Cardiovascular disease worsens the maternal prognosis of COVID-19

Codes and outputs

08/20/2021

Description

This file presents the documentation of the analysis of article "Cardiovascular disease worsens the maternal prognosis of COVID-19" with authors XXXXXX. ATUALIZAR AQUI COM AS QUESTÕES DOS AUTORES

R packages used, functions and dataset import

The data are analyzed using the free-software R (https://www.R-project.org) in version 4.0.3. Next, we present and load the libraries used in the data analysis process.

```
loadlibrary <- function(x) {</pre>
  if (!require(x, character.only = TRUE)) {
    install.packages(x, dependencies = T)
    if (!require(x, character.only = TRUE))
      stop("Package not found")
}
packages <-
  c(
    "dplyr",
    "MatchIt",
    "lubridate",
    "readr",
    "readxl",
    "ggplot2",
    "kableExtra",
    "tables",
    "questionr",
    "car",
    "data.table",
    "magrittr",
    "tidyverse",
    "readxl",
    "summarytools",
    "modelsummary",
    "RColorBrewer",
```

```
"zoo",
   "WeightIt",
   "jtools",
   "cobalt",
   "weights"
)
lapply(packages, loadlibrary)
```

One can see below the functions that will be used in the data analysis.

```
#functions for summary measures
media <- function(x)
  mean(x, na.rm = TRUE)
mediana <- function(x)
  median(x, na.rm = TRUE)
DP <- function(x)</pre>
  sd(x, na.rm = TRUE)
minimo <- function(x)</pre>
  base::min(x, na.rm = TRUE)
maximo <- function(x)</pre>
  base::max(x, na.rm = TRUE)
q25 <- function(x)
  stats::quantile(x, p = 0.25, na.rm = TRUE)
q75 <- function(x)
  stats::quantile(x, p = 0.75, na.rm = TRUE)
IQR <- function(x)</pre>
  round(q75(x) - q25(x), 2)
n <- function(x)</pre>
  sum(!is.na(x))
```

The Influenza Epidemiological Surveillance Information System, SIVEP-Gripe (Sistema de Informação de Vigilância Epidemiológica da Gripe), is a nationwide surveillance database used to monitor severe acute respiratory infections in Brazil.

Notification is mandatory for Influenza Syndrome (characterized by at least two of the following signs and symptoms: fever, even if referred, chills, sore throat, headache, cough, runny nose, olfactory or taste disorders) and who has dyspnea/respiratory discomfort or persistent pressure in the chest or O2 saturation less than 95% in room air or bluish color of the lips or face. Asymptomatic individuals with laboratory confirmation by molecular biology or immunological examination for COVID-19 infection are also reported.

For notifications in Sivep-Gripe, hospitalized cases in both public and private hospitals and all deaths due to severe acute respiratory infections regardless of hospitalization must be considered.

The search was limited to the first notified case of COVID-19 in February 2020 until the 15th epidemiological week of 2021 (up to April 17, 2021). The datasets were obtained on May 5, 2021, on the site https://opendatasus.saude.gov.br/dataset. The first period (8th to 53rd epidemiological week of 2020) and the second period (1st to 15th epidemiological week of 2021) datasets can be obtained at https://drive.google.com/file/d/1jts4h0ovdwFh86SdKyslMLSG9rOy3UjX/view?usp=sharing and at https://drive.google.com/file/d/1gQSy_dcUkd1UrDEcsrDbyGl4gEvcI8z_/view?usp=sharing, respectively. The data are loaded below:

```
";",
  escape_double = FALSE,
  locale = locale(encoding = "ISO-8859-2"),
  trim ws = TRUE
#2020
data_2020 <- read_delim(
  "INFLUD-03-05-2021.csv",
  ";",
  escape_double = FALSE,
  locale = locale(encoding = "ISO-8859-2"),
  trim_ws = TRUE
sem <- 15 #limit of epidemiological week of 2021
#memory.limit(999999)
#### Concatenating 2020 and 2021 data #############
data_all <- rbind(data_2020, data_2021)</pre>
#Creating the case year variable
data_all <- data_all %>%
  dplyr::mutate(
    dt_sint = as.Date(DT_SIN_PRI, format = "%d/%m/%Y"),
    year_case = lubridate::year(dt_sint),
  )
```

There are 1905854 observations. The case selection is presented in the following according to the flowchart presented in the article.

Case selection and data treatment

The first filter consists of selecting the cases from 8th epidemiological week of 2020 to 15th epidemiological week of 2021.

```
#Cases from the 8th epidemiological week of 2020 to 15th of 2021
data1 <- data_all %>%
filter((year_case==2020 & SEM_PRI >=8) | year_case ==2021)
```

There are 12563 cases in 2021 in epidemiological week 53 of 2020. These are cases from the first two days of 2021, which are still part of the last epidemiological week of 2020 (http://portalsinan.saude.gov.br/calendario-epidemiologico?layout=edit&id=168). However, these cases belong to the 53rd week of 2020 and we corrected as follows:

```
data1 <- data1 %>%
mutate(year_case = ifelse(year_case ==2021 & SEM_PRI ==53, 2020, year_case)) %>%
filter(year_case==2020 | (year_case ==2021 & SEM_PRI <= sem))</pre>
```

There are 1876953 observations.

The next selection consists of selecting only confirmed cases of COVID-19:

```
##Filtering confirmed cases of COVID-19
data2 <- data1 %>%
    filter(CLASSI_FIN == 5)
```

There are 1174350 observations.

The next step is to identify hospitalized cases (HOSPITAL=1) and then select only those cases.

```
#Filtering only hospitalized cases
data3 <- data2 %>%
filter(HOSPITAL == 1)
```

Now we have 1114042 observations.

The next selection consists of selecting females cases:

```
##Filtering female notifications
data4 <- filter(data3, CS_SEX0 == "F")</pre>
```

There are 495698 observations.

The next selection consists of selecting women between 10 and 55 years old.

```
#Filtering women aged 10-55
data5 <- data4 %>%
filter(NU_IDADE_N > 9 & NU_IDADE_N <= 55)</pre>
```

There are 174525 observations.

The next step is to identify pregnant and postpartum people (variable classi_gesta_puerp) and then select only those cases.

```
#Creating the classification variable if pregnant, postpartum and not (neither pregnant nor postpartum)
data5 <- data5 %>%
  mutate(
    classi_gesta_puerp = case_when(
      CS_GESTANT == 1 ~ "1tri", #1st trimester
     CS_GESTANT == 2 ~ "2tri", #2nd trimester
      CS_GESTANT == 3 ~ "3tri", #3rd trimester
     CS_GESTANT == 4 ~ "IG_ig", #ignored gestational Age
      CS_GESTANT == 5 &
       PUERPERA == 1 ~ "puerp", #puerperium
     CS_GESTANT == 9 & PUERPERA == 1 ~ "puerp", #puerperium
     TRUE ~ "no" #neither pregnant nor postpartum
   )
  )
#Filtering only pregnant and postpartum women
data6 <- data5 %>%
 filter(classi_gesta_puerp != "no")
```

There are 10635 observations.

The variable identified as EVOLUCAO is categorized with 1 - cure, 2 - death by SARS, 3 - death by other cause, 9 - ignored.

```
with(data6, freq(EVOLUCAO))
```

```
## Frequencies
## data6$EVOLUCAO
## Type: Numeric
##
##
                           % Valid
                                      % Valid Cum.
                                                       % Total
                                                                  % Total Cum.
                   Freq
##
                                                                          75.27
##
              1
                   8005
                             85.38
                                              85.38
                                                         75.27
              2
                     979
                             10.44
                                              95.82
                                                          9.21
                                                                         84.48
##
##
              3
                     15
                              0.16
                                              95.98
                                                          0.14
                                                                         84.62
##
              9
                     377
                              4.02
                                             100.00
                                                          3.54
                                                                         88.16
##
           <NA>
                   1259
                                                         11.84
                                                                        100.00
##
                  10635
                            100.00
                                             100.00
                                                        100.00
                                                                        100.00
         Total
```

Now we are going to select only the finalized cases:

Frequencies

```
#Filtering only finalized cases
data7 <- data6 %>%
  filter((EVOLUCA0 == 1 | EVOLUCA0 == 2 | EVOLUCA0 == 3) & !is.na(EVOLUCA0))

#Creating the evolution variable
data7 <- data7 %>%
  mutate(evolution = case_when(
    EVOLUCA0 == 1 ~ "cure",
    EVOLUCA0 == 2 ~ "death",
    EVOLUCA0 == 3 ~ "death"
))

with(data7, freq(evolution))
```

```
## data7$evolution
## Type: Character
##
##
                          % Valid
                                     % Valid Cum.
                                                      % Total
                                                                % Total Cum.
##
##
                  8005
                            88.95
                                             88.95
                                                        88.95
                                                                        88.95
          cure
                   994
                                                                       100.00
##
         death
                            11.05
                                            100.00
                                                        11.05
##
           <NA>
                     0
                                                         0.00
                                                                       100.00
##
                  8999
                           100.00
                                            100.00
                                                       100.00
                                                                       100.00
         Total
```

There are 8999 confirmed and hospitalized cases of COVID-19 in pregnant and postpartum women aged 10 to 55 years. We identify 60.42% of these notifications without any information about the presence or absence of cardiovascular diseases (CVD). The next selection consists of selecting cases where the CVD variable (CARDIOPATI) indicates yes or no:

```
with(data7, freq(CARDIOPATI))
## Frequencies
## data7$CARDIOPATI
## Type: Numeric
##
##
               Freq % Valid % Valid Cum. % Total % Total Cum.
                      16.55 16.55
81.39 97.94
2.06 100.00
##
          1 602 16.55
                                              6.69
                                                            6.69
         2 2960 81.39
9 75 2.06
                                             32.89
                                                          39.58
                                                          40.42
##
                                             0.83
        <NA> 5362
                                             59.58
                                                         100.00
               8999 100.00 100.00 100.00
       Total
                                                          100.00
#Creating the CVD variable
data7 <- data7 %>%
 mutate(CVD = case_when(
   CARDIOPATI == 1 ~ "yes",
   CARDIOPATI == 2 ~ "no",
   TRUE ~ NA_character_
```

Now we present the final dataset.

```
#Filtering cases with information about CVD
data <- data7 %>%
filter(!is.na(CVD))
```

```
questionr::freq(
  data$CVD,
    cum = FALSE,
  total = TRUE,
  na.last = FALSE,
  valid = FALSE
) %>%
  kable(caption = "Frequency of CVD", digits = 2) %>%
  kable_styling(latex_options = "HOLD_position")
```

Table 1: Frequency of CVD

	n	%
no	2960	83.1
yes	602	16.9
Total	3562	100.0

Epidemiologic characteristics

Ethnicity

```
# Ethnicity
data <- data %>%
 mutate(
   ethnicity = case when(
     CS_RACA == 1 ~ "white",
     CS_RACA == 2 ~ "black",
     CS_RACA == 3 ~ "yellow",
     CS RACA == 4 ~ "brown",
     CS_RACA == 5 ~ "indigenous",
     TRUE ~ NA_character_
   )
 )
with(data, ctable(ethnicity, CVD, prop = "c", useNA = "no", chisq = FALSE))
## Cross-Tabulation, Column Proportions
## ethnicity * CVD
## Data Frame: data
##
## ---
                                           yes
##
               CVD
                              no
                                                         Total
##
   ethnicity
       black
brown
                   167 ( 6.5%) 46 ( 9.1%) 213 ( 6.9%)
##
##
                    1375 (53.5%) 244 (48.1%) 1619 (52.6%)
                    30 ( 1.2%) 1 ( 0.2%) 31 ( 1.0%)
## indigenous
                     975 ( 37.9%) 214 ( 42.2%) 1189 ( 38.6%)
##
        white
                      24 ( 0.9%)
##
        yellow
                                    2 ( 0.4%)
                                                   26 ( 0.8%)
##
                    2571 (100.0%) 507 (100.0%) 3078 (100.0%)
        Total
fisher.test(data$ethnicity, data$CVD)
##
## Fisher's Exact Test for Count Data
## data: data$ethnicity and data$CVD
## p-value = 0.007096
## alternative hypothesis: two.sided
```

