206
                                                                         0.4
## # i 140 more rows
## # i 1 more variable: .std.resid <dbl>
```





```
lapply(seq(1,5), function(x) {
  paste(x, "é", ifelse(x\\2==0, "par", "impar"))
})
## [[1]]
## [1] "1 é impar"
##
## [[2]]
## [1] "2 é par"
##
## [[3]]
## [1] "3 é impar"
##
## [[4]]
## [1] "4 é par"
##
## [[5]]
## [1] "5 é impar"
```



```
library(purrr)
map(seq(1,5), function(x) {
  paste(x, "é", ifelse(x\\2==0, "par", "impar"))
})
## [[1]]
## [1] "1 é impar"
##
## [[2]]
## [1] "2 é par"
##
## [[3]]
## [1] "3 é impar"
##
## [[4]]
  [1] "4 é par"
##
## [[5]]
  [1] "5 é ímpar"
```



```
map_dfr(seq(1,5), function(x) {
  tibble(par=paste(x, "é", ifelse(x\%2==0, "par", "impar")))
})
## # A tibble: 5 x 1
## par
## <chr>
## 1 1 é impar
## 2 2 é par
## 3 3 é impar
## 4 4 é par
## 5 5 é impar
```





```
map_dfc(tibble(um=1, dois=2, tres=3), function(x) {
  x * seq(1,5)
})
## # A tibble: 5 x 3
##
           dois tres
        um
     <dbl> <dbl> <dbl>
##
## 1
               2
                      3
## 2
         3
               6
                      9
## 3
## 4
         4
               8
                     12
         5
## 5
               10
                     15
```



library(agridat) caribbean.maize

##		isle	site	block	plot	trt	ears	yield
##	1	Antigua	DBAN	B1	1	T111	42	4.96
##	2	Antigua	DBAN	B1	2	T000	41	3.94
##	3	Antigua	DBAN	B1	3	T311	49	6.35
##	4	Antigua	DBAN	B1	4	T202	48	5.56
##	5	Antigua	DBAN	B1	5	T111	45	5.36
##	6	Antigua	DBAN	B1	6	T220	46	6.18
##	7	Antigua	DBAN	B1	7	T113	42	4.71
##	8	Antigua	DBAN	B1	8	T131	44	6.03
##	9	Antigua	DBAN	B1	9	T022	42	2.88
##	10	Antigua	DBAN	B2	10	T222	44	5.68
##	11	Antigua	DBAN	B2	11	T311	42	5.80
##	12	Antigua	\mathtt{DBAN}	B2	12	T020	42	4.16
##	13	Antigua	DBAN	B2	13	T200	46	4.90
##	14	Antigua	DBAN	B2	14	T111	44	5.25
##	15	Antigua	DBAN	B2	15	T131	48	5.80
##	16	Antiqua	DRAN	R2	16	T002	46	2 18



```
caribbean.maize %>%
 group_split(isle)
## <list_of<
##
    tbl_df<
##
      isle : factor<2a195>
##
      site: factor<459f2>
##
      block: factor<cfca5>
##
   plot : integer
      trt : factor<ed0b9>
##
##
      ears : integer
##
      vield: double
##
    >
## >[2]>
## [[1]]
## # A tibble: 288 x 7
##
     isle site block plot trt ears yield
##
     <fct> <fct> <fct> <int> <fct> <int> <dbl>
   1 Antigua DBAN B1
##
                            1 T111
                                      42 4.96
   O A I C DDAN D4
                            0 5000
```

3 B1

4 B1

5 B1

6 B1

7 B1

8 B1

9 B1

##

##

##

##

##

##

3

4

5

6

7

8

9

6.35

5.56

5.36

6.18

4.71

6.03

2.88



0.83

2.44

1.34

2.06

3.02

2.18

1.3

5.53

6.61

2.97

4.47

4.06

5.02

 $4.56_{15}6.9$

7.7

6.7

6.5

6.4

6.7

6.7

```
caribbean.maize %>%
  group_split(isle) %>%
  map(function(x) x %>%
        pivot wider(id cols=c(block,plot), names from=site,
                     values_from=yield))
## [[1]]
   # A tibble: 36 x 10
##
##
      block plot DBAN
                          LFAN
                                 TEAN
                                       WEAN
                                              WLAN
                                                    NSAN
                                                          OVAN
                                                                 ORA
      <fct> <int> <dbl> <
##
    1 B1
                    4.96
                          2.92 1.27
                                       4.02
                                                    2.43
                                                          2.04
                                                                 5.2
##
                 1
                                             2
    2 B1
                                                                 6.8
##
                 2
                    3.94
                          1.68
                                 2.1
                                       5.8
                                             2.39
                                                    1.28
                                                          3.88
```

4.5

4.66

2.94

1.6

1.82

4.74 2.08

4.44 1.74

2.37

3.16

4.55

3.2

3.69

2.16

5.31

5.12

5.98

5.46

3.45

4.96

2.04

1.64

1.83

2.08

3.13

1.96

1.3

##

##

##

##

##

4 B1

5 B1

6 B1

7 B1

8 B1

9 B1

4

5

6

7

8

9

5.56

5.36

6.18

4.71

6.03

2.88



5.2

6.8

7.7

6.7

6.5

6.4

6.7

6.7

