Plasma

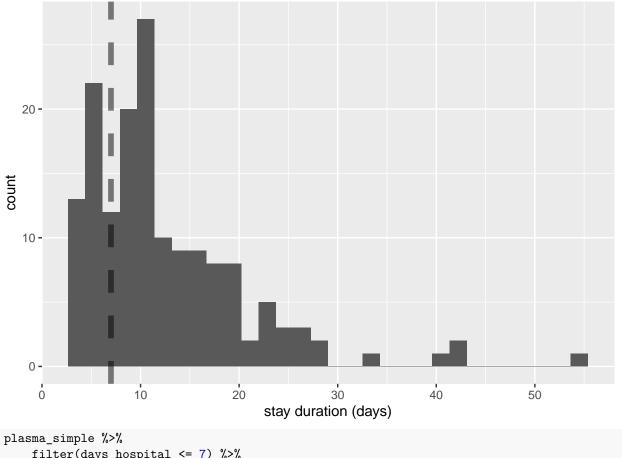
Federico Zertuche

```
library(tidyverse)
library(lubridate)
library(survival)
plasma_simple <- read_csv('./plasma_simple.csv')</pre>
##
## -- Column specification -----
## cols(
##
     .default = col_double(),
##
     hospital = col_character(),
     name = col_character(),
##
##
     sex = col_character(),
##
     vital_st = col_character(),
     hta = col_character(),
##
     dm = col_character(),
     obesity = col_character(),
##
##
     overweight = col_character()
## )
## i Use `spec()` for the full column specifications.
Hospital Stays:
  • 49 patients had stays of 7 or less days.
```

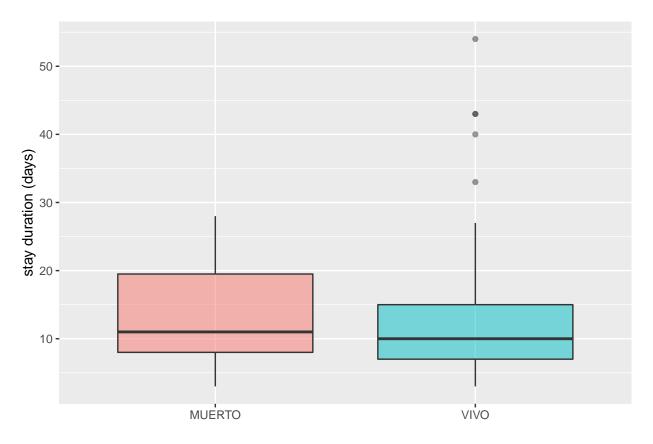
• Stays lasted 12.4 days in average (sd 8.18).

```
plasma_simple %>%
    ggplot() +
    geom_histogram(aes(x = days_hospital)) +
    geom_vline(xintercept = 7, linetype = 2, size = 2, alpha = 0.5) +
    labs(x = 'stay duration (days)')
```

`stat_bin()` using `bins = 30`. Pick better value with `binwidth`.



```
filter(days_hospital <= 7) %>%
    count()
## # A tibble: 1 x 1
##
         n
##
     <int>
## 1
        47
plasma_simple %>%
    summarise(mean(days_hospital), sd(days_hospital))
## # A tibble: 1 x 2
##
     `mean(days_hospital)` `sd(days_hospital)`
##
                                         <dbl>
                     <dbl>
                                          8.19
## 1
                      12.4
plasma_simple %>%
    ggplot() +
    geom_boxplot(aes(x = vital_st, y = days_hospital, fill = vital_st), alpha = 0.5) +
    theme(legend.position = "None") +
    labs(x = '', y = 'stay duration (days)')
```

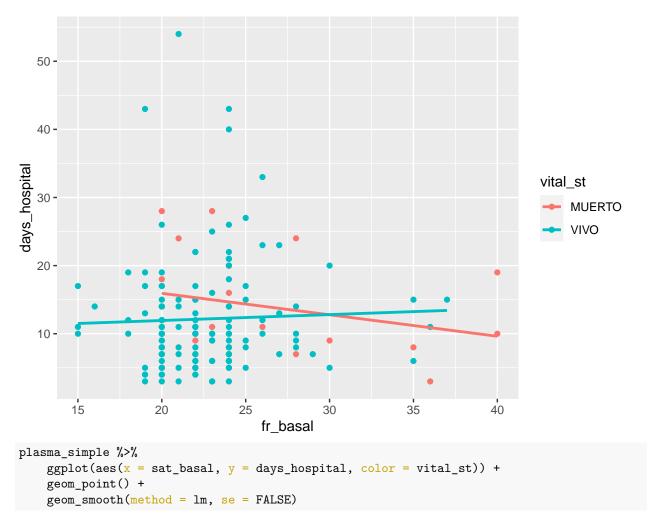


Predict hospital stay duration from saturation or respiratory frequency?