Education

```
data$education <- factor(data$education,</pre>
levels = c("No schooling","1st to 5th grade", "6th to 9th grade",
"Middle school", "Superior"))
with(data, ctable(education, CVD, prop = "c", useNA = "no", chisq = FALSE))
## Cross-Tabulation, Column Proportions
## education * CVD
## Data Frame: data
##
                     CVD
                                                               Total
##
         education
      No schooling
                            11 ( 0.7%)
                                          2 ( 0.7%)
                                                        13 ( 0.7%)
##
##
   1st to 5th grade
                          123 ( 7.9%) 33 ( 12.2%) 156 ( 8.5%)
## 6th to 9th grade
                          307 (19.7%) 52 (19.2%) 359 (19.6%)
##
       Middle school
                          857 (55.0%) 140 (51.7%) 997 (54.5%)
                           260 ( 16.7%) 44 ( 16.2%)
                                                      304 ( 16.6%)
##
           Superior
              Total
##
                          1558 (100.0%) 271 (100.0%)
                                                       1829 (100.0%)
```

fisher.test(data\$education, data\$CVD)

```
##
## Fisher's Exact Test for Count Data
##
## data: data$education and data$CVD
## p-value = 0.2405
## alternative hypothesis: two.sided
```

Age (Years)

	n	media	DP	mediana	minimo	maximo	q25	q75	IQR
no	2960.00	30.00	7.43	30.00	13.00	55.00	24.00	35.00	11.00
yes	602.00	34.76	8.52	34.00	15.00	55.00	29.00	39.75	10.75

```
t.test(NU_IDADE_N ~ CVD, data = data)
```

```
##
## Welch Two Sample t-test
##
## data: NU_IDADE_N by CVD
## t = -12.745, df = 797.4, p-value < 2.2e-16
## alternative hypothesis: true difference in means is not equal to 0</pre>
```

```
## 95 percent confidence interval:
## -5.485124 -4.021042
## sample estimates:
## mean in group no mean in group yes
## 30.00439 34.75748
```

Age group

```
data <- data %>%
 mutate(
   age_group = case_when(
    NU_IDADE_N <= 19 ~ "<20",
    NU IDADE N >= 20
    & NU_IDADE_N <= 34 ~ "20-34",
    NU_IDADE_N > 34 ~ ">=35",
    TRUE ~ NA_character_
   )
 )
data$age_group <-
 factor(data$age_group, levels = c("<20", "20-34", ">=35"))
with(data, ctable(age_group, CVD, prop = "c", useNA = "no", chisq = FALSE))
## Cross-Tabulation, Column Proportions
## age_group * CVD
## Data Frame: data
##
## ----- ---- ----
##
            CVD
                                       yes
                                                    Total
                           no
  age_group
                  237 ( 8.0%) 10 ( 1.7%) 247 ( 6.9%)
##
       <20
                 1895 ( 64.0%) 292 ( 48.5%) 2187 ( 61.4%)
      20-34
                   828 ( 28.0%) 300 ( 49.8%) 1128 ( 31.7%)
##
       >=35
      Total 2960 (100.0%) 602 (100.0%) 3562 (100.0%)
```

```
with(data,fisher.test(age_group, CVD))
```

```
##
## Fisher's Exact Test for Count Data
##
## data: age_group and CVD
## p-value < 2.2e-16
## alternative hypothesis: two.sided</pre>
```

Residence area

```
data <- data %>%
 mutate(zone = case_when(CS_ZONA ==1 ~ "Urban",
                           CS_ZONA == 2 ~ "Rural",
                           CS_ZONA == 3 ~ "Periurban",
                            TRUE ~ NA_character_))
with(data, ctable( zone, CVD, prop = "c", useNA = "no", chisq = FALSE))
## Cross-Tabulation, Column Proportions
## zone * CVD
## Data Frame: data
               CVD
##
                              no
                                            yes
                                                         Total
##
         zone
                     10 ( 0.4%) 3 ( 0.5%) 13 ( 0.4%)
##
   Periurban
                    159 ( 5.7%) 24 ( 4.4%) 183 ( 5.5%)
##
       Rural
                   2618 (93.9%) 522 (95.1%) 3140 (94.1%)
##
        Urban
                    2787 (100.0%) 549 (100.0%) 3336 (100.0%)
        Total
fisher.test(data$zone, data$CVD)
##
## Fisher's Exact Test for Count Data
## data: data$zone and data$CVD
## p-value = 0.3673
## alternative hypothesis: two.sided
```

Gestational moment

```
ctable(data$classi_gesta_puerp, data$CVD, prop="c", OR=TRUE, useNA = "no", chisq = TRUE)
## Cross-Tabulation, Column Proportions
## classi_gesta_puerp * CVD
## Data Frame: data
##
##
  ##
                   CVD
                                            yes
                                                        Total
                                 no
##
   classi_gesta_puerp
                         161 ( 5.4%) 34 ( 5.6%) 195 ( 5.5%)
##
              1tri
##
                         471 ( 15.9%) 120 ( 19.9%) 591 ( 16.6%)
               2tri
              3tri
                        1199 ( 40.5%) 248 ( 41.2%) 1447 ( 40.6%)
##
                         84 ( 2.8%) 30 ( 5.0%)
##
             IG_ig
                                                 114 ( 3.2%)
                        1045 ( 35.3%) 170 ( 28.2%) 1215 ( 34.1%)
##
             puerp
                       2960 (100.0%) 602 (100.0%) 3562 (100.0%)
             Total
```

```
## ## Chi.squared df p.value ## ------ ## 19.4898 4 6e-04 ## -----
```

Hospital-acquired infection

```
data <- data %>%
 mutate(hospital_infection = case_when(NOSOCOMIAL ==1 ~ "yes",
                          NOSOCOMIAL == 2 ~ "no",
                          TRUE ~ NA_character_))
with(data, ctable( hospital_infection, CVD, prop = "c", useNA = "no", chisq = FALSE))
## Cross-Tabulation, Column Proportions
## hospital_infection * CVD
## Data Frame: data
##
                      CVD no yes
##
                                                                Total
   {\tt hospital\_infection}
##
                           2551 ( 98.0%) 448 ( 98.0%) 2999 ( 98.0%)
                             53 ( 2.0%) 9 ( 2.0%)
##
                 yes
                                                         62 ( 2.0%)
                           2604 (100.0%) 457 (100.0%) 3061 (100.0%)
fisher.test(data$hospital_infection, data$CVD)
```

```
##
## Fisher's Exact Test for Count Data
##
## data: data$hospital_infection and data$CVD
## p-value = 1
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.4163122 1.9955578
## sample estimates:
## odds ratio
## 0.9669453
```

Symptoms

Fever

```
data <- data %>%
 mutate(fever = case_when(FEBRE == 1 ~ "yes",
                   FEBRE == 2 \sim "no",
                   TRUE ~ NA_character_))
with(data, ctable(CVD, fever, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * fever
## Data Frame: data
##
## ----- ----- ------ ------
       fever
                  no
                                    Total
##
                           yes
##
    CVD
            1289 (44.2%) 1625 (55.8%) 2914 (100.0%)
##
   no
##
             209 (38.3%) 337 (61.7%)
                                546 (100.0%)
   yes
            1498 (43.3%) 1962 (56.7%) 3460 (100.0%)
##
  Total
## ----- -----
##
## -----
## Chi.squared df p.value
## -----
  6.4048
          1 0.0114
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.28
           1.06
                  1.54
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
   1.16
         1.16 1.16
## -----
Cough
```

```
## Data Frame: data
##
##
## ----- ----- ------ ------
##
       cough
                   no
                             yes
                                      Total
##
   CVD
##
    no
            1002 (34.4%) 1913 (65.6%) 2915 (100.0%)
   yes
             138 (24.6%) 424 (75.4%)
##
                                 562 (100.0%)
          1140 (32.8%) 2337 (67.2%) 3477 (100.0%)
##
   Total
## ----
 Chi.squared df p.value
## -----
   20.1693 1 0
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
   1.61
        1.31
                 1.98
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.40
           1.40
                 1.40
```

Sore throat

```
data <- data %>%
 mutate(sore_throat = case_when(GARGANTA == 1 ~ "yes",
                         GARGANTA == 2 ~ "no",
                         TRUE ~ NA_character_))
with(data, ctable(CVD, sore_throat, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * sore_throat
## Data Frame: data
##
##
## ----- ----- ----- ------ ------
                              no
                                         yes
##
          sore_throat
##
     CVD
                      2245 (78.5%) 616 (21.5%)
                                              2861 (100.0%)
##
     no
##
                      367 (74.7%) 124 (25.3%)
                                              491 (100.0%)
     yes
                     2612 (77.9%) 740 (22.1%) 3352 (100.0%)
## -----
##
```

```
## Chi.squared df p.value
## -----
   3.1648
        1 0.0752
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
   1.23
         0.99
##
               1.54
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
   1.05 1.05 1.05
##
```

Dyspnea

```
with(data, ctable(CVD, dyspnea, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
```

```
## Cross-Tabulation, Row Proportions
## CVD * dyspnea
## Data Frame: data
##
##
## ----- -----
      dyspnea no yes
                                   Total
   CVD
##
##
          1361 (46.7%) 1553 (53.3%) 2914 (100.0%)
   no
            154 (27.7%) 401 (72.3%) 555 (100.0%)
##
   yes
            1515 (43.7%) 1954 (56.3%) 3469 (100.0%)
  Total
## ----- -----
## -----
## Chi.squared df p.value
## -----
   67.3439
         1
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
    2.28 1.87 2.79
```

```
## ## ------ Lo - 0% Hi - 0% ## ------ ## 1.68 1.68 1.68 ## -----
```

Respiratory distress

```
data <- data %>%
 mutate(resp_dist = case_when(DESC_RESP == 1 ~ "yes",
                    DESC_RESP == 2 ~ "no",
                    TRUE ~ NA_character_))
with(data, ctable(CVD, resp_dist, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * resp_dist
## Data Frame: data
##
## ----- -----
       resp_dist
                                yes
                                         Total
                      no
##
    CVD
                1541 (53.1%) 1361 (46.9%) 2902 (100.0%)
##
    no
                213 (40.3%) 316 (59.7%) 529 (100.0%)
##
    yes
               1754 (51.1%) 1677 (48.9%) 3431 (100.0%)
  Total
## ----- -----
##
 -----
 Chi.squared df p.value
## -----
   28.9948 1
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.68
           1.39
                   2.03
## -----
##
## Risk Ratio Lo - 0% Hi - 0%
##
   1.32
           1.32
                 1.32
## -----
```

Saturation

```
data <- data %>%
 mutate(saturation = case_when(SATURACAO == 1 ~ "yes",
                    SATURACAO == 2 ~ "no",
                    TRUE ~ NA_character_))
with(data, ctable(CVD, saturation, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * saturation
## Data Frame: data
##
## ----- ----- ------
       saturation no
##
                                 yes
                                          Total
##
    CVD
              1873 (65.1%) 1003 (34.9%) 2876 (100.0%)
##
    no
##
                 231 (43.4%) 301 (56.6%)
                                    532 (100.0%)
    yes
                2104 (61.7%) 1304 (38.3%) 3408 (100.0%)
##
  Total
## ----- ----- -----
##
## -----
## Chi.squared df p.value
## -----
  88.612
          1
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
    2.43
           2.02
                   2.94
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.50
          1.50 1.50
## -----
```

Diarrhea

```
## Data Frame: data
##
##
## ----- ----- -----
##
       diarrhea
                    no
                            yes
                                      Total
##
   CVD
##
             2557 (88.8%) 324 (11.2%) 2881 (100.0%)
   no
              421 (87.0%) 63 (13.0%)
##
   yes
                                484 (100.0%)
            2978 (88.5%) 387 (11.5%) 3365 (100.0%)
##
  Total
## ----
## Chi.squared df p.value
## -----
  1.1081 1 0.2925
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
          0.88
##
   1.18
                1.58
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.02
          1.02
                1.02
```

Vomit

```
data <- data %>%
 mutate(vomit = case_when(VOMITO == 1 ~ "yes",
                          VOMITO == 2 ~ "no",
                          TRUE ~ NA_character_))
with(data, ctable(CVD, vomit, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * vomit
## Data Frame: data
##
##
                         no
                                     yes
##
          vomit
##
     CVD
                2589 (90.2%) 282 (9.8%) 2871 (100.0%)
##
     no
##
                  423 (87.9%) 58 (12.1%)
                                          481 (100.0%)
     yes
              3012 (89.9%) 340 (10.1%) 3352 (100.0%)
## -----
##
```

```
## Chi.squared df p.value
## -----
##
   2.021
         1 0.1551
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
         0.93
##
   1.26
               1.70
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
   1.03 1.03 1.03
##
```