```
caribbean.maize %>%
  group split(isle) %>%
  map(\(x) x \%)
        pivot wider(id cols=c(block,plot), names from=site,
                     values_from=yield))
## [[1]]
   # A tibble: 36 x 10
##
##
      block plot DBAN
                          LFAN
                                TEAN
                                       WEAN
                                             WLAN
                                                   NSAN
                                                          OVAN
                                                                ORA
      <fct> <int> <dbl> <
##
    1 B1
                   4.96
                          2.92 1.27
                                       4.02
                                                   2.43
                                                         2.04
##
                1
                                             2
    2 B1
##
                2
                   3.94
                          1.68
                                2.1
                                       5.8
                                             2.39
                                                   1.28
                                                         3.88
    3 B1
                3
                   6.35
                                                   0.83
                                                         5.53
##
                          4.5
                                2.37
                                       2.16
                                             2.04
```

4.74 2.08

4.44 1.74

3.16

4.55

3.2

3.69

4.66

2.94

1.6

1.82

5.31

5.12

5.98

5.46

3.45

4.96

1.64

1.83

2.08

3.13

1.96

1.3

2.44

1.34

2.06

3.02

2.18

1.3

6.61

2.97

4.47

4.06

5.02

 $4.56_{16}6.9$

##

##

8 B1

9 B1

8

9

6.03

2.88

1.6

1.82



ORA

6.7

```
caribbean.maize %>%
  group split(isle) %>%
  map(. %>%
        pivot wider(id cols=c(block,plot), names from=site,
                     values_from=yield))
## [[1]]
   # A tibble: 36 x 10
##
##
      block plot
                    DBAN
                          LFAN
                                 TEAN
                                       WEAN
                                              WLAN
                                                    NSAN
                                                           OVAN
      <fct> <int> <dbl> <
##
    1 B1
                    4.96
                          2.92 1.27
                                       4.02
                                                           2.04
##
                 1
                                              2
                                                    2.43
```

```
5.2
    2 B1
                                                               6.8
##
                2
                   3.94
                         1.68
                                2.1
                                      5.8
                                            2.39
                                                  1.28
                                                        3.88
    3 B1
                3
                   6.35
                                                  0.83
                                                        5.53
                                                               7.7
##
                         4.5
                                2.37
                                      2.16
                                            2.04
##
    4 B1
                4
                   5.56
                         4.74
                               2.08
                                      5.31
                                            1.64
                                                  2.44
                                                        6.61
                                                               6.7
    5 B1
                5
                   5.36
                         4.44 1.74
                                      5.12
                                            1.83
                                                  1.3
                                                        2.97
                                                               6.5
##
##
    6 B1
                6
                   6.18
                         4.66
                               3.16
                                      5.98
                                            2.08
                                                  1.34
                                                        4.47
                                                               6.4
##
    7 B1
                7
                   4.71
                         2.94
                                4.55
                                      5.46
                                            3.13
                                                  2.06
                                                        4.06
                                                               6.7
```

3.2

3.69

3.45

4.96

1.96

1.3

3.02

2.18

5.02

 $4.56_{17}6.9$



```
iris %>%
  select(-Species) %>%
  as.list() %>%
  map(\(x) x \%)
        enframe() %>%
        cbind(Species=iris$Species) %>%
        aov(value ~ Species, data=.) %>%
        residuals() %>%
        shapiro.test() %>%
        tidy()) %>%
  bind rows(.id="Variável")
```



```
iris %>%
  select(starts_with("Sepal")) %>%
  as.list() %>%
  map(. %>%
      enframe() %>%
      cbind(Species=iris$Species) %>%
      aov(value ~ Species, data=.) %>%
      tidy()) %>%
  bind_rows(.id="Variável")
```

```
## # A tibble: 4 x 7
##
    Variável
                term
                           df sumsq meansq statistic
                                                    p.val
##
    <chr>>
                <chr>
                        <dbl> <dbl> <dbl>
                                             <dbl>
                                                      <db
                                             119.
                                                   1.67e-
  1 Sepal.Length Species
                            2 63.2 31.6
  2 Sepal.Length Residuals 147 39.0 0.265
                                              NA
                                                  NA
## 3 Sepal.Width Species
                          2 11.3 5.67
                                              49.2 4.49e-
## 4 Sepal.Width Residuals
                          147 17.0
                                    0.115
                                              NΑ
                                                  NA
```



```
iris %>%
 select(starts with("Sepal")) %>%
 as.list() %>%
 map dfr(. %>%
          enframe() %>%
          cbind(Species=iris$Species) %>%
          aov(value ~ Species, data=.) %>%
          tidy())
## # A tibble: 4 x 6
##
                df sumsq meansq statistic p.value
    term
##
             <dbl> <dbl> <dbl>
                                  <dbl>
                                           <dbl>
    <chr>
  1 Species
                   63.2 31.6
                                  119.
                                        1.67e-31
##
  2 Residuals 147 39.0 0.265
                                  NA
                                       NA
## 3 Species 2 11.3 5.67
                                  49.2 4.49e-17
## 4 Residuals 147 17.0 0.115
                                   NA
                                       NA
```