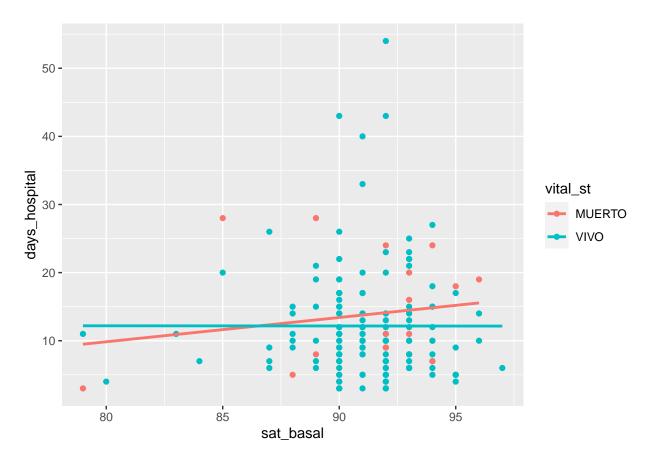
No.

```
plasma_simple %>%
    ggplot(aes(x = fr_basal, y = days_hospital, color = vital_st)) +
    geom_point() +
    geom_smooth(method = lm, se = FALSE)
```

$geom_smooth()$ using formula 'y ~ x'



`geom_smooth()` using formula 'y ~ x'



First models:

- Differences were computed by (last measurement basal).
- All models controll for age, sex, death or alive, hospital, number of days from: symptoms to hospitalization and symptoms to treatment.
- Treatment has a positive effect on saturation difference and respiratory frequency difference.

```
sat_mod <- lm(sat_diff ~ age + sex + group + vital_st + hospital +</pre>
   nd_sint_tto + nd_sint_ing, data = plasma_simple)
sat_mod %>%
    summary()
##
## Call:
## lm(formula = sat_diff ~ age + sex + group + vital_st + hospital +
##
       nd_sint_tto + nd_sint_ing, data = plasma_simple)
##
## Residuals:
        Min
##
                  1Q
                       Median
                                     3Q
                                             Max
## -23.3454 -2.2764 -0.1697
                                1.5997 14.5166
##
## Coefficients:
                 Estimate Std. Error t value Pr(>|t|)
##
                            2.620488
                                      -1.570 0.11870
## (Intercept) -4.113478
                            0.026989 -2.739 0.00695 **
## age
                -0.073926
```

```
## sexMASCULINO -0.219291
                           0.749964 -0.292 0.77041
                0.006302 0.702030 0.009 0.99285
## group
## vital stVIVO 5.967895
                                      4.555 1.12e-05 ***
                          1.310061
                           1.180252
                                    -0.184 0.85403
## hospitalHPAS -0.217539
## hospitalIESS 1.816690
                           1.100843
                                      1.650 0.10110
## nd sint tto
                0.106569
                           0.074420
                                      1.432 0.15434
## nd_sint_ing
                0.003670
                           0.015091
                                      0.243 0.80822
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 4.11 on 142 degrees of freedom
    (7 observations deleted due to missingness)
## Multiple R-squared: 0.2361, Adjusted R-squared: 0.1931
## F-statistic: 5.486 on 8 and 142 DF, p-value: 4.853e-06
fr_mod <- lm(fr_diff ~ age + sex + group + vital_st + hospital +</pre>
  nd_sint_tto + nd_sint_ing, data = plasma_simple)
fr_mod %>%
   summary()
##
## Call:
## lm(formula = fr_diff ~ age + sex + group + vital_st + hospital +
##
      nd_sint_tto + nd_sint_ing, data = plasma_simple)
##
## Residuals:
##
       Min
                 1Q
                      Median
                                   30
## -16.6416 -2.2233 0.3685
                               2.1403 22.4224
##
## Coefficients:
                Estimate Std. Error t value Pr(>|t|)
## (Intercept) -1.097761
                           3.195110 -0.344 0.73167
                0.002846
                          0.032907
                                      0.086 0.93121
## age
## sexMASCULINO 0.251937
                          0.914417
                                     0.276 0.78332
## group
               -0.979610
                          0.855971 -1.144 0.25437
## vital_stVIVO -7.292668
                           1.597332
                                     -4.566 1.07e-05 ***
## hospitalHPAS 1.986333
                           1.439058
                                      1.380 0.16966
## hospitalIESS 2.985088
                          1.342236
                                      2.224 0.02773 *
## nd_sint_tto
                0.282053
                           0.090738
                                      3.108 0.00227 **
## nd_sint_ing
                0.018287
                           0.018400
                                      0.994 0.32198
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5.011 on 142 degrees of freedom
     (7 observations deleted due to missingness)
## Multiple R-squared: 0.2443, Adjusted R-squared: 0.2017
## F-statistic: 5.738 on 8 and 142 DF, p-value: 2.486e-06
```