Abdominal pain

```
data <- data %>%
 mutate(abd_pain = case_when(DOR_ABD == 1 ~ "yes",
                     DOR\_ABD == 2 \sim "no",
                     TRUE ~ NA_character_))
with(data, ctable(CVD, abd_pain, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * abd_pain
## Data Frame: data
##
##
## ----- -----
        abd_pain no yes
                                          Total
    CVD
##
##
                1877 (91.1%) 183 (8.9%) 2060 (100.0%)
    no
##
                291 (89.0%) 36 (11.0%) 327 (100.0%)
    yes
               2168 (90.8%) 219 (9.2%) 2387 (100.0%)
   Total
## ----- -----
 _____
## Chi.squared df p.value
## -----
##
    1.2858
           1
               0.2568
## ------
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
##
   1.27 0.87 1.85
```

```
## ## Risk Ratio Lo - 0% Hi - 0% ## ------ ## 1.02 1.02 1.02
```

Fatigue

```
data <- data %>%
 mutate(fatigue = case_when(FADIGA == 1 ~ "yes",
                    FADIGA == 2 \sim "no",
                    TRUE ~ NA_character_))
with(data, ctable(CVD, fatigue, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * fatigue
## Data Frame: data
##
## ----- ----- ----- -----
        fatigue
                      no
                               yes
                                         Total
##
    CVD
              1609 (77.8%) 460 (22.2%) 2069 (100.0%)
##
    no
               238 (69.6%) 104 (30.4%)
                                   342 (100.0%)
##
   yes
          1847 (76.6%) 564 (23.4%) 2411 (100.0%)
  Total
## ----- -----
##
## -----
## Chi.squared df p.value
## -----
   10.4972 1 0.0012
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.53
            1.19
                   1.97
## -----
##
## Risk Ratio Lo - 0% Hi - 0%
##
   1.12
           1.12
                  1.12
```

Loss of smell

```
data <- data %>%
 mutate(loss_smell = case_when(PERD_OLFT == 1 ~ "yes",
                    PERD_OLFT == 2 ~ "no",
                    TRUE ~ NA_character_))
with(data, ctable(CVD, loss_smell, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * loss_smell
## Data Frame: data
##
## ----- ----- -----
        loss_smell no
##
                                 yes
                                          Total
##
    CVD
                1683 (81.4%) 384 (18.6%) 2067 (100.0%)
##
    no
##
                 250 (74.2%) 87 (25.8%)
                                    337 (100.0%)
    yes
                1933 (80.4%) 471 (19.6%) 2404 (100.0%)
##
   Total
## ----- -----
##
##
## Chi.squared df p.value
## -----
   9.1828
           1 0.0024
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.53
            1.17
                   1.99
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.10
          1.10 1.10
```

Loss of taste

```
## Data Frame: data
##
##
 ##
                     no
##
       loss_taste
                             yes
                                     Total
   CVD
##
              1694 (82.1%) 369 (17.9%) 2063 (100.0%)
##
    no
               250 (74.0%) 88 (26.0%)
##
    yes
                                338 (100.0%)
##
  Total
               1944 (81.0%) 457 (19.0%)
                               2401 (100.0%)
##
##
##
 Chi.squared df p.value
## -----
   11.9908 1 5e-04
## -----
##
 -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
##
   1.62
          1.24
               2.11
## -----
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.11
         1.11
               1.11
```

Any respiratory symptom

CVD * resp_symp

Now we are going to create the variable resp_symp indicating the presence of at least one respiratory symptom (Dyspnea, Fadigue, Saturation, Respiratory distress). Empty cells are considered missing data (<NA>). The variable qt_resp_symp indicates the number of respiratory symptoms of each case.

```
## Data Frame: data
##
##
## ----- ----- -----
##
                                    yes
                                              Total
        resp_symp
                        no
##
   CVD
                  935 (31.9%) 1992 (68.1%) 2927 (100.0%)
##
    no
                             484 (83.7%) 578 (100.0%)
##
   yes
                   94 (16.3%)
                 1029 (29.4%) 2476 (70.6%) 3505 (100.0%)
##
   Total
## ----
  Chi.squared df p.value
## -----
    56.4759 1 0
##
 -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
    2.42
         1.91
                   3.05
## -----
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
     1.96
            1.96
                    1.96
data <- data %>%
 mutate(qt_resp_symp = case_when(qt_sintomas_resp_aux == 4 ~ "4 symptoms",
                         qt_sintomas_resp_aux == 3 ~ "3 symptoms",
                         qt_sintomas_resp_aux == 2 ~ "2 symptoms",
                         qt_sintomas_resp_aux == 1 ~ "1 symptom",
                         qt_sintomas_resp_aux == 0 ~ "No respiratory symptom",
                       TRUE ~ NA_character_))
with(data, ctable(qt_resp_symp, CVD, prop="c", useNA = "no", chisq = TRUE))
## Cross-Tabulation, Column Proportions
## qt_resp_symp * CVD
## Data Frame: data
##
##
## ----- ---- ----
##
                     CVD
                                             yes
                                                       Total
                                  no
##
           qt_resp_symp
                           640 ( 21.9%) 116 ( 20.1%)
##
                                                756 ( 21.6%)
            1 symptom
##
            2 symptoms
                          528 ( 18.0%) 151 ( 26.1%) 679 ( 19.4%)
                          615 ( 21.0%) 164 ( 28.4%)
                                                779 ( 22.2%)
##
            3 symptoms
                          209 ( 7.1%) 53 ( 9.2%)
##
            4 symptoms
                                                  262 ( 7.5%)
## No respiratory symptom
                          935 (31.9%) 94 (16.3%) 1029 (29.4%)
                         2927 (100.0%) 578 (100.0%) 3505 (100.0%)
         Total
## ------ ---- ----
```

```
## ## ------ df p.value ## ------ ## 71.8703 4 0
```

Any symptom

```
df <- select(data, dyspnea, fatigue, saturation, resp_dist, fever, cough, sore_throat, diarrhea, vomit,
soma <- function(x){</pre>
 if (sum(is.na(x)) == 12)
   return(NA_character_)
   return(sum(!is.na(x) & x=="yes"))
data$qt_symp_aux <- apply(df,1,soma)</pre>
data <- data %>%
 mutate(symp = case_when(qt_symp_aux >= 1 ~ "yes",
                       qt_symp_aux == 0 ~ "no",
                       TRUE ~ NA_character_))
with(data, ctable(CVD, symp, prop="r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * symp
## Data Frame: data
##
## ----- -----
                       no
##
          \operatorname{\mathtt{symp}}
                                     yes
                                                 Total
##
      CVD
##
                256 (8.7%) 2685 (91.3%) 2941 (100.0%)
      no
                 22 (3.7%) 574 (96.3%)
      yes
                                          596 (100.0%)
                278 (7.9%) 3259 (92.1%) 3537 (100.0%)
##
    Total
##
##
  Chi.squared df p.value
                    0
##
    16.513
              1
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
                1.59
```

```
## Risk Ratio Lo - 0% Hi - 0%
## ------
## 2.36 2.36 2.36
## -----
```

Comorbities

```
#Hematologic
data <- data %>%
mutate(hematologic = case when(HEMATOLOGI == 1 ~ "yes",
                              HEMATOLOGI == 2 ~ "no",
                              TRUE ~ NA character ))
#Immunodepression
data <- data %>%
mutate(imunodepre = case_when(IMUNODEPRE == 1 ~ "yes",
                              IMUNODEPRE == 2 ~ "no",
                              TRUE ~ NA_character_))
#Hepatic
data <- data %>%
mutate(hepatic = case_when(HEPATICA == 1 ~ "yes",
                            HEPATICA== 2 ~ "no",
                            TRUE ~ NA_character_))
#Asthma
data <- data %>%
mutate(asthma = case_when(ASMA == 1 ~ "yes",
                        ASMA == 2 \sim "no",
                        TRUE ~ NA character ))
#Diabetes
data <- data %>%
mutate(diabetes = case_when(DIABETES == 1 ~ "yes",
                            DIABETES == 2 ~ "no",
                            TRUE ~ NA_character_))
#Neurologic
data <- data %>%
mutate(neurologic = case_when(NEUROLOGIC == 1 ~ "yes",
                              NEUROLOGIC == 2 ~ "no",
                              TRUE ~ NA_character_))
#Pneumologic
data <- data %>%
mutate(pneumologic = case_when(PNEUMOPATI == 1 ~ "yes",
                              PNEUMOPATI == 2 ~ "no",
                              TRUE ~ NA_character_))
#Renal
data <- data %>%
mutate(renal = case_when(RENAL == 1 ~ "yes",
                         RENAL == 2 \sim "no",
                         TRUE ~ NA_character_))
#Obesity
data <- data %>%
mutate(obesity = case_when(OBESIDADE == 1 ~ "yes",
                              OBESIDADE == 2 ~ "no",
                              TRUE ~ NA_character_))
```

Hematologic

```
with(data, ctable(CVD, hematologic, prop = "r", useNA = "no", chisq = TRUE, OR=TRUE))
## Cross-Tabulation, Row Proportions
## CVD * hematologic
## Data Frame: data
##
## ------ ------
##
        hematologic
                        no yes
                                           Total
##
    CVD
                  2919 (98.9%) 31 (1.1%) 2950 (100.0%)
##
    no
##
    yes
                  428 (97.9%)
                            9 (2.1%)
                                     437 (100.0%)
##
                  3347 (98.8%) 40 (1.2%)
                                     3387 (100.0%)
   Total
##
## -----
 Chi.squared df p.value
    2.5101
          1 0.1131
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.98
            0.94
                    4.19
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.01
           1.01
```

Diabetes

```
2610 (88.6%)
                     337 (11.4%)
                            2947 (100.0%)
##
   no
##
             312 (66.4%) 158 (33.6%)
                            470 (100.0%)
   yes
             2922 (85.5%) 495 (14.5%) 3417 (100.0%)
##
  Total
## ----- ----- -----
## -----
 Chi.squared df p.value
## -----
##
  159.2143
         1
##
##
## -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
   3.92
         3.14
               4.90
## -----
##
 -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.33
        1.33
             1.33
## -----
```

Obesity

```
ctable(data$CVD, data$obesity, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")
```

```
## Cross-Tabulation, Row Proportions
## CVD * obesity
## Data Frame: data
##
##
## ----- ----- ------ ------
##
        obesity
                      no
                               yes
                                         Total
##
    CVD
    no
              2651 (91.2%) 255 (8.8%)
                                  2906 (100.0%)
##
                344 (76.6%) 105 (23.4%)
                                   449 (100.0%)
    yes
              2995 (89.3%) 360 (10.7%) 3355 (100.0%)
##
   Total
## ----- ---- ----- ----- ----- -----
##
## -----
  Chi.squared df p.value
 _____
##
   85.149
           1
##
##
 _____
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
    3.17
            2.46
                    4.09
##
```

Asthma

```
ctable(data$CVD, data$asthma, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")
## Cross-Tabulation, Row Proportions
## CVD * asthma
## Data Frame: data
##
##
## ----- -----
##
                            yes
       asthma
                                       Total
                    no
    CVD
##
             2734 (93.2%) 199 (6.8%) 2933 (100.0%)
##
    no
    yes
              396 (90.8%)
                        40 (9.2%)
##
                                 436 (100.0%)
##
  Total
             3130 (92.9%) 239 (7.1%) 3369 (100.0%)
## ----- ----- -----
##
## Chi.squared df p.value
## -----
           1 0.0866
##
   2.9356
## -----
##
## Odds Ratio Lo - 95% Hi - 95%
##
   1.39
           0.97
                   1.98
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.03
           1.03
## -----
```