```
cor(iris[-5])

## Sepal.Length Sepal.Width Petal.Length Petal.Widt
## Sepal.Length 1.0000000 -0.1175698 0.8717538 0.817941
## Sepal.Width -0.1175698 1.0000000 -0.4284401 -0.366125
## Petal.Length 0.8717538 -0.4284401 1.0000000 0.962865
## Petal.Width 0.8179411 -0.3661259 0.9628654 1.000000
cor.test(iris[-5])

## Error in cor.test.default(iris[-5]): 'X' deve ser um vetor nu
```





[1,] 0.000000e+00 1.518983e-01 1.038667e-47 2.325498e-37 [2,] 1.518983e-01 0.000000e+00 4.513314e-08 4.073229e-06 [3,] 1.038667e-47 4.513314e-08 0.000000e+00 4.675004e-86 [4,] 2.325498e-37 4.073229e-06 4.675004e-86 0.000000e+00





```
expand_grid(Var1=names(iris)[-5],
           Var2=names(iris)[-5]) %>%
 pmap_chr(\(Var1, Var2){
   if(Var1==Var2) return("-")
   cor.test(iris[,Var1], iris[,Var2]) %>%
     .$p.value %>%
     scales::pvalue(acc=.0001, dec=",")
 }) %>%
 matrix(nrow=4)
       [,1] \qquad [,2]
                       [,3] [,4]
##
## [1,] "-"
              "0.1519" "<0.0001" "<0.0001"
## [2,] "0,1519" "-"
                          "<0.0001" "<0.0001"
## [3,] "<0.0001" "<0.0001" "-" "<0.0001"
## [4,] "<0,0001" "<0,0001" "<0,0001" "-"
```

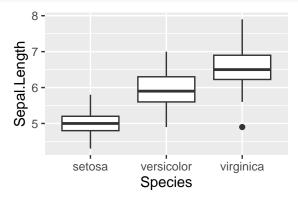




```
map2(c(1,2,3,4), c("a","b","c","d"),
     \(num, chr) paste0("Número: ", num, "; Caractere: ", chr))
## [[1]]
## [1] "Número: 1; Caractere: a"
##
## [[2]]
## [1] "Número: 2; Caractere: b"
##
## [[3]]
## [1] "Número: 3; Caractere: c"
##
## [[4]]
## [1] "Número: 4; Caractere: d"
```



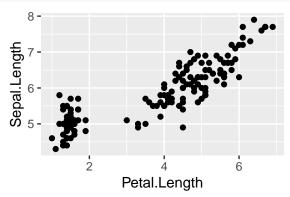
```
library(ggplot2)
iris %>%
    ggplot(aes(x=Species, y=Sepal.Length)) +
    geom_boxplot()
```





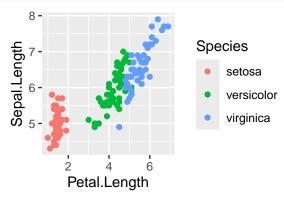


```
iris %>%
  ggplot(aes(x=Petal.Length, y=Sepal.Length)) +
  geom_point()
```

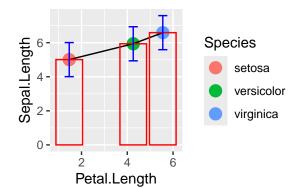






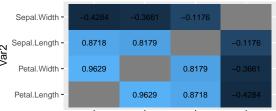








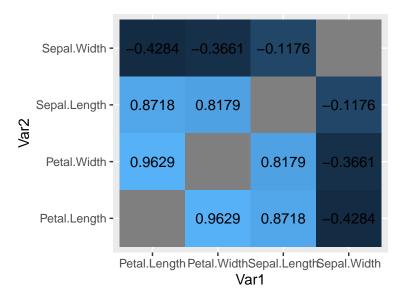
```
iris %>%
  select(-Species) %>%
  cor() %>%
  as.data.frame() %>%
  rownames_to_column("Var1") %>%
  pivot_longer(-Var1, names_to="Var2", values_to="Cor") %>%
  mutate(Cor = ifelse(Var1==Var2, NA, Cor)) %>%
  ggplot(aes(x=Var1, y=Var2, fill=Cor)) +
  geom_tile(show.legend=F) +
  geom_text(aes(label=round(Cor, 4)), size=2) +
  theme_grey(base_size = 8)
```



Petal.Length Petal.Width Sepal.Length Sepal.Width Var1







Exercício

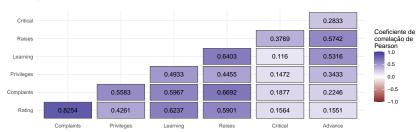
Exercício



Personalizar o gráfico a seguir:

```
cor(attitude) %>%
  as.data.frame() %>%
  rownames_to_column("Var1") %>%
  pivot_longer(-Var1, names_to="Var2") %>%
  ggplot(aes(x=Var1, y=Var2, fill=value)) +
  geom_tile()
```

Exemplo:



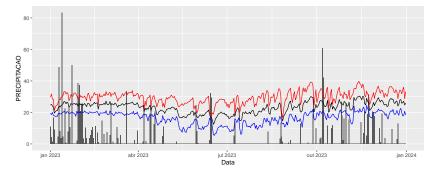
ggplot2 (continuação)



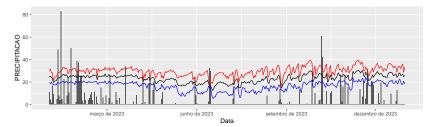
```
linhas <- read lines(
  "http://www.leb.esalq.usp.br/leb/exceldados/DCE2023.TXT")
inicio <- which(str_starts(linhas, "="))[c(1, seq(3, 37, 3))]</pre>
final <- which(str_starts(linhas, "="))[c(2, seq(5,37,3), 37)]
pular <- sapply(seq(1,13), function(i){</pre>
  seq(inicio[i], final[i])
})
pular2 <- which(linhas=="")</pre>
linhas2 <- linhas[-c(unlist(pular).pular2)]
dados metereologicos <- as tibble(linhas2) %>%
  separate(value.
           c("No", "ANO", "DIA", "MES", "R.GLOBA",
             "INSOLAÇÃO", "PRECIPITAÇÃO", "UMIDADE RELATIV",
             "VENTO MAXIMO", "VENTO MEDIO", "TEMPER MAXIMA",
             "TEMPER MINIMA", "TEMPER MEDIA", "EVAPORACAO"),
           sep=" +") %>%
  unite(Data, DIA, MES, ANO) %>%
  mutate(Data=lubridate::dmy(Data)) %>%
  mutate_at(vars(-Data), str_replace, ",", ".") %>%
  mutate at(vars(-Data), parse number) %>%
  filter(!is.na(Data))
```



```
dados_metereologicos %>%
   ggplot(aes(x=Data)) +
   geom_col(aes(y=PRECIPITACAO)) +
   geom_line(aes(y=`TEMPER MEDIA`)) +
   geom_line(aes(y=`TEMPER MINIMA`), col="blue") +
   geom_line(aes(y=`TEMPER MAXIMA`), col="red")
```

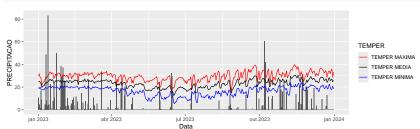




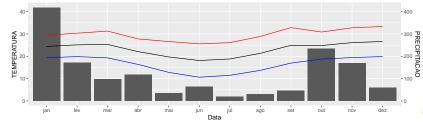




```
dados_metereologicos %>%
  select(Data, PRECIPITACAO, starts_with("TEMPER")) %>%
  pivot_longer(starts_with("TEMPER"), names_to="TEMPER") %>%
  ggplot(aes(x=Data)) +
  geom_bar(stat="unique", aes(y=PRECIPITACAO)) +
  geom_line(aes(y=value, color=TEMPER)) +
  scale_color_manual(values=c("red", "black", "blue"))
```



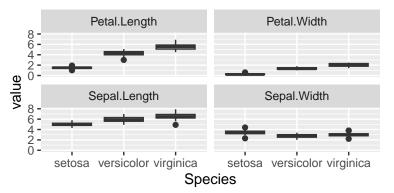








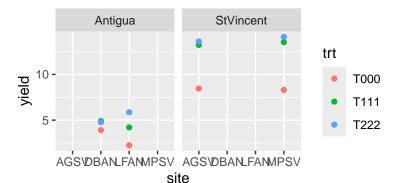
```
iris %>%
  pivot_longer(-Species) %>%
  ggplot(aes(x=Species, y=value)) +
  geom_boxplot() +
  facet_wrap(~name)
```







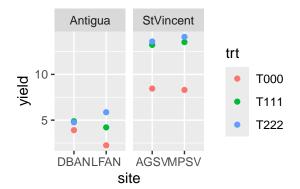
```
agridat::caribbean.maize %>%
  filter(site %in% c("DBAN", "LFAN", "MPSV", "AGSV")) %>%
  filter(trt %in% c("T000", "T111", "T222")) %>%
  group_by(isle, site, trt) %>%
  summarise_at(vars(yield), mean) %>%
  ggplot(aes(x=site, col=trt, y=yield)) +
  geom_point() +
  facet_wrap(~isle)
```



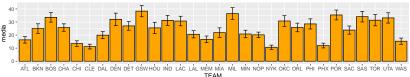