Distributions per Hospital

```
plasma_simple %>%
   ggplot() +
   geom_density(aes(sat_diff, fill = hospital), position = 'dodge', alpha = 0.3) +
```

```
labs(x = 'saturation difference')

## Warning: Removed 7 rows containing non-finite values (stat_density).

## Warning: Width not defined. Set with 'position_dodge(width = ?)'

0.15-

| hospital HGDC HPAS IESS
```

```
plasma_simple %>%
   ggplot() +
   geom_density(aes(fr_diff, fill = hospital), position = 'dodge', alpha = 0.3) +
   labs(x = 'respiratory frequency difference')
```

0

10

Warning: Removed 7 rows containing non-finite values (stat_density).

-10

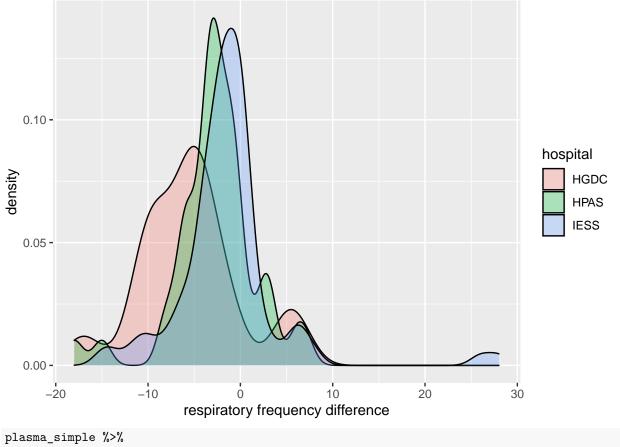
saturation difference

Warning: Width not defined. Set with `position_dodge(width = ?)`

0.00 -

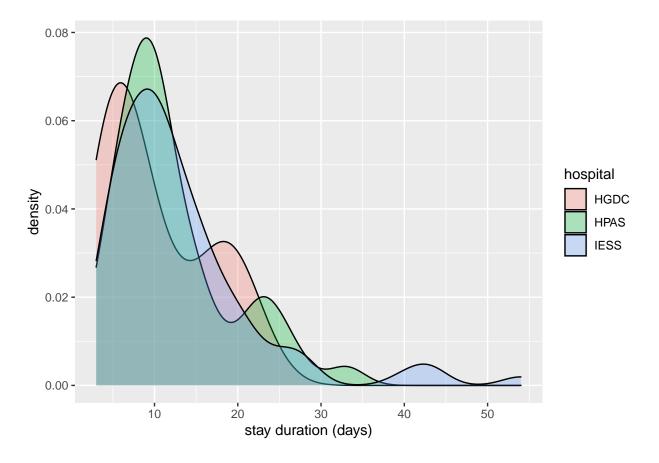
-30

-20



```
plasma_simple %>%
    ggplot() +
    geom_density(aes(days_hospital, fill = hospital), position = 'dodge', alpha = 0.3) +
    labs(x = 'stay duration (days)')
```