Hepatic

```
ctable(data$CVD, data$hepatic, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")

## Cross-Tabulation, Row Proportions
## CVD * hepatic
## Data Frame: data
##
```

```
##
## ----- -----
##
      hepatic
                 no
                      yes
                              Total
   CVD
##
##
   no
           2921 ( 99.6%) 12 (0.4%) 2933 (100.0%)
           426 ( 98.8%) 5 (1.2%)
                          431 (100.0%)
##
  yes
          3347 (99.5%) 17 (0.5%) 3364 (100.0%)
 Total
## ----- -----
##
## -----
 Chi.squared df p.value
## -----
       1 0.0912
  2.8535
## ------
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
   2.86
         1.00
              8.15
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
  1.01
        1.01
             1.01
## -----
```

fisher.test(data\$hepatic, data\$CVD)

```
##
## Fisher's Exact Test for Count Data
##
## data: data$hepatic and data$CVD
## p-value = 0.05627
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.7841874 8.7612195
## sample estimates:
## odds ratio
## 2.855734
```

Neurologic

```
ctable(data$CVD, data$neurologic, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")

## Cross-Tabulation, Row Proportions
## CVD * neurologic
## Data Frame: data
##
##
##
##
```

```
##
       neurologic
                                  Total
             no yes
##
   CVD
             2915 (99.0%) 28 (1.0%)
##
   no
                            2943 (100.0%)
              422 (97.9%)
                      9 (2.1%)
                             431 (100.0%)
##
   yes
              3337 (98.9%) 37 (1.1%)
  Total
                            3374 (100.0%)
## ----- ----- -----
## -----
## Chi.squared df p.value
 -----
   3.4923 1 0.0617
## -----
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
   2.22
         1.04
##
                4.74
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
         1.01
              1.01
   1.01
## -----
```

fisher.test(data\$neurologic, data\$CVD)

```
##
## Fisher's Exact Test for Count Data
##
## data: data$neurologic and data$CVD
## p-value = 0.04511
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.9148775 4.8799321
## sample estimates:
## odds ratio
## 2.219605
```

Pneumologic

```
ctable(data$CVD, data$pneumologic, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")
## Cross-Tabulation, Row Proportions
## CVD * pneumologic
```

```
2905 (98.9%)
                       31 (1.1%) 2936 (100.0%)
##
    no
##
               424 (97.0%) 13 (3.0%) 437 (100.0%)
   yes
##
               3329 (98.7%) 44 (1.3%) 3373 (100.0%)
## ----- ----- -----
## -----
 Chi.squared df p.value
## -----
         1 0.0021
##
   9.4404
##
##
 Odds Ratio Lo - 95% Hi - 95%
## -----
   2.87
         1.49
## -----
##
 -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.02
         1.02
              1.02
## -----
```

Immunodepression

```
ctable(data$CVD, data$imunodepre, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")
```

```
## Cross-Tabulation, Row Proportions
## CVD * imunodepre
## Data Frame: data
##
##
## ----- ----- -----
##
        imunodepre
                               yes
                                        Total
                       no
##
    CVD
                 2874 (98.0%) 58 (2.0%)
                                  2932 (100.0%)
    no
##
                 418 (96.5%)
                          15 (3.5%)
                                   433 (100.0%)
    yes
                 3292 (97.8%)
##
                          73 (2.2%) 3365 (100.0%)
   Total
## ----- ----- -----
  Chi.squared df p.value
 -----
##
   3.2567
           1 0.0711
##
##
 _____
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.78
           1.00
                   3.17
##
```

```
## ------
## Risk Ratio Lo - 0% Hi - 0%
## ------
## 1.02 1.02 1.02
```

Renal

```
ctable(data$CVD, data$renal, chisq=TRUE, prop="r", OR=TRUE, useNA = "no")
## Cross-Tabulation, Row Proportions
## CVD * renal
## Data Frame: data
##
##
##
##
         renal
                                            Total
                       no
                                 yes
##
     CVD
                2890 (98.8%)
                            36 (1.2%) 2926 (100.0%)
##
     no
     yes
##
                414 (95.6%)
                            19 (4.4%)
                                      433 (100.0%)
                                    3359 (100.0%)
##
   Total
                3304 (98.4%)
                            55 (1.6%)
##
##
  Chi.squared df p.value
  -----
             1
##
    21.431
  _____
##
  Odds Ratio Lo - 95% Hi - 95%
##
     3.68
              2.09
                       6.48
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
     1.03
              1.03
```

The variable comorb_group indicates the number o comorbidities of each notification.

Any comorbidity

```
df <- select(data,obesity,hematologic,hepatic,asthma,diabetes,neurologic,pneumologic,imunodepre,renal)
soma <- function(x){
  if (sum(is.na(x))==9)
    return(NA_character_)</pre>
```

```
return(sum(!is.na(x) & x=="yes"))
data$qt_comorb_aux <- apply(df,1,soma)</pre>
data <- data %>%
 mutate(qt_comorb = case_when(qt_comorb_aux== 9 ~ "9 comorbidities",
                             qt_comorb_aux == 7 ~ "7 comorbidities",
                             qt comorb aux== 6 ~ "6 comorbidities",
                             qt_comorb_aux == 5 ~ "5 comorbidities",
                             qt_comorb_aux== 4 ~ "4 comorbidities",
                             qt_comorb_aux == 3 ~ "3 comorbidities",
                             qt_comorb_aux == 2 ~ "2 comorbidities",
                             qt_comorb_aux == 1 ~ "1 comorbidity",
                             qt_comorb_aux == 0 ~ "No comorbidity",
                          TRUE ~ NA_character_))
data <- data %>%
 mutate(comorb_group = case_when(qt_comorb_aux == 0 ~ "No comorbidity",
                             qt_comorb_aux == 1 ~ "1 comorbidity",
                             qt_comorb_aux == 2 ~ "2 comorbidities",
                             qt_comorb_aux > 2 ~ ">2 comorbidities",
                             TRUE ~ NA_character_))
data$comorb_group <-</pre>
 factor(data$comorb_group, levels = c("No comorbidity", "1 comorbidity", "2 comorbidities", ">2 comorb
with(data, ctable(comorb_group, CVD, prop="c", useNA = "no", chisq = TRUE, OR = TRUE))
## Cross-Tabulation, Column Proportions
## comorb_group * CVD
## Data Frame: data
##
##
## ----- ---- ----
##
                   CVD
                                               yes
                                  no
                                                           Total
##
      comorb_group
    No comorbidity
##
                        2126 (71.8%) 234 (46.8%) 2360 (68.2%)
##
     1 comorbidity
                         701 ( 23.7%) 186 ( 37.2%) 887 ( 25.6%)
## 2 comorbidities
                         116 ( 3.9%) 64 ( 12.8%) 180 ( 5.2%)
                          17 ( 0.6%) 16 ( 3.2%)
                                                    33 ( 1.0%)
##
  >2 comorbidities
             Total
                        2960 (100.0%) 500 (100.0%) 3460 (100.0%)
##
## ----- ----
##
## -----
## Chi.squared df p.value
## -----
## 165.5306 3 0
## -----
```

Outcome

ICU

```
#tabela cruzada de uti
with(data, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: data
##
##
##
        icu
                              yes
                                        Total
    CVD
            2043 (71.8%) 802 (28.2%) 2845 (100.0%)
356 (63.0%) 209 (37.0%) 565 (100.0%)
##
     no
##
    yes
            2399 (70.4%) 1011 (29.6%) 3410 (100.0%)
   Total
##
## -----
## Chi.squared df p.value
## -----
    17.0872
           1
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.50
            1.24
                    1.81
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.14
           1.14
                  1.14
```

1st trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "1tri")
with(df4, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df4
##
##
## ----- ---- -----
##
       icu no yes
    CVD
##
          121 (77.1%) 36 (22.9%) 157 (100.0%)
##
    no
           18 (58.1%) 13 (41.9%) 31 (100.0%)
##
   yes
           139 (73.9%) 49 (26.1%) 188 (100.0%)
  Total
## ----- ----
## -----
 Chi.squared df p.value
   3.9164
          1 0.0478
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    2.43
         1.09
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.33
          1.33
                1.33
```

2nd trimester

```
df4 <- data %>%
   filter(classi_gesta_puerp == "2tri")
with(df4, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))

## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df4
##
##
```

```
##
       icu
               no
                        yes
                                Total
##
   CVD
          315 (68.8%) 143 (31.2%)
                            458 (100.0%)
##
    no
    yes
##
           66 (58.4%)
                    47 (41.6%)
                            113 (100.0%)
          381 (66.7%) 190 (33.3%)
                            571 (100.0%)
##
  Total
  Chi.squared df p.value
 -----
   3.9355
          1 0.0473
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
   1.57
          1.03
                 2.39
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
    1.18
          1.18
                1.18
## -----
```

3rd trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "3tri")
with(df4, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df4
## ----- ---- -----
##
         icu
                no
                                            Total
                                  yes
     CVD
##
              883 (76.6%)
                          270 (23.4%) 1153 (100.0%)
##
     no
               159 (68.2%)
                           74 (31.8%)
    yes
                                      233 (100.0%)
              1042 (75.2%)
                          344 (24.8%)
                                       1386 (100.0%)
##
    Total
##
##
## Chi.squared df p.value
## -----
##
    6.7894
             1 0.0092
##
```

Puerperium

```
df4 <- data %>%
 filter(classi_gesta_puerp == "puerp")
with(df4, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df4
##
## ----- ----
##
        icu
                  no
                           yes
                                      Total
##
    CVD
##
            678 (66.9%) 335 (33.1%)
                               1013 (100.0%)
    no
    yes
##
             94 (58.4%)
                      67 (41.6%)
                                161 (100.0%)
##
            772 (65.8%) 402 (34.2%)
                               1174 (100.0%)
   Total
## ----- ---- -----
##
 Chi.squared df p.value
## -----
##
           1
   4.1332
               0.042
## -----
##
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
    1.44
            1.03
                    2.03
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
    1.15
          1.15 1.15
```

Duration of hospitalization in ICU

n	media	DP	mediana	minimo	maximo	q25	q75	IQR
514.00 114.00			7.00 10.50	0.00 0.00	94.00 105.00		15.00 17.00	

```
wilcox.test(icu_days ~ CVD, data = df)
```

```
## Wilcoxon rank sum test with continuity correction
##
## data: icu_days by CVD
## W = 25486, p-value = 0.02941
## alternative hypothesis: true location shift is not equal to 0
```

Ventilatory support

```
with(data, ctable(CVD, ven_support, prop="r", useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * ven_support
## Data Frame: data
##
                    No yes, invasive yes, noninvasive
##
        ven_support
                                                                 Total
##
     CVD
                   1470 (52.7%) 443 (15.9%)
                                                877 (31.4%) 2790 (100.0%)
##
     no
##
                    226 (40.7%)
                                110 (19.8%)
                                               219 (39.5%) 555 (100.0%)
     yes
                    1696 (50.7%) 553 (16.5%) 1096 (32.8%) 3345 (100.0%)
##
   Total
##
## -----
## Chi.squared df p.value
## -----
    26.5268 2
## -----
```

Intubation

```
data <- data %>%
 mutate(intubation = case_when(SUPORT_VEN == 1 ~ "yes",
                      SUPORT VEN == 2 ~ "no",
                      SUPORT VEN == 3 ~ "no",
                      TRUE ~ NA_character_)
with(data, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: data
##
##
## ----- ----- -----
                        no
        intubation
                                 yes
                                            Total
##
    CVD
##
                  2347 (84.1%) 443 (15.9%) 2790 (100.0%)
    no
                  445 (80.2%) 110 (19.8%)
##
                                      555 (100.0%)
   yes
                  2792 (83.5%) 553 (16.5%) 3345 (100.0%)
   Total
## ----- -----
## -----
## Chi.squared df p.value
   4.9304 1 0.0264
## -----
##
 -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
            1.04
##
                    1.65
    1.31
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
    1.05
           1.05
                   1.05
## -----
1st trimester
df4 <- data %>%
 filter(classi_gesta_puerp == "1tri")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = FALSE))
```

Cross-Tabulation, Row Proportions

```
## CVD * intubation
## Data Frame: df4
##
##
## ----- ---- ----- ----- ----- -----
         intubation no yes
##
                                                 Total
     CVD
##
                    135 (88.8%) 17 (11.2%)
##
     no
                                          152 (100.0%)
                                          29 (100.0%)
##
                     23 (79.3%) 6 (20.7%)
    yes
                    158 (87.3%) 23 (12.7%) 181 (100.0%)
##
   Total
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
                     5.80
            0.74
     2.07
## ----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
     1.12
            1.12 1.12
## -----
fisher.test(df4$CVD, df4$intubation)
##
## Fisher's Exact Test for Count Data
##
## data: df4$CVD and df4$intubation
## p-value = 0.2182
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.601484 6.248095
## sample estimates:
## odds ratio
## 2.061721
2nd trimester
df4 <- data %>%
 filter(classi_gesta_puerp == "2tri")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
##
```