```
agridat::caribbean.maize %>%
  filter(site %in% c("DBAN", "LFAN", "MPSV", "AGSV")) %>%
  filter(trt %in% c("T000", "T111", "T222")) %>%
  group_by(isle, site, trt) %>%
  summarise_at(vars(yield), mean) %>%
  ggplot(aes(x=site, col=trt, y=yield)) +
  geom_point() +
  facet_wrap(~isle, scales = "free_x")
```



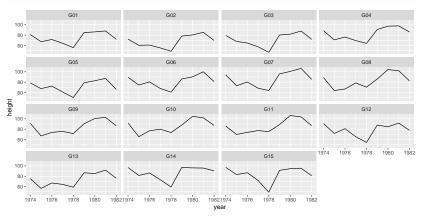






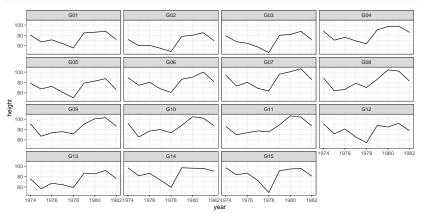


```
agridat::aastveit.barley.height %>%
ggplot(aes(x=year, y=height)) +
facet_wrap(~gen) +
geom_line()
```

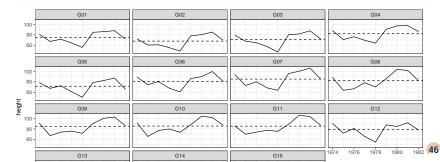




```
agridat::aastveit.barley.height %>%
ggplot(aes(x=year, y=height)) +
facet_wrap(~gen) +
geom_line() +
theme_bw()
```

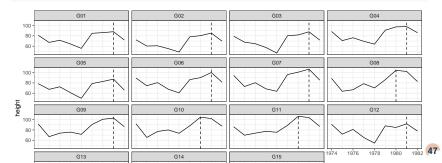






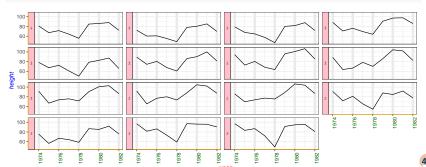


```
agridat::aastveit.barley.height %>%
  ggplot(aes(x=year, y=height)) +
  facet_wrap(~gen) +
  geom_line() +
  geom_vline(
   data=aastveit.barley.height %>%
      group_by(gen) %>%
      filter(height==max(height)),
      aes(xintercept=year), linetype="dashed") +
  theme_bw()
```





```
agridat::aastveit.barley.height %>%
  ggplot(aes(x=year, y=height)) +
  facet_wrap(-gen, strip.position = "left") +
  geom_line() +
  theme_bw() +
  theme(axis.title.x = element_text(color="red", face="bold"),
        axis.title.y = element_text(color="blue", face="italic"),
        axis.text.x = element_text(color="darkgreen", angle=90),
        strip.background = element_rect(fill="pink"),
        strip.text = element_text(size=3),
        axis.line = element_line(color="orange"),
        panel.grid.major.x = element_line(linewidth=2),
        panel.grid.minor.y = element_line(linetype="dashed"))
```



Exemplos

Conjuntos de dados do governo



https://dados.gov.br/dados/conjuntos-dados

New names:



Fonte: https://dados.gov.br/dados/conjuntos-dados/plantas-alternativas-as-plantas-exoticas-invasoras-rj

* `Nome científico` -> `Nome científico...5`

```
## * `Nativa (N) ou Exótica (E)` -> `Nativa (N) ou Exótica (E)..
## * `Nome científico` -> `Nome científico...7`
## * `Nativa (N) ou Exótica (E)` -> `Nativa (N) ou Exótica (E)..
## * `Nome científico` -> `Nome científico...9`
```

* `Nativa (N) ou Exótica (E)` -> `Nativa (N) ou Exótica (E)..
A tibble: 261 x 12

`Código de uso` `Família EEI` `Nome científico EEI`



Fonte: https://dados.gov.br/dados/conjuntos-dados/plantas-alternativas-as-plantas-exoticas-invasoras-rj

```
library(readxl)
invasoras <- read xlsx(
  paste0("../dados/listagem-de-plantas-alternativas-as-",
         "plantas-exoticas-invasoras-listadas-para-o-",
         "estado-do-rio-de.xlsx").
  col_names=c("Codigo", "Familia_EEI", "NomeCient_EEI",
              "NomePop EEI",
              "NomeCient Ombro", "NouE_Ombro",
              "NomeCient Semidec", "NouE Semidec",
              "NomeCient InfMar", "NouE InfMar",
              "Ecossist EEI", "Categoria"),
  range="A8:L268")
```

##

##



```
## # A tibble: 1,566 x 9
                 Familia_EEI NomeCient_EEI NomePop_EEI Ecossist_
##
##
      <chr>
                 <chr>
                             <chr>
                                           <chr>
                                                        <chr>
##
    1 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
    2 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
##
    3 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
##
##
    4 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
##
    5 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
```

6 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações

7 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações

8 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
9 ornamental Acanthaceae Asystasia ga~ asistásia,~ Formações
10 ornamental Acanthaceae Asystasia ga~ asistásia,~ Form53,ões

invasoras %>%