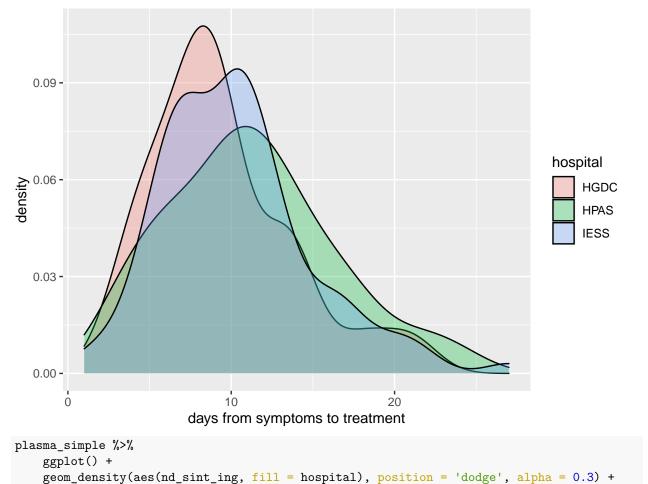
Warning: Width not defined. Set with `position_dodge(width = ?)`



Treatment per Hospital

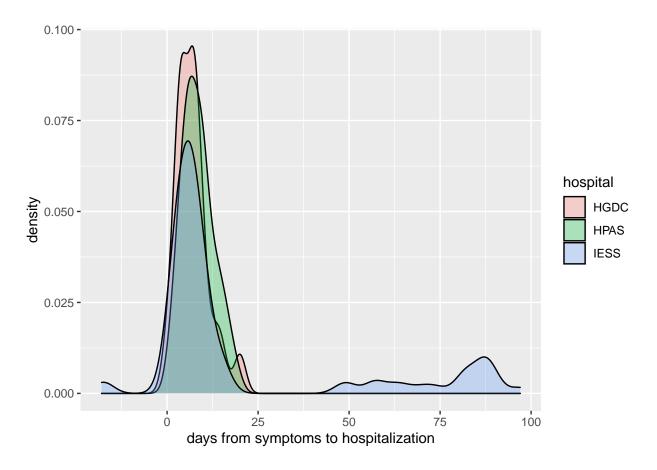
```
plasma_simple %>%
    ggplot() +
    geom_density(aes(nd_sint_tto, fill = hospital), position = 'dodge', alpha = 0.3) +
    labs(x = 'days from symptoms to treatment')
```

Warning: Width not defined. Set with `position_dodge(width = ?)`



labs(x = 'days from symptoms to hospitalization')

Warning: Width not defined. Set with `position_dodge(width = ?)`



Missing data.

```
plasma_simple %>%
    filter(is.na(sat_final)) %>%
    count()
## # A tibble: 1 x 1
##
##
     <int>
## 1
# missing data:
plasma_simple %>%
    group_by(hospital) %>%
    mutate(n_{hosp} = n()) \%
    filter(is.na(sat_final)) %>%
    group_by(hospital, group) %>%
    summarise(n_na = n(), n_hosp = max(n_hosp)) %>%
    mutate(percent = 100*(n_na / n_hosp)) %>%
    select(hospital, group, percent) %>%
    pivot_wider(names_from = group, values_from = percent)
## `summarise()` has grouped output by 'hospital'. You can override using the `.groups` argument.
## # A tibble: 2 x 3
               hospital [2]
## # Groups:
              `0` `1`
    hospital
```

```
## <chr> <dbl> <dbl> <dbl> =# 1 HGDC 8.70 13.0 ## 2 HPAS 4.65 NA
```