----- ----- -----

```
##
       intubation no
                                  Total
                           yes
##
    CVD
##
   no
              373 (85.0%)
                      66 (15.0%)
                             439 (100.0%)
               88 (79.3%)
                       23 (20.7%)
                             111 (100.0%)
##
    yes
                             550 (100.0%)
##
  Total
              461 (83.8%) 89 (16.2%)
## ----- ----- -----
## -----
 Chi.squared df p.value
 -----
   1.7139 1 0.1905
##
 _____
## -----
 Odds Ratio Lo - 95% Hi - 95%
    1.48
          0.87
##
                2.51
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.07
        1.07
               1.07
## ------
```

3rd trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "3tri")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
## ----- ----- -----
          intubation
                                        yes
##
     CVD
                     1007 (89.0%) 125 (11.0%)
                                             1132 (100.0%)
##
     no
                     195 (84.4%) 36 (15.6%)
##
                                           231 (100.0%)
    yes
                    1202 (88.2%) 161 (11.8%) 1363 (100.0%)
## ----- ----
##
##
  Chi.squared df p.value
    3.3759
             1
                  0.0662
##
```

```
## Odds Ratio Lo - 95% Hi - 95%
## ------
## 1.49 1.00 2.22
## -----
## #
## Risk Ratio Lo - 0% Hi - 0%
## ------
## 1.05 1.05 1.05
```

Puerperium

```
df4 <- data %>%
 filter(classi_gesta_puerp == "puerp")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
## ----- ----- -----
##
        intubation
                       no
                                yes
##
    CVD
##
    no
                778 (77.6%)
                          224 (22.4%) 1002 (100.0%)
##
                118 (75.2%)
                          39 (24.8%)
                                   157 (100.0%)
    yes
                 896 (77.3%) 263 (22.7%) 1159 (100.0%)
##
   Total
## ----- -----
##
##
## Chi.squared df p.value
## -----
   0.3468
           1 0.5559
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
    1.15
            0.78
                   1.70
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
   1.03
         1.03
                 1.03
```

Outcome - cure x death

```
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: data
##
## ----- ----- -----
##
                                 death
        evolution
                       cure
                                             Total
##
    CVD
                 2559 (86.5%) 401 (13.5%) 2960 (100.0%)
##
    no
                  488 (81.1%) 114 (18.9%) 602 (100.0%)
##
    yes
                3047 (85.5%) 515 (14.5%) 3562 (100.0%)
  Total
##
## ----- ----- -----
##
## Chi.squared df p.value
## -----
   11.3176 1
                8e-04
##
 Odds Ratio Lo - 95% Hi - 95%
## -----
            1.19
    1.49
                    1.88
##
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.07
            1.07
## -----
1st trimester
df4 <- data %>%
 filter(classi_gesta_puerp == "1tri")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = FALSE))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
##
##
         evolution
                      cure
                                death
                                          Total
##
    CVD
```

with(data, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))

```
##
                  146 (90.7%) 15 (9.3%) 161 (100.0%)
     no
##
                   30 (88.2%) 4 (11.8%) 34 (100.0%)
     yes
##
                  176 (90.3%) 19 ( 9.7%) 195 (100.0%)
## ----- ----- -----
## -----
## Odds Ratio Lo - 95% Hi - 95%
     1.30
             0.40
                      4.18
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
           1.03 1.03
    1.03
fisher.test(df4$CVD, df4$evolution)
##
## Fisher's Exact Test for Count Data
## data: df4$CVD and df4$evolution
## p-value = 0.7494
## alternative hypothesis: true odds ratio is not equal to 1
## 95 percent confidence interval:
## 0.2926084 4.4639958
## sample estimates:
## odds ratio
## 1.295924
2nd trimester
df4 <- data %>%
 filter(classi_gesta_puerp == "2tri")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
## ----- -----
         evolution cure death
                                             Total
##
     CVD
                  410 (87.0%) 61 (13.0%)
##
                                         471 (100.0%)
     no
                   90 (75.0%) 30 (25.0%)
##
    yes
                                         120 (100.0%)
##
                  500 (84.6%) 91 (15.4%)
                                         591 (100.0%)
  Total
## ----- --
##
```

```
## Chi.squared df p.value
## -----
   9.753
##
        1 0.0018
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
   2.24
##
         1.37
               3.67
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
   1.16 1.16 1.16
##
```

3rd trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "3tri")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
##
## ----- ----- -----
        evolution cure death
    CVD
##
##
    no
               1071 (89.3%) 128 (10.7%) 1199 (100.0%)
##
                212 (85.5%) 36 (14.5%) 248 (100.0%)
   yes
               1283 (88.7%) 164 (11.3%) 1447 (100.0%)
  Total
## ------ ------
##
## -----
## Chi.squared df p.value
## -----
##
   2.6461
           1 0.1038
 -----
##
## Odds Ratio Lo - 95% Hi - 95%
## -----
                  2.12
    1.42
           0.95
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
```

```
## 1.04 1.04 1.04
## -----
```

Puerperium

```
df4 <- data %>%
 filter(classi_gesta_puerp == "puerp")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
        evolution cure death Total
##
     CVD
##
                 856 (81.9%) 189 (18.1%) 1045 (100.0%)
    no
                 133 (78.2%) 37 (21.8%) 170 (100.0%)
##
    yes
##
                 989 (81.4%) 226 (18.6%) 1215 (100.0%)
   Total
## -----
##
## -----
## Chi.squared df p.value
## -----
##
    1.0751
            1 0.2998
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
          0.85 1.87
     1.26
##
## Risk Ratio Lo - 0% Hi - 0%
    1.05 1.05 1.05
```

Time elapsed between the start of symptoms and the outcome (days)

```
data <- data %>%
   dplyr::mutate(
   dt_sin_pri = as.Date(DT_SIN_PRI, format = "%d/%m/%Y"),
   dt_evoluca = as.Date(DT_EVOLUCA, format = "%d/%m/%Y"),
   days_symp_evol = as.numeric(dt_evoluca-dt_sin_pri)
)
```

	n	media	DP	mediana	minimo	maximo	q25	q75	IQR
no	2863.00	16.00	15.80	13.00	0.00	244.00	8.00	19.00	11.00
yes	586.00	17.10	12.80	14.00	0.00	113.00	9.00	20.75	11.75

```
wilcox.test(days_symp_evol ~ CVD, data = df)
```

```
##
## Wilcoxon rank sum test with continuity correction
##
## data: days_symp_evol by CVD
## W = 751451, p-value = 6.809e-05
## alternative hypothesis: true location shift is not equal to 0
```

Considering only death cases

n	media	DP	mediana	minimo	maximo	q25	q75	IQR
399.00 114.00			18.00 17.00		158.00 113.00			

```
wilcox.test(days_symp_evol ~ CVD, data = df)
```

```
##
## Wilcoxon rank sum test with continuity correction
##
## data: days_symp_evol by CVD
## W = 23410, p-value = 0.6331
## alternative hypothesis: true location shift is not equal to 0
```

Considering only cure cases

n	media	DP	mediana	minimo	maximo	q25	q75	IQR
2464.00 472.00			12.00 13.00	$0.00 \\ 0.00$			18.00 19.00	

```
##
## Wilcoxon rank sum test with continuity correction
##
## data: days_symp_evol by CVD
## W = 513735, p-value = 5.814e-05
## alternative hypothesis: true location shift is not equal to 0
```

Distribution of number of hospitalizations due to COVID-19 infection by month of first symptoms, stratified by groups with and without CVD

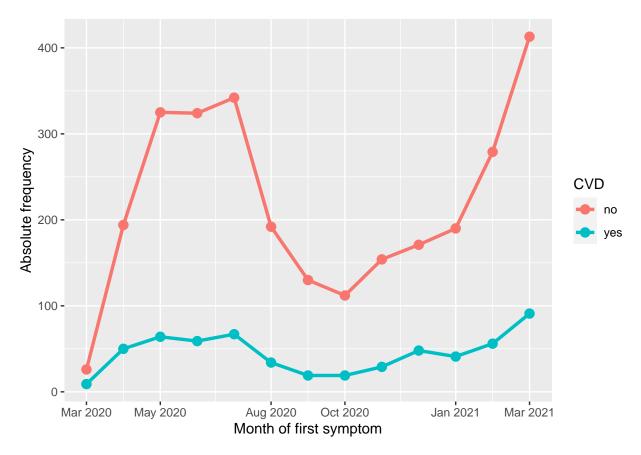
```
# Date of the first symptoms
data <- data %>%
mutate(dt_first_symp = as.Date(DT_SIN_PRI, format = "%d/%m/%Y")
)
```

The next graph presents the temporal evolution of hospitalizations by month of first symptoms, stratified by groups with and without cardiovascular disease.

```
## Creating the variable month_year
d1 <- data[data$dt_first_symp < as.Date("01/04/2021",format="%d/%m/%Y"),]
d1 <- d1 %>%
mutate(month_year = paste(formatC(month(dt_first_symp), width=2, format="d", flag="0"),year(dt_first_symp)
d<- table(factor(d1$month_year, levels = c("03/2020", "04/2020", "05/2020", "06/2020", "07/2020", "08/2
G <- data.table(d)
Sys.setlocale("LC_TIME","C")

## [1] "C"

G$V1 <- as.yearmon(G$V1, format = "%m/%Y")
ggplot(G, aes(x=V1, y=N, colour=factor(V2), group=V2)) +
geom_line(size=1.2) + geom_point(size=3) +
labs(x="Month of first symptom", y="Absolute frequency", color = "CVD")</pre>
```



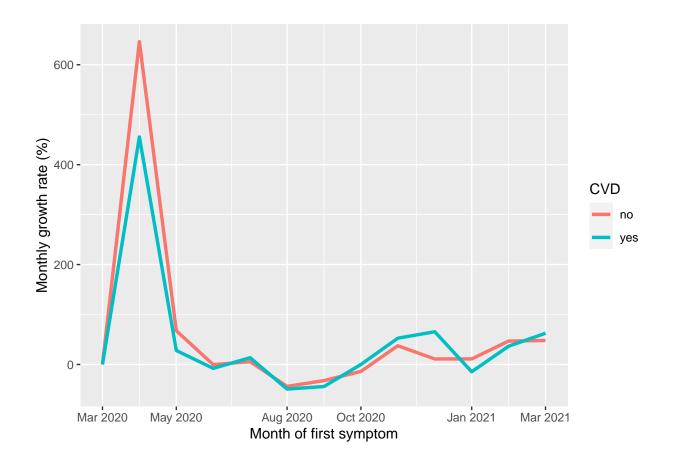
Frequency table of number of hospitalizations due to COVID-19 infection by month of first symptoms, stratified by groups with and without CVD.

```
## Cross-Tabulation, Column Proportions
## month_year * CVD
## Data Frame: d1
##
##
##
                  CVD
                                                                Total
                                   no
                                                  yes
##
     month_year
##
        01/2021
                         190 (
                                6.7%)
                                          41 ( 7.0%)
                                                         231 (
                                                                6.7%)
        02/2021
                         279 (
                                9.8%)
                                          56 ( 9.6%)
                                                         335 (
                                                                9.7%)
##
##
        03/2020
                          26 ( 0.9%)
                                          9 ( 1.5%)
                                                          35 ( 1.0%)
                         413 ( 14.5%)
##
        03/2021
                                          91 (15.5%)
                                                         504 (14.7%)
##
        04/2020
                         194 ( 6.8%)
                                          50 ( 8.5%)
                                                         244 ( 7.1%)
##
        05/2020
                         325 (11.4%)
                                          64 (10.9%)
                                                         389 (11.3%)
##
        06/2020
                         324 (11.4%)
                                          59 (10.1%)
                                                         383 (11.1%)
##
        07/2020
                         342 ( 12.0%)
                                          67 (11.4%)
                                                         409 (11.9%)
##
        08/2020
                         192 ( 6.7%)
                                                         226 ( 6.6%)
                                          34 ( 5.8%)
##
        09/2020
                         130 ( 4.6%)
                                          19 (
                                               3.2%)
                                                         149 ( 4.3%)
##
        10/2020
                         112 ( 3.9%)
                                          19 (
                                               3.2%)
                                                         131 ( 3.8%)
##
        11/2020
                         154 ( 5.4%)
                                          29 (
                                               4.9%)
                                                         183 ( 5.3%)
        12/2020
                         171 ( 6.0%)
                                          48 (8.2%)
                                                         219 ( 6.4%)
##
```

```
## Total 2852 (100.0%) 586 (100.0%) 3438 (100.0%)
```