```
pivot_longer(c(ends_with("Ombro"),
                 ends_with("Semidec"),
                 ends with("InfMar"))) %>%
  separate(name, c("Var", "Ecossist")) %>%
  pivot_wider(id_cols=c(Codigo,ends_with("EEI"),Categoria,Ecossist),
              names_from=Var)
## # A tibble: 354 x 9
##
      Codigo Familia_EEI NomeCient_EEI NomePop_EEI Ecossist_EEI Catego
##
      <chr>
              <chr>>
                          <chr>>
                                        <chr>>
                                                     <chr>>
                                                                      <d
##
    1 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    2 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
    3 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
##
    4 orname~ Acanthaceae Brillantaisi~ erva-de-bi~ Floresta Om~
##
    5 orname~ Acanthaceae Brillantaisi~ erva-de-bi~ Floresta Om~
##
    6 orname~ Acanthaceae Brillantaisi~ erva-de-bi~ Floresta Om~
##
    7 orname~ Acanthaceae Thunbergia a~ amarelinha~ Floresta Om~
    8 orname~ Acanthaceae Thunbergia a~ amarelinha~ Floresta Om~
##
   9 orname~ Acanthaceae Thunbergia a~ amarelinha~ Floresta Om~
##
## 10 orname~ Acanthaceae Thunbergia g~ tumbérgia-~ Floresta Om~ 54
```

invasoras %>%



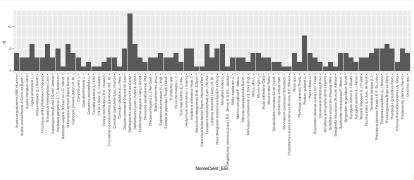
```
pivot_longer(c(ends_with("Ombro"),
                 ends_with("Semidec"),
                 ends with("InfMar"))) %>%
  separate(name, c("Var", "Ecossist")) %>%
  pivot_wider(id_cols=c(Codigo,ends_with("EEI"),Categoria,Ecossist),
              names from=Var) %>%
  unnest_longer(c(NomeCient, NouE))
## # A tibble: 783 x 9
##
     Codigo Familia_EEI NomeCient_EEI NomePop_EEI Ecossist_EEI Catego
##
      <chr> <chr>
                          <chr>>
                                        <chr>
                                                    <chr>
                                                                      <d
##
    1 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
    2 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    3 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    4 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
##
    5 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    6 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    7 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    8 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
   9 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~ 55
##
```



```
## # A tibble: 504 \times 9
##
      Codigo Familia_EEI NomeCient_EEI NomePop_EEI Ecossist_EEI Catego
##
      <chr>>
               <chr>>
                            <chr>>
                                            <chr>>
                                                         <chr>>
                                                                            <d
    1 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
##
    2 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    3 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    4 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    5 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
    6 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    7 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~
##
    8 orname~ Acanthaceae Asystasia ga~ asistásia,~ Formações P~ \[ \begin{aligned} \frac{56}{2} \end{aligned} \]
##
```

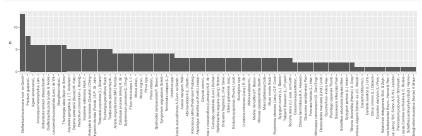


```
invasoras %>%
  group_by(NomeCient_EEI) %>%
  count() %>%
  ungroup() %>%
  ggplot(aes(x=NomeCient_EEI, y=n)) +
  geom_col() +
  theme_grey(base_size = 8) +
  theme(axis.text.x = element_text(angle=90, hjust=1, vjust=.5))
```





```
invasoras %>%
  group_by(NomeCient_EEI) %>%
  count() %>%
  ungroup() %>%
  mutate(NomeCient_EEI = fct_reorder(NomeCient_EEI, desc(n))) %>%
  ggplot(aes(x=NomeCient_EEI, y=n)) +
  geom_col() +
  theme_grey(base_size = 8) +
  theme(axis.text.x = element_text(angle=90, hjust=1, vjust=.5))
```







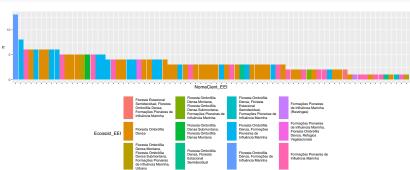


```
invasoras %>%
  group_by(NomeCient_EEI, Ecossist_EEI) %>%
  count() %>%
  group by (NomeCient EEI) %>%
  count() %>%
  arrange(desc(n))
## # A tibble: 77 x 2
## # Groups: NomeCient_EEI [77]
##
      NomeCient_EEI
                                                   n
      <chr>
                                               <int>
##
    1 Acacia angustissima (Mill.) Kuntze
##
##
   2 Acacia mangium Wild
##
    3 Dieffenbachia amoena hort, ex Gentil
    4 Elaeis guineensis Jacq.
##
##
    5 Melinis minutiflora P. Beauv.
    6 Mimusops commersonii (G. Don) Engl.
##
##
   7 Thunbergia grandiflora Roxb.
                                                   2
   8 Acacia auriculiformis A.Cunn. ex Benth.
##
```