Hospitalization Days and Probability of Survival

```
glm(days_hospital ~ age + sex + group + vital_st + hospital +
   nd_sint_tto + nd_sint_ing, data=plasma_simple,
   family = poisson(link='log')) %>%
   summary()
##
## Call:
## glm(formula = days_hospital ~ age + sex + group + vital_st +
      hospital + nd_sint_tto + nd_sint_ing, family = poisson(link = "log"),
##
      data = plasma_simple)
##
## Deviance Residuals:
      Min
                1Q
                    Median
                                 3Q
                                         Max
## -3.5011 -1.5109 -0.5000
                            0.7748
                                      8.5560
## Coefficients:
##
                Estimate Std. Error z value Pr(>|z|)
                ## (Intercept)
## age
                0.008894
                         0.001849
                                    4.810 1.51e-06 ***
## sexMASCULINO -0.057516
                          0.051802 -1.110
                                            0.2669
               -0.077812
## group
                          0.047261 -1.646
                                             0.0997 .
## vital_stVIVO -0.110625
                         0.070596 -1.567
                                             0.1171
## hospitalHPAS 0.150331
                          0.079593
                                    1.889
                                             0.0589 .
## hospitalIESS 0.302426
                          0.072200
                                    4.189 2.81e-05 ***
## nd_sint_tto
                0.034565
                           0.004703
                                     7.350 1.99e-13 ***
                           0.001214 -6.042 1.53e-09 ***
## nd_sint_ing -0.007332
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## (Dispersion parameter for poisson family taken to be 1)
##
      Null deviance: 711.58 on 157 degrees of freedom
## Residual deviance: 588.23 on 149 degrees of freedom
## AIC: 1268.4
##
## Number of Fisher Scoring iterations: 5
coxph(Surv(days_hospital, status) ~ age + sex + group + hospital +
   nd_sint_tto + nd_sint_ing + hta + obesity + dm,
   data=plasma_simple) %>%
   summary()
## Call:
## coxph(formula = Surv(days_hospital, status) ~ age + sex + group +
##
      hospital + nd_sint_tto + nd_sint_ing + hta + obesity + dm,
##
      data = plasma_simple)
##
    n= 127, number of events= 15
##
##
      (31 observations deleted due to missingness)
```

```
##
                   coef exp(coef) se(coef)
##
                                                z Pr(>|z|)
                0.11154
                          1.11800 0.03746 2.977 0.00291 **
## sexMASCULINO 2.94985 19.10306
                                   1.21499 2.428
                                                   0.01519 *
## group
                0.61220
                          1.84449
                                   0.67647 0.905
                                                   0.36547
## hospitalHPAS -1.42113
                         0.24144 0.87274 -1.628
                                                   0.10345
## hospitalIESS -2.45772
                          0.08563
                                   1.07514 -2.286
                                                   0.02226 *
## nd_sint_tto -0.11253
                          0.89357
                                   0.06816 -1.651
                                                   0.09873
## nd_sint_ing 0.04525
                          1.04628
                                   0.01702 2.658
                                                   0.00786 **
## htaSI
               -0.61279
                          0.54184
                                   0.88428 -0.693
                                                   0.48832
## obesitySI
               -0.14497
                          0.86505 0.94739 -0.153 0.87838
                0.49126
                          1.63438 0.81802 0.601 0.54814
## dmSI
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
##
                exp(coef) exp(-coef) lower .95 upper .95
## age
                            0.89445
                 1.11800
                                      1.03885
                                                 1.2032
## sexMASCULINO 19.10306
                            0.05235
                                      1.76563
                                              206.6833
## group
                 1.84449
                            0.54216
                                      0.48985
                                                 6.9453
## hospitalHPAS
                 0.24144
                            4.14182
                                      0.04364
                                                 1.3356
## hospitalIESS
                 0.08563
                          11.67815 0.01041
                                                 0.7043
                 0.89357
## nd_sint_tto
                           1.11911
                                      0.78183
                                                 1.0213
## nd sint ing
                 1.04628
                            0.95576
                                      1.01195
                                                 1.0818
                                      0.09576
## htaSI
                 0.54184
                            1.84557
                                                 3.0660
## obesitySI
                 0.86505
                            1.15601
                                      0.13509
                                                 5.5394
## dmSI
                 1.63438
                            0.61185
                                      0.32889
                                                 8.1218
##
## Concordance= 0.893 (se = 0.033)
## Likelihood ratio test= 27.81 on 10 df,
                                            p=0.002
                                            p=0.06
## Wald test
                       = 17.76 on 10 df,
## Score (logrank) test = 25.27 on 10 df,
                                            p=0.005
coxph(Surv(days_hospital, status) ~ group,
    data=plasma_simple) %>%
    summary()
## Call:
## coxph(formula = Surv(days_hospital, status) ~ group, data = plasma_simple)
##
    n= 158, number of events= 19
##
##
##
            coef exp(coef) se(coef)
                                        z Pr(>|z|)
## group 0.003121 1.003126 0.477064 0.007
                                             0.995
##
##
         exp(coef) exp(-coef) lower .95 upper .95
## group
                      0.9969
            1.003
                                0.3938
                                           2.555
##
## Concordance= 0.57 (se = 0.07)
## Likelihood ratio test= 0 on 1 df,
## Wald test
                       = 0 on 1 df,
                                       p=1
## Score (logrank) test = 0 on 1 df,
                                       p=1
survfit(Surv(days_hospital, status) ~ group, data=plasma_simple) %>%
   broom::tidy() %>%
    ggplot(aes(time, estimate, color = strata)) +
```