Now, to understand better the monthly growth rate of hospitalized cases by COVID-19, we present the following graph:

```
D1<- table(d1$month_year, d1$CVD) %>%
      data.table()
D1$V1 <- as.yearmon(D1$V1, format = \%m/\%Y")
# Calculating the monthly growth rate
D1 <- D1 %>%
  arrange(D1)
dif<-c()
dif[1:2]<-0
for(i in 1:nrow(D1)){
  if( i %% 2 == 0){
    dif[i] < -D1$N[i+2] -D1$N[i]
  }
 else{
    dif[i] < -D1$N[i+2] -D1$N[i]
}
D1$dif<-dif
taxa_cresc<-c()</pre>
taxa_cresc[1:2]<-0</pre>
for(i in 1:(nrow(D1)-2)){
  if(i \% 2 == 0){
    taxa_cresc[i+2]<-(dif[i])/(D1$N[i])</pre>
  }
 else{
    taxa_cresc[i+2]<-(dif[i])/(D1$N[i])</pre>
  }
D1$growth_rate <-taxa_cresc*100
ggplot(D1) + geom_line(aes(x=V1, y = growth_rate, color = V2), size=1.2) + labs(x="Month of first symp
```



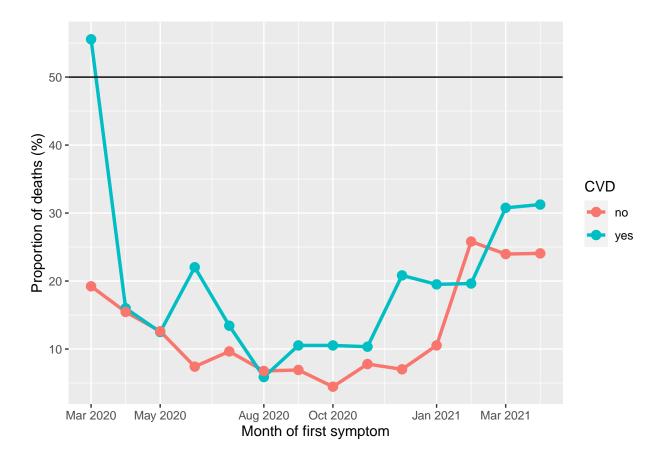
Proportion of deaths in the CVD and non-CVD groups, fixed the month of the first symptom.

```
# Proportion of deaths in the group with presence of CVD
d1 <- data
d1 <- d1 %>%
  filter(CVD == 'yes')
d1 <- d1 %>%
  mutate(month_year = paste(formatC(month(dt_first_symp), width=2, format="d", flag="0"), year(dt_first_
d<- prop.table(table(d1$month_year, d1$evolution),1)</pre>
G3 <- as.data.frame(d)
G3 <- G3[G3$Var2 == "death",]
G3$Freq <- round((G3$Freq)*100, 2)
G3$Var1 \leftarrow as.yearmon(G3$Var1, format = "%m/%Y")
# Proportion of deaths in the group with absence of CVD
d2 <- data
d2 <- d2 %>%
 filter(CVD == 'no')
d2 <- d2 %>%
  mutate(month_year = paste(formatC(month(dt_first_symp), width=2, format="d", flag="0"), year(dt_first_
d2<- prop.table(table(d2$month_year, d2$evolution),1)</pre>
G32 <- as.data.frame(d2)
```

```
G32 <- G32[G32$Var2 == "death",]
G32$Freq <- round((G32$Freq)*100, 2)
G32$Var1 <- as.yearmon(G32$Var1, format = "%m/%Y")

# Concatening the proportion of deaths in both groups, with and without CVD
G33 <- rbind(G3, G32)
cardio <- c(rep("yes", nrow(G3)), rep("no", nrow(G32)))
G33 <- cbind(G33, cardio)

ggplot(data=G33,aes(x = Var1, y=Freq, color = cardio)) + geom_line(size=1.2) +
geom_point(size=3) +
geom_hline(yintercept = 50) +
labs(x="Month of first symptom", y="Proportion of deaths (%)", color = "CVD")
```



Propensity Score Matching (PSM):

The analysis that will be made in this section aims to understand the effect of having cardiovascular disease or not on some variables of interest such as symptoms and outcome. In order to make the CVD and non-CVD groups similar with respect to the distribution of some variables that may bring confusion to the study, we will make use of the Propensity Score Matching (PSM) technique. The variables we will control in the analysis are: age, race, diabetes and obesity.

First, we present the difference result between the groups regarding the control variables before the PSM. We consider as "balanced" the cases with mean difference greater than 0.05. As we can see, for most categories of control variables the groups are not balanced before the matching.

```
##
                                             M.Threshold.Un
                           Type Diff.Un
## NU_IDADE_N
                        Contin. 0.5949 Not Balanced, >0.05
## ethnicity_black
                       Binary 0.0258
                                            Balanced, <0.05
## ethnicity_brown
                         Binary -0.0535 Not Balanced, >0.05
## ethnicity_indigenous Binary -0.0097
                                            Balanced, <0.05
                                            Balanced, <0.05
## ethnicity white
                         Binary 0.0429
## ethnicity_yellow
                         Binary -0.0054
                                            Balanced, <0.05
## ethnicity:<NA>
                         Binary 0.0264
                                            Balanced, <0.05
## diabetes_yes
                         Binary 0.2218 Not Balanced, >0.05
## diabetes:<NA>
                         Binary 0.2149 Not Balanced, >0.05
## obesity yes
                         Binary 0.1461 Not Balanced, >0.05
## obesity:<NA>
                         Binary 0.2359 Not Balanced, >0.05
## Balance tally for mean differences
##
                       count
## Balanced, <0.05
                           5
## Not Balanced, >0.05
                           6
##
## Variable with the greatest mean difference
##
      Variable Diff.Un
                            M.Threshold.Un
  NU_IDADE_N 0.5949 Not Balanced, >0.05
##
##
## Sample sizes
##
       Control Treated
## All
          2960
As one can see, all variables categories are balanced after PSM.
## Estimating the weights of each sampling unit
fit <- weightit(cardio ~ NU_IDADE_N + ethnicity + diabetes + obesity, use.mlogit = FALSE, data = data,
                  method = "ps", estimand = "ATE")
data$weight <- fit$weights</pre>
bal.tab(fit, m.threshold = 0.05, disp.v.ratio = TRUE)
## Call
   weightit(formula = cardio ~ NU_IDADE_N + ethnicity + diabetes +
##
       obesity, data = data, method = "ps", estimand = "ATE", use.mlogit = FALSE)
##
## Balance Measures
##
                            Type Diff.Adj
                                              M. Threshold V. Ratio. Adj
## prop.score
                        Distance 0.0394 Balanced, <0.05
                                                                1.3504
## NU IDADE N
                         Contin.
                                   0.0041 Balanced, <0.05
                                                                0.9849
## ethnicity_black
                         Binary -0.0003 Balanced, <0.05
## ethnicity_brown
                          Binary -0.0088 Balanced, <0.05
## ethnicity_indigenous
                         Binary 0.0056 Balanced, <0.05
```

bal.tab(cardio ~ NU_IDADE_N + ethnicity + diabetes + obesity, data = data, estimand = "ATE", m.threshol

data\$cardio <- ifelse(data\$CVD=="yes",1,0)</pre>

Balance Measures

```
## ethnicity_white
                       Binary 0.0088 Balanced, <0.05
## ethnicity_yellow
                       Binary -0.0052 Balanced, <0.05
                       Binary 0.0144 Balanced, <0.05
## ethnicity:<NA>
                        Binary -0.0036 Balanced, <0.05
## diabetes_yes
                        Binary 0.0117 Balanced, <0.05
## diabetes:<NA>
## obesity yes
                        Binary 0.0064 Balanced, <0.05
## obesity:<NA>
                        Binary 0.0133 Balanced, <0.05
##
## Balance tally for mean differences
##
                     count
## Balanced, <0.05
                        12
## Not Balanced, >0.05
                         0
## Variable with the greatest mean difference
         Variable Diff.Adj
                            M.Threshold
## ethnicity:<NA> 0.0144 Balanced, <0.05
##
## Effective sample sizes
            Control Treated
## Unadjusted 2960. 602.
## Adjusted 2332.6 331.15
```

Symptom analysis after PSM:

----- Lo - 95% Hi - 95% ## ----- ----

1 0.2275

0.97

1.17

1.4563

1.06

##

##

##

```
with(data, ctable(CVD, fever, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
Fever
## Cross-Tabulation, Row Proportions
## CVD * fever
## Data Frame: data
##
## ----- ----- -----
         fever
                        no
                                    yes
                                                Total
##
    CVD
               1535.7 (44.5%) 1914.5 (55.5%) 3450.2 (100.0%)
##
     no
##
    yes
               1510.0 (43.0%) 1997.8 (57.0%) 3507.8 (100.0%)
  Total
           3045.8 (43.8%) 3912.2 (56.2%) 6958.0 (100.0%)
## ----- ----- -----
##
## -----
## Chi.squared df p.value
```

```
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
                 1.03
    1.03
         1.03
## -----
with(data, ctable(CVD, cough, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
Cough
## Cross-Tabulation, Row Proportions
## CVD * cough
## Data Frame: data
##
##
 ##
        cough
                                            Total
                     no
                                 yes
##
    CVD
##
             1161.8 (33.7%) 2290.8 (66.3%) 3452.6 (100.0%)
    no
##
   yes
             1124.8 (32.0%) 2395.0 (68.0%) 3519.8 (100.0%)
##
          2286.6 (32.8%) 4685.8 (67.2%) 6972.4 (100.0%)
   Total
## ----- ----- ------ ------
##
```

```
with(data, ctable(CVD, sore_throat, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Sore throat

```
## Cross-Tabulation, Row Proportions
## CVD * sore_throat
## Data Frame: data
##
## ----- ----- ----- ------ ------
                                     yes
       sore_throat
                       no
                                               Total
##
    CVD
##
    no
                  2651.6 (78.2%) 738.2 (21.8%) 3389.8 (100.0%)
##
                 2588.7 (76.6%) 790.8 (23.4%)
                                         3379.6 (100.0%)
   yes
   Total
                 5240.3 (77.4%) 1529.0 (22.6%)
                                         6769.3 (100.0%)
## ----- --- ---- ---- ----- ------
 -----
 Chi.squared df p.value
   2.4586
           1 0.1169
##
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
          0.98
    1.10
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
   1.02
           1.02
                 1.02
```

```
with(data, ctable(CVD, dyspnea, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Dyspnea

```
## Cross-Tabulation, Row Proportions
## CVD * dyspnea
## Data Frame: data
##
##
## ----- ----- -----
##
         dyspnea
                                                 Total
                         no
                                     yes
##
    CVD
##
               1574.0 (45.6%) 1876.2 (54.4%) 3450.2 (100.0%)
     no
               1196.9 (33.9%) 2337.6 (66.1%) 3534.4 (100.0%)
##
    yes
            2770.8 (39.7%) 4213.8 (60.3%) 6984.6 (100.0%)
##
##
## -----
## Chi.squared df p.value
## -----
```

```
100.3512 1 0
## -----
##
## -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
      1.49
  1.64
## -----
##
## -----
 Risk Ratio Lo - 0% Hi - 0%
## -----
       1.35
  1.35
           1.35
## -----
```

```
with(data, ctable(CVD, resp_dist, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Respiratory distress

```
## Cross-Tabulation, Row Proportions
## CVD * resp_dist
## Data Frame: data
##
## ----- ----- -----
       resp_dist
                                           Total
                       no
                                 yes
    CVD
##
##
    no
               1789.1 (52.1%) 1645.7 (47.9%)
                                    3434.8 (100.0%)
               1578.1 (45.4%) 1900.2 (54.6%)
                                    3478.3 (100.0%)
##
    yes
               3367.2 (48.7%) 3545.9 (51.3%)
                                    6913.1 (100.0%)
  Total
##
##
 -----
 Chi.squared df p.value
## -----
##
   30.9549 1
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
           1.19
    1.31
                 1.44
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
   1.15
          1.15
                1.15
```