```
ecossist <- invasoras %>%
  mutate_at(vars(Ecossist_EEI), replace_na, "?") %>%
  group_by(NomeCient_EEI, Ecossist_EEI) %>%
  count() %>%
  group_by(NomeCient_EEI) %>%
  arrange(Ecossist_EEI) %>%
  summarise(Ecossist_EEI = (function(x) {
    if(length(x)==1) return(x)
    else {
      if(is.na(x[1])) return(x[2])
      if (is.na(x[2])) return(x[1])
      if(grepl(x[1], x[2])) return(x[2])
      if(grepl(x[2], x[1])) return(x[1])
      return(paste(x, collapse=", "))
  })(Ecossist_EEI),
 n = sum(n)) %>%
  mutate_at(vars(Ecossist_EEI), ~ifelse(
    .=="Formações Pioneiras de Influência Marinha, Floresta Ombrófila D
    "Floresta Ombrófila Densa, Formações Pioneiras de Influência Marinh
```







Fonte: https://dados.gov.br/dados/conjuntos-dados/programa-fomento-rural

```
library(readr)
fomentoRural <- read csv(</pre>
  paste0("https://aplicacoes.mds.gov.br/sagi/",
         "servicos/misocial/?",
         "fg=anomes s:2023*&",
         "fq=formento_qtd_total_familias_benef_i:*&",
         "q=*:*&",
         "rows=1000000&",
         "wt=csv"))
head(fomentoRural)
```

```
## # A tibble: 6 x 1,077
     paa_qtd_agricul_familiar_modal~1 qtd_pes_pbf_idade_7_~2 bolsa_atle
##
##
     <lgl>
                                                          <dbl> <lgl>
     NA
                                                            118 NA
## 1
## 2 NA
                                                            128 NA
                                                             82 NA
## 3 NA
                                                                     63
## 4 NA
                                                            148 NA
```



```
fomentoRural %>%
  select_if(~!all(is.na(.))) %>%
  select_if(~!all(.==0))
```

```
## # A tibble: 662 x 348
##
      qtd_pes_pbf_idade_7_a_15_sexo~1 cadunico_tot_fam_ext~2 igd_pbf_qt
                                 dbl>
                                                          db1>
##
                                    118
                                                           7707
##
    1
##
                                    128
                                                             NΑ
##
                                    82
                                                           4030
                                    148
##
                                                             NA
##
    5
                                    79
                                                             NA
##
    6
                                    85
                                                           5489
##
                                     83
                                                           5522
##
    8
                                     24
                                                            756
##
                                     25
                                                           3006
## 10
                                    103
                                                           1309
## # i 652 more rows
## # i abbreviated names: 1: qtd_pes_pbf_idade_7_a_15_sexo_feminino_i,
       2: cadunico_tot_fam_ext_pob_i, 3: igd_pbf_qtd_total_publico_saud
## #
```

i 345 more variables:



```
fomentoRural %>%
  select(anomes s, matches( paste0(
    "formento\\ qtd\\ fam\\ benef\\ .*\\ i")))
## # A tibble: 662 x 15
##
      anomes_s formento_qtd_fam_ben~1 formento_qtd_fam_ben~2 for
##
         <dbl>
                                 <dbl>
                                                        <dbl>
        202310
##
    1
##
    2
        202312
    3
        202311
##
        202312
##
    4
        202312
##
    5
    6
        202310
##
##
    7
        202311
##
        202306
    8
        202308
##
##
   10
        202311
                                                             0
  # i 652 more rows
##
## # i abbreviated names:
                                                             65
```

#



```
fomentoRural %>%
  select(anomes s, matches(
    "formento\\_qtd\\_fam\\_benef\\_.*\\_i")) %>%
  group_by(anomes_s) %>%
  summarise_all(sum, na.rm=T)
## # A tibble: 10 x 15
##
      anomes_s formento_qtd_fam_ben~1 formento_qtd_fam_ben~2 formento_q
##
         <dbl>
                                <dbl>
                                                        <dbl>
        202301
##
    1
##
        202303
   3 202304
##
      202306
##
    4
##
    5
      202307
      202308
##
    6
##
    7
        202309
##
   8
        202310
##
      202311
                                    5
        202312
##
   10
##
       abbreviated names:
```

1: formento_qtd_fam_benef_catadores_material_reciclado_i,

66

select(anomes_s, matches(

fr <- fomentoRural %>%



```
"formento\\ gtd\\ fam\\ benef\\ .*\\ i")) %>%
  group_by(anomes_s) %>%
  summarise_all(sum, na.rm=T) %>%
 pivot longer(-anomes s)
head(fr)
## # A tibble: 6 x 3
##
                                                                     val
     anomes s name
##
        <dbl> <chr>
                                                                     <db
## 1
       202301 formento_qtd_fam_benef_catadores_material_reciclado_i
       202301 formento_qtd_fam_benef_comun_terreiro_i
## 2
## 3
       202301 formento_qtd_fam_benef_ciganas_i
## 4
       202301 formento_qtd_fam_benef_quilombolas_i
## 5
       202301 formento_qtd_fam_benef_assentados_ref_agraria_i
## 6
       202301 formento_qtd_fam_benef_agrilcultores_familiares_i
```