```
with(data, ctable(CVD, saturation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
Saturation
## Cross-Tabulation, Row Proportions
## CVD * saturation
## Data Frame: data
##
##
        saturation
                          no
                                      yes
                                                 Total
    CVD
##
##
                 2164.0 (63.5%) 1241.8 (36.5%) 3405.8 (100.0%)
    no
                 1810.9 (52.3%) 1654.5 (47.7%) 3465.4 (100.0%)
    yes
   Total
                 3974.9 (57.8%) 2896.3 (42.2%) 6871.2 (100.0%)
##
## ----- ----- ------ ------ ------
##
 Chi.squared df p.value
## -----
   89.2193
          1
## -----
##
 Odds Ratio Lo - 95% Hi - 95%
## -----
##
    1.59
            1.45
                    1.75
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.22
            1.22
## -----
with(data, ctable(CVD, diarrhea, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
Diarrhea
## Cross-Tabulation, Row Proportions
## CVD * diarrhea
## Data Frame: data
##
##
```

3021.0 (88.6%) 389.7 (11.4%) 3410.7 (100.0%)

yes

Total

##

##

##

CVD

no

diarrhea

```
##
             3002.5 (89.0%) 371.5 (11.0%) 3374.1 (100.0%)
   ves
             6023.5 (88.8%) 761.2 (11.2%) 6784.7 (100.0%)
##
  Total
## ----- -----
##
 Chi.squared df p.value
## -----
   0.2517 1 0.6159
 _____
##
 _____
 Odds Ratio Lo - 95% Hi - 95%
##
         0.82
##
  0.96
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
   1.00
         1.00
               1.00
## -----
```

```
with(data, ctable(CVD, vomit, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Vomit

```
## Cross-Tabulation, Row Proportions
## CVD * vomit
## Data Frame: data
##
##
## ----- -----
##
        vomit
                    no
                                         Total
                              yes
##
    CVD
    no
             3062.7 (90.1%) 337.3 (9.9%) 3400.1 (100.0%)
##
             2998.6 (89.3%) 359.5 (10.7%) 3358.1 (100.0%)
    yes
            6061.3 (89.7%) 696.8 (10.3%) 6758.1 (100.0%)
##
  Total
## ----- ----- -----
## -----
 Chi.squared df p.value
 _____
   1.0397
##
          1 0.3079
##
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
   1.09
           0.93
                  1.27
##
```

```
## Risk Ratio Lo - 0% Hi - 0%
## -----
## 1.01 1.01 1.01
## ----
```

```
with(data, ctable(CVD, fatigue, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Fatigue

```
## Cross-Tabulation, Row Proportions
## CVD * fatigue
## Data Frame: data
##
##
## ----- -----
##
        fatigue
                                            Total
                                 yes
                      no
    CVD
##
                                     2471.8 (100.0%)
##
              1892.0 (76.5%) 579.8 (23.5%)
    no
              1733.2 (74.8%) 583.8 (25.2%) 2317.0 (100.0%)
##
    yes
##
  Total
              3625.2 (75.7%) 1163.6 (24.3%) 4788.8 (100.0%)
## ----- ----- -----
##
 Chi.squared df p.value
## -----
    1.871
           1 0.1714
##
## -----
##
## Odds Ratio Lo - 95% Hi - 95%
           0.96
##
    1.10
                   1.25
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
    1.02
           1.02
```

```
with(data, ctable(CVD, loss_smell, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Loss of smell

```
## Cross-Tabulation, Row Proportions
## CVD * loss_smell
```

```
##
## ----- ----- -----
                                       yes
##
         loss_smell
                                                   Total
                           no
##
   CVD
                   2002.3 (81.1%) 467.5 (18.9%)
                                          2469.8 (100.0%)
##
    no
                  1787.6 (77.5%) 520.4 (22.5%) 2308.0 (100.0%)
##
    yes
##
   Total
                   3789.9 (79.3%) 987.9 (20.7%) 4777.7 (100.0%)
##
##
  Chi.squared df p.value
## -----
          1 0.0023
   9.2983
##
  -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
            1.08
##
    1.25
                   1.43
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
    1.05
            1.05
                    1.05
with(data, ctable(CVD, loss_taste, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
Loss of taste
## Cross-Tabulation, Row Proportions
## CVD * loss taste
## Data Frame: data
##
##
                                       yes
##
         loss_taste
                                                   Total
##
     CVD
                   2008.7 (81.5%) 455.8 (18.5%)
                                          2464.6 (100.0%)
##
    no
                   1792.7 (77.6%) 518.6 (22.4%)
##
                                          2311.3 (100.0%)
     yes
                   3801.4 (79.6%) 974.4 (20.4%)
                                          4775.9 (100.0%)
   Total
##
  _____
  Chi.squared df p.value
## -----
   11.1723
            1
                 8e-04
## -----
```

Data Frame: data

##

```
##
## -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
   1.27
        1.11
             1.47
 _____
##
##
 Risk Ratio Lo - 0% Hi - 0%
## -----
  1.05
       1.05
            1.05
## -----
```

```
with(data, ctable(CVD, abd_pain, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Abdominal pain

```
## Cross-Tabulation, Row Proportions
## CVD * abd_pain
## Data Frame: data
##
##
 ##
        abd_pain
                               yes
                      no
                                          Total
##
    CVD
##
               2243.3 (91.3%) 214.6 (8.7%)
                                   2457.9 (100.0%)
    no
               2102.5 (92.1%) 180.7 (7.9%)
                                  2283.2 (100.0%)
##
    yes
   Total
              4345.8 (91.7%) 395.3 (8.3%)
                                  4741.2 (100.0%)
##
##
  Chi.squared df p.value
##
 _____
   0.9274
          1 0.3355
 _____
##
##
 -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
##
    0.90
           0.73
                  1.10
## -----
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
    0.99
         0.99 0.99
```

Any respiratory symptom Now we are going to create the variable resp_symp indicating the presence of at least one respiratory symptom (Dyspnea, Fadigue, Saturation, Respiratory distress). Empty cells are

considered missing data (<NA>). The variable qt_resp_symp indicates the number of respiratory symptoms of each case.

```
with(data, ctable(CVD, resp_symp, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * resp_symp
## Data Frame: data
##
##
## -----
##
        resp_symp
                         no
                                    yes
                                                Total
    CVD
                1056.4 (30.5%) 2408.6 (69.5%) 3465.1 (100.0%)
##
    no
    yes
                 738.7 (20.7%)
                            2834.6 (79.3%)
                                         3573.4 (100.0%)
##
##
                 1795.2 (25.5%) 5243.3 (74.5%) 7038.5 (100.0%)
   Total
##
  _____
  Chi.squared df p.value
 -----
    88.6786
           1
##
##
  Odds Ratio Lo - 95% Hi - 95%
    1.68
            1.51
                    1.88
## -----
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.47
            1.47
                   1.47
## -----
with(data, ctable(CVD, qt_resp_symp, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * qt_resp_symp
## Data Frame: data
##
##
 ##
                      1 symptom
                                  2 symptoms
                                             3 symptoms
                                                       4 symptoms No respir
        qt_resp_symp
##
    CVD
                                          768.8 (22.2%)
##
                    763.3 (22.0%)
                              616.1 (17.8%)
                                                       260.4 (7.5%)
                                                                       1
    no
##
    yes
                   709.1 (19.8%) 892.6 (25.0%) 949.9 (26.6%)
                                                       283.0 (7.9%)
                   1472.4 (20.9%) 1508.7 (21.4%) 1718.8 (24.4%) 543.3 (7.7%)
##
##
## -----
## Chi.squared df p.value
```

```
## ------
## 127.2795 4 0
## -----
```

Any symptom Similar to the analysis of any respiratory symptom, we are going to create the variable symp indicating the presence of at least one symptom (Dyspnea, Fatigue, Saturation, Respiratory distress, Fever, Cough, Sore throat, Diarrhea, Vomit, Abdominal pain, Loss of smell, Loss of taste). Empty cells are considered missing data (<NA>).

```
with(data, ctable(CVD, symp, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

```
## Cross-Tabulation, Row Proportions
## CVD * symp
## Data Frame: data
##
##
## ----- ---- ----- ------ ------
             no
                             yes
##
       \operatorname{\mathtt{symp}}
                                       Total
##
    CVD
##
           285.8 (8.2%) 3197.3 (91.8%) 3483.1 (100.0%)
    no
##
           135.9 (3.8%) 3458.6 (96.2%) 3594.5 (100.0%)
    yes
           421.7 (6.0%) 6655.9 (94.0%) 7077.6 (100.0%)
##
   Total
  _____ _____
##
  -----
  Chi.squared df p.value
##
  _____
##
   61.0734
##
          1
                Ω
 -----
##
## -----
 Odds Ratio Lo - 95% Hi - 95%
           1.85
                  2.81
##
    2.28
##
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    2.17 2.17 2.17
## -----
```

Outcome analysis after PSM:

ICU

```
with(data, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
### Cross-Tabulation, Row Proportions
## CVD * icu
```

```
##
##
## ----- ---- -----
##
        icu
                     no
                                yes
                                            Total
##
   CVD
            2421.5 (71.6%) 959.7 (28.4%) 3381.2 (100.0%)
##
    no
            2318.3 (67.5%) 1117.4 (32.5%) 3435.7 (100.0%)
##
   yes
##
   Total
            4739.9 (69.5%) 2077.1 (30.5%) 6817.0 (100.0%)
## ----- ---- ----
##
## -----
## Chi.squared df p.value
   13.5876 1 2e-04
##
 -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
##
    1.22
            1.10
                  1.35
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
    1.06
           1.06
                  1.06
df43 <- data %>%
 filter(classi_gesta_puerp == "1tri")
with(df43, ctable(CVD, icu, prop="r", useNA = "no", OR=TRUE, chisq = TRUE, weights = weight))
1st trimester
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df43
##
##
## ----- ---- -----
##
        icu
                   no
                              yes
                                        Total
##
    CVD
            137.3 (76.8%) 41.4 (23.2%)
##
                                  178.7 (100.0%)
    no
             83.2 (63.3%) 48.2 (36.7%)
                                  131.4 (100.0%)
   yes
            220.5 (71.1%) 89.6 (28.9%) 310.2 (100.0%)
   Total
## ----- ---- -----
##
## -----
## Chi.squared df p.value
```

Data Frame: data

```
6.0614 1 0.0138
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
          1.17
    1.92
## -----
##
 _____
## Risk Ratio Lo - 0% Hi - 0%
## -----
       1.21 1.21
##
   1.21
## -----
df43 <- data %>%
 filter(classi_gesta_puerp == "2tri")
with(df43, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
2nd trimester
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df43
##
##
## ----- ---- -----
##
                                    Total
      icu
                no
                          yes
##
   CVD
          362.9 (68.1%) 169.9 (31.9%) 532.9 (100.0%)
##
    no
           377.5 (61.4%) 237.4 (38.6%)
    yes
##
                              614.9 (100.0%)
          740.4 (64.5%) 407.4 (35.5%) 1147.8 (100.0%)
##
  Total
## ----- ----
##
## -----
## Chi.squared df p.value
        1 0.0208
   5.3395
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
        1.05
   1.34
## -----
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
```

```
df43 <- data %>%
 filter(classi_gesta_puerp == "3tri")
with(df43, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
3rd trimester
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df43
##
##
## ----- ---- -----
##
        icu
                                 yes
                                            Total
                     no
##
    CVD
##
            1058.1 (76.8%) 318.8 (23.2%) 1376.8 (100.0%)
    no
             1252.9 (74.1%) 437.2 (25.9%) 1690.1 (100.0%)
##
    yes
##
            2311.0 (75.4%) 755.9 (24.6%) 3066.9 (100.0%)
  Total
## ----- ---- -----
##
## Chi.squared df p.value
## -----
    2.8646
            1 0.0905
## -----
##
## Odds Ratio Lo - 95% Hi - 95%
##
    1.16
            0.98
                    1.37
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
           1.04
    1.04
## -----
df43 <- data %>%
 filter(classi_gesta_puerp == "puerp")
```

Puerperium

1.11 1.11 1.11

with(df43, ctable(CVD, icu, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))

```
## Cross-Tabulation, Row Proportions
## CVD * icu
## Data Frame: df43
##
## ----- ---- ----
                  no
       icu
                                      Total
                            yes
   CVD
##
##
    no
           802.7 (66.3%) 407.6 (33.7%) 1210.3 (100.0%)
           540.2 (60.3%)
                      355.7 (39.7%) 896.0 (100.0%)
##
    yes
  Total
          1342.9 (63.8%)
                      763.4 (36.2%) 2106.3 (100.0%)
## ----- ---- ----
## -----
 Chi.squared df p.value
   7.8264
          1 0.0051
##
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
         1.08
    1.30
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## ----- ----
   1.10
          1.10
```

Days of hospitalization in ICU

```
d_yes <- data %>%
  filter(CVD == "yes" & !is.na(icu_days))
d_no <- data %>%
  filter(CVD == "no" & !is.na(icu_days))
wtd.t.test(d_yes$icu_days, d_no$icu_days, weight = d_yes$weight, weighty = d_no$weight)
## $test
## [1] "Two Sample Weighted T-Test (Welch)"
## $coefficients
##
                df
                             p.value
     t.value
    1.2896729 164.5233473 0.1989742
##
## $additional
## Difference Mean.x
                           Mean.y Std. Err
## 1.437357 11.941962 10.504606 1.114513
```

Ventilatory support

```
with(data, ctable(CVD, ven_support, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * ven_support
## Data Frame: data
##
No yes, invasive yes, noninvasive
        ven_support
                                                               Total
##
    CVD
##
                  1693.0 (51.3%) 540.7 (16.4%) 1068.2 (32.4%)
                                                       3301.9 (100.0%)
    no
                  1611.5 (47.2%) 603.3 (17.7%) 1197.4 (35.1%) 3412.2 (100.0%)
##
    yes
                  3304.6 (49.2%) 1144.0 (17.0%) 2265.6 (33.7%) 6714.2 (100.0%)
  Total
## ----- ----- ----- ------
##
## Chi.squared df p.value
## -----
## 10.9876 2 0.0041
## -----
```

Intubation

1.10

##

0.97

1.25

```
with(data, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: data
##
## ----- ----- ------ ------ ------
        intubation
                                     yes
                                                Total
                          nο
##
    CVD
                2761.2 (83.6%) 540.7 (16.4%) 3301.9 (100.0%)
##
    no
##
   yes
                 2809.0 (82.3%) 603.3 (17.7%) 3412.2 (100.0%)
  Total
                5570.2 (83.0%) 1144.0 (17.0%) 6714.2 (100.0%)
## ----- ----- -----
##
 -----
 Chi.squared df p.value
    1.927 1 0.1651
##
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
```

```
## ------
## Risk Ratio Lo - 0% Hi - 0%
## ------
## 1.02 1.02 1.02
```

1st trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "1tri")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
##
## ----- ----- ------
##
        intubation
                         no
                                    yes
                                              Total
##
    CVD
##
                 153.4 (88.7%) 19.6 (11.3%) 173.0 (100.0%)
    no
                  97.1 (80.6%) 23.4 (19.4%) 120.5 (100.0%)
    yes
                 250.5 (85.3%) 43.0 (14.7%) 293.5 (100.0%)
   Total
##
##
  -----
 Chi.squared df p.value
         1 0.0776
   3.1142
##
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.89
            0.98
## -----
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.10
           1.10
                   1.10
```

${\bf 2nd\ trimester}$

```
df4 <- data %>%
  filter(classi_gesta_puerp == "2tri")
```

```
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
## ----- ----- -----
        intubation
                         no
                                     yes
##
   CVD
##
    no
                429.5 (84.4%) 79.6 (15.6%) 509.2 (100.0%)
                 493.7 (81.6%) 111.4 (18.4%) 605.1 (100.0%)
##
   yes
  Total
                 923.2 (82.9%) 191.1 (17.1%) 1114.3 (100.0%)
## ----- -
## Chi.squared df p.value
## -----
##
    1.31
            1 0.2524
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
    1.22
          0.89
##
## Risk Ratio Lo - 0% Hi - 0%
## -----
    1.03
           1.03 1.03
```

3rd trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "3tri")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
## ------ ------
                                          yes
##
          intubation
                                                        Total
                             no
##
    CVD
                   1205.3 (89.1%) 147.2 (10.9%) 1352.5 (100.0%)
##
     no
##
                    1479.2 (86.3%) 234.6 (13.7%) 1713.7 (100.0%)
    yes
```

```
3066.3 (100.0%)
           2684.5 (87.5%) 381.8 (12.5%)
## ----- ----- -----
##
## -----
## Chi.squared df p.value
## -----
 5.1945
      1 0.0227
## -----
##
##
 _____
 Odds Ratio Lo - 95% Hi - 95%
## -----
     1.04
  1.30
            1.62
## -----
##
## -----
 Risk Ratio Lo - 0% Hi - 0%
## -----
       1.03
  1.03
            1.03
## -----
```

Puerperium

```
df4 <- data %>%
 filter(classi_gesta_puerp == "puerp")
with(df4, ctable(CVD, intubation, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * intubation
## Data Frame: df4
##
##
##
          intubation
                              no
                                          yes
                                                         Total
     CVD
##
                      903.0 (76.3%) 280.6 (23.7%) 1183.6 (100.0%)
     no
##
    yes
                     654.0 (75.2%) 215.5 (24.8%)
                                                869.5 (100.0%)
##
                    1557.0 (75.8%) 496.2 (24.2%) 2053.1 (100.0%)
##
  -----
  Chi.squared df p.value
## -----
    0.2619 1 0.6088
##
##
## Odds Ratio Lo - 95% Hi - 95%
## -----
     1.06
               0.86
                        1.30
```

```
## ## Risk Ratio Lo - 0% Hi - 0% ## ------ ## 1.01 1.01 1.01 ## -----
```

Outcome - cure x death

```
with(data, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: data
##
                        cure
                                                   Total
         evolution
                                     \mathtt{death}
##
     CVD
##
                  3020.3 (86.0%) 491.3 (14.0%)
                                           3511.6 (100.0%)
    no
                  2974.7 (82.4%) 636.8 (17.6%) 3611.5 (100.0%)
##
    yes
   Total
                  5995.0 (84.2%) 1128.1 (15.8%) 7123.1 (100.0%)
## ----- ----- ----- ----- ------ -----
##
## -----
## Chi.squared df p.value
## -----
    17.433
            1
                  0
##
##
## Odds Ratio Lo - 95% Hi - 95%
## -----
           1.16 1.50
     1.32
##
## Risk Ratio Lo - 0% Hi - 0%
##
    1.04
            1.04
## -----
```

1st trimester

```
df4 <- data %>%
  filter(classi_gesta_puerp == "1tri")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
```

Cross-Tabulation, Row Proportions

```
## CVD * evolution
## Data Frame: df4
##
##
 ##
       evolution
                  cure death
                                     Total
##
   CVD
              165.5 (90.4%) 17.5 ( 9.6%)
##
   no
                               183.1 (100.0%)
##
             121.1 (88.1%) 16.4 (11.9%)
                               137.4 (100.0%)
   yes
             286.6 (89.4%) 33.9 (10.6%) 320.5 (100.0%)
##
  Total
##
## ------
## Chi.squared df p.value
       1 0.6267
  0.2366
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
## -----
          0.62
## -----
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
   1.03 1.03 1.03
## -----
```

2nd trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "2tri")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
##
##
         evolution
                        cure
                                   death
                                                 Total
##
    CVD
                  471.8 (86.2%) 75.7 (13.8%)
##
                                         547.5 (100.0%)
    no
                  484.2 (76.2%) 151.0 (23.8%) 635.3 (100.0%)
    yes
                 956.0 (80.8%) 226.7 (19.2%) 1182.7 (100.0%)
   Total
##
## ----- ----- -----
##
## -----
## Chi.squared df p.value
```

3rd trimester

```
df4 <- data %>%
 filter(classi_gesta_puerp == "3tri")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
##
## ----- ----- ------
        evolution
                       cure
                                 death
                                             Total
##
    CVD
                1279.2 (89.5%) 150.6 (10.5%)
                                      1429.8 (100.0%)
    no
                1565.4 (86.8%) 237.9 (13.2%)
##
                                      1803.3 (100.0%)
    yes
   Total
                2844.6 (88.0%) 388.5 (12.0%)
                                      3233.1 (100.0%)
## ----- ----- ------
## -----
## Chi.squared df p.value
## -----
   5.0783
         1 0.0242
## -----
##
 -----
 Odds Ratio Lo - 95% Hi - 95%
## -----
##
    1.29
           1.04
                   1.60
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
   1.03
           1.03
##
                 1.03
```

Puerperium

```
df4 <- data %>%
 filter(classi_gesta_puerp == "puerp")
with(df4, ctable(CVD, evolution, prop="r", OR=TRUE, useNA = "no", chisq = TRUE, weights = weight))
## Cross-Tabulation, Row Proportions
## CVD * evolution
## Data Frame: df4
##
##
##
                    cure death
        evolution
##
    CVD
##
    no
                 1009.5 (81.0%) 237.0 (19.0%) 1246.5 (100.0%)
##
                 709.0 (77.1%) 210.6 (22.9%) 919.6 (100.0%)
   yes
                 1718.5 (79.3%) 447.5 (20.7%) 2166.1 (100.0%)
   Total
## ----- ---- ----- ------ ------
## -----
## Chi.squared df p.value
    4.6463
           1
                0.0311
## -----
##
## -----
## Odds Ratio Lo - 95% Hi - 95%
    1.27 1.03 1.56
##
## -----
## Risk Ratio Lo - 0% Hi - 0%
## -----
##
    1.05 1.05 1.05
```

Time elapsed between the start of symptoms and the outcome (days) after PSM

```
d_yes <- data %>%
   filter(CVD == "yes" & !is.na(days_symp_evol))
d_no <- data %>%
   filter(CVD == "no" & !is.na(days_symp_evol))

# Performing weighted Student's t-tests to compare the CVD and non-CVD groups with respect to the time
wtd.t.test(d_yes$days_symp_evol, d_no$days_symp_evol, weight = d_yes$weight, weighty = d_no$weight)
```

```
## $test
## [1] "Two Sample Weighted T-Test (Welch)"
## $coefficients
##
        t.value
                          df
                                  p.value
      0.1099695 1063.6837985
                                0.9124543
##
## $additional
## Difference
                    Mean.x
                                Mean.y
                                          Std. Err
## 0.06197113 16.09992557 16.03795444 0.56353001
Considering only death cases
df <- data[!is.na(data$days_symp_evol) & !is.na(data$CVD),]</pre>
df <- df %>%
filter(evolution == "death")
d_yes <- df %>%
   filter(CVD == "yes" & !is.na(days_symp_evol))
d_no <- df %>%
   filter(CVD == "no" & !is.na(days_symp_evol))
# Fixed the death cases, we perform a weighted Student's t-tests to compare the CVD and non-CVD groups
wtd.t.test(d_yes$days_symp_evol, d_no$days_symp_evol, weight = d_yes$weight, weighty = d_no$weight)
## $test
## [1] "Two Sample Weighted T-Test (Welch)"
##
## $coefficients
                               p.value
##
       t.value
                        df
                             0.2086723
## -1.2611092 209.4593902
##
## $additional
                             Mean.y Std. Err
## Difference
                  {\tt Mean.x}
## -1.603168 18.525606 20.128775
Considering only cure cases
df <- data[!is.na(data$days_symp_evol) & !is.na(data$CVD),]</pre>
df <- df %>%
filter(evolution == "cure")
d yes <- df %>%
```

filter(CVD == "yes" & !is.na(days_symp_evol))

filter(CVD == "no" & !is.na(days_symp_evol))

d_no <- df %>%

Fixed the cure cases, we perform a weighted Student's t-tests to compare the CVD and non-CVD groups w wtd.t.test(d_yes\$days_symp_evol, d_no\$days_symp_evol, weight = d_yes\$weight, weighty = d_no\$weight)

```
## $test
## [1] "Two Sample Weighted T-Test (Welch)"
##
## $coefficients
## t.value df p.value
## 0.3550950 844.4701387 0.7226071
##
## $additional
## Difference Mean.x Mean.y Std. Err
## 0.2215881 15.5695137 15.3479256 0.6240248
```