```
fr <- fr %>%
 mutate_at(vars(anomes_s), as.character) %>%
  mutate_at(vars(anomes_s), fast_strptime, format="%Y%m") %>%
  mutate_at(vars(anomes_s), as_date)
head(fr)
## # A tibble: 6 x 3
##
     anomes_s
               name
##
     <date>
               <chr>
## 1 2023-01-01 formento_qtd_fam_benef_catadores_material_reciclado_i
## 2 2023-01-01 formento_qtd_fam_benef_comun_terreiro_i
## 3 2023-01-01 formento_qtd_fam_benef_ciganas_i
## 4 2023-01-01 formento_qtd_fam_benef_quilombolas_i
## 5 2023-01-01 formento_qtd_fam_benef_assentados_ref_agraria_i
## 6 2023-01-01 formento_qtd_fam_benef_agrilcultores_familiares_i
```



```
fr <- fr %>%
 mutate(GPTE = str_remove(name, "formento\\_qtd\\_fam\\_benef\\_"),
        GPTE = str_remove(GPTE, "\\_i$")) %>%
 mutate_at(vars(GPTE), factor)
head(fr)
## # A tibble: 6 x 4
##
    anomes_s
              name
##
     <date> <chr>
## 1 2023-01-01 formento_qtd_fam_benef_catadores_material_reciclado_i
## 2 2023-01-01 formento_qtd_fam_benef_comun_terreiro_i
## 3 2023-01-01 formento_qtd_fam_benef_ciganas_i
## 4 2023-01-01 formento_qtd_fam_benef_quilombolas_i
## 5 2023-01-01 formento_qtd_fam_benef_assentados_ref_agraria_i
## 6 2023-01-01 formento_qtd_fam_benef_agrilcultores_familiares_i
```

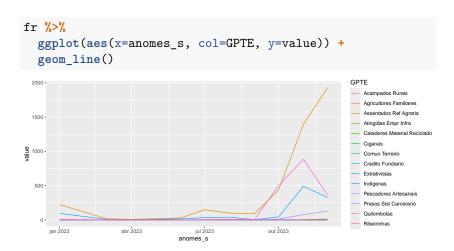


```
fr <- fr %>%
  mutate_at(vars(GPTE), fct_relabel,
            str_replace_all, "\\_", " ") %>%
  mutate_at(vars(GPTE), fct_relabel,
            str to title)
levels(fr$GPTE)
    [1] "Acampados Rurais"
                                        "Agrilcultores Familiares
##
##
    [3] "Assentados Ref Agraria"
                                        "Atingidas Empr Infra"
    [5] "Catadores Material Reciclado"
                                        "Ciganas"
##
##
    [7] "Comun Terreiro"
                                        "Credito Fundiario"
##
    [9] "Extrativistas"
                                        "Indinegas"
   [11] "Pescadores Artesanais"
                                        "Presos Sist Carcerario"
##
   [13] "Quilombolas"
                                        "Ribeirinhas"
```



```
fr <- fr %>%
  mutate at(vars(GPTE), fct relabel,
            str_replace,
            "Agrilcultores", "Agricultores") %>%
  mutate at(vars(GPTE), fct_relabel,
            str_replace,
            "Indinegas", "Indigenas")
levels(fr$GPTE)
                                        "Agricultores Familiares"
##
    [1] "Acampados Rurais"
    [3] "Assentados Ref Agraria"
##
                                        "Atingidas Empr Infra"
    [5] "Catadores Material Reciclado" "Ciganas"
##
    [7] "Comun Terreiro"
                                        "Credito Fundiario"
##
    [9] "Extrativistas"
                                        "Indígenas"
##
   [11] "Pescadores Artesanais"
                                        "Presos Sist Carcerario"
   [13] "Quilombolas"
                                        "Ribeirinhas"
##
```





Exercícios

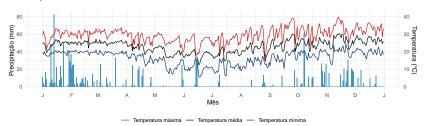
Exercício 1



Personalizar o gráfico a seguir:

```
dados_metereologicos %>%
  select(Data, PRECIPITACAO, starts_with("TEMPER")) %>%
  pivot_longer(starts_with("TEMPER")) %>%
  ggplot(aes(x=Data)) +
  geom_bar(stat="unique", aes(y=PRECIPITACAO)) +
  geom_line(aes(y=value, color=name))
```

Exemplo:



Exercício 2

1500 -



- ► Encontrar o conjunto de dados dos Cursos Ativos da Graduação da UFSCar (https://dados.gov.br/dados/conjuntos-dados/)
- ► Processar os dados
- ▶ Obter a soma de vagas para cada uma das duas modalidades

```
## Rows: 78 Columns: 6
## -- Column specification -----
## Delimiter: ","
## chr (4): Curso, Turno, Modalidade, Campus
## dbl (2): Codigo E-mec, Vagas
##
## i Use `spec()` to retrieve the full column specification for
## i Specify the column types or set `show col types = FALSE` to
    Modalidade
##
## 1
            EAD 1405
## 2 Presencial 3027
```

► Fazer um gráfico, baseando-se no gráfico a seguir: