STAT796_HW1

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January 23, 2019

1. Import Data

2. rename and recode

3. summary stats for entire cohort

3a. average, max, min, sc of age (years)

3b. count of subjects with each service type

```
##
## medical surgical
## 93 107
```

3c. count of subjects with procedure type

```
##
## elective emergency
## 53 147
```

3d. count of subjects of each race and each sex

```
##
## male female
## white 108 67
## black 10 5
## other 6 4
```

3e. count of subject by consciousness

```
## ## no_coma_or_stupor stupor coma ## 185 5 10
```

4. summary stats for cohort, grouped by by vital status

4a. average, max, min, sd age

```
## # A tibble: 2 x 6
## Died n mean_age max_age min_age sd_age
## <int> <dbl> <dbl> <dbl>
```

```
## 1 0 160 55.650 91 16 20.42818
## 2 1 40 65.125 92 19 16.64900
```

4b. count by service type

```
##
## elective emergency
## 53 147
```

4c. count by procedure type

```
##
## elective emergency
## 53 147
```

4d. count by race and sex

```
##
## male female
## white 108 67
## black 10 5
## other 6 4
```

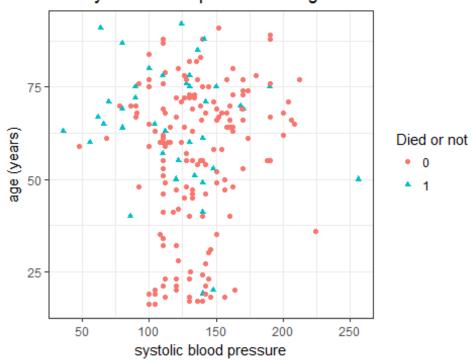
4e. count by consciousness

```
##
## no_coma_or_stupor stupor coma
## 185 5 10
```

5. graphical summaries

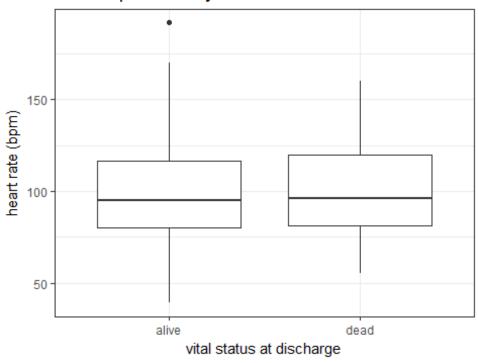
5a. scatterplot of sbp against age

5a - systolic blood pressure vs age



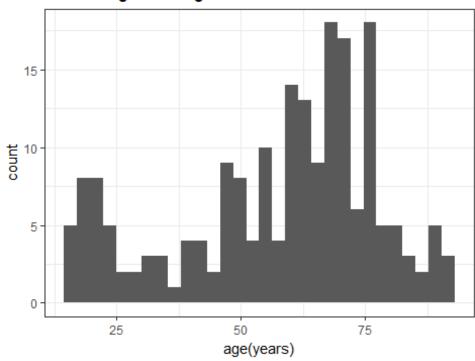
5b. boxplots of hr, grouped by consciousness

5b - boxplot of hr by consciousness



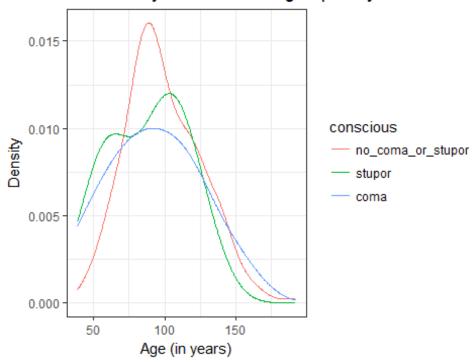
5c. histograms of age

5c - histogram of age



5d. densit estimates of hr, grouped by consciousness

5d - density estimate of hr, grouped by consciousnes



STAT796_HW1_code

Colleen Brents

January 23, 2019

```
library(ggplot2)
library(dplyr)
library(tidyr)
library(readr)
library(knitr)
```

1. Import Data

```
url <-
paste0("https://raw.githubusercontent.com/cbrents/stat796logreg/master/hw1/IC
U_raw.txt")
icu_raw<-read_table2(url)</pre>
```

2. rename and recode

```
icu<- icu raw %>%
    transmute(id=ID,
              died=STA,
              age=AGE,
              female=GENDER,
              race=factor(RACE,
                           levels=1:3,
                           labels=c("white", "black", "other")),
              service=factor(SER,
                              levels=0:1,
                              labels=c("medical", "surgical")),
              cancer=CAN,
              renal=CRN,
              infect=INF,
              cpr=CPR,
              sbp=SYS,
              hr=HRA,
              prv admit=PRE,
              type=factor(TYP,
                           levels=0:1,
                           labels=c("elective", "emergency")),
              fracture=FRA,
              po2_g60=P02,
              '1-P02'= po2 g60,
              pco2_g45=PCO,
              bicarb_g18=BIC,
              '1 - BIC' = bicarb_g18,
              creat_g2=CRE,
```

```
conscious=factor(LOC,
    levels=0:2,
    labels=c("no_coma_or_stupor", "stupor", "coma")))
```

3. summary stats for entire cohort

```
3a. average, max, min, sc of age (years)
```

3b. count of subjects with each service type

```
a<- table (icu$service)
print(a)
##
## medical surgical
## 93 107</pre>
```

3c. count of subjects with procedure type

```
b<- table(icu$type)
print(b)

##

## elective emergency
## 53 147</pre>
```

3d. count of subjects of each race and each sex

```
#count(icu, race, female)
z<- table(icu$race, icu$female)</pre>
colnames(z)<- c("male", "female")</pre>
print(z)
##
##
           male female
##
     white 108
                    67
##
     black 10
                     5
##
     other 6
                      4
```

```
3e. count of subject by consciousness
```

```
#count(icu, conscious)
x<- table(icu$conscious)
print(x)
##
## no_coma_or_stupor stupor coma
## 185 5 10</pre>
```

4. summary stats for cohort, grouped by by vital status

```
4a. average, max, min, sd age
```

```
age table <- icu %>%
 group by(Died=died) %>%
 summarize(n=n(),
          mean_age = mean(age),
           max_age = max(age),
           min_age = min(age),
           sd_age =sd(age)) # died == 0 alive == 1
print(age_table)
## # A tibble: 2 x 6
##
     Died
              n mean_age max_age min_age
                                           sd age
##
    <int> <int>
                  <dbl> <dbl> <dbl> <dbl>
                                         <dbl>
            160
## 1
        0
                  55.650
                              91
                                      16 20.42818
## 2
       1
             40
                  65.125
                              92
                                    19 16.64900
```

4b. count by service type

```
vitals <- icu %>%
   group_by(died)
y<- table(vitals$type)
print(y)

##
## elective emergency
## 53 147</pre>
```

4c. count by procedure type

```
a<- table(vitals$type)
print(a)
##
## elective emergency
## 53 147</pre>
```

4d. count by race and sex

```
z<- table(vitals$race, vitals$female)
colnames(z)<- c("male", "female")
print(z)</pre>
```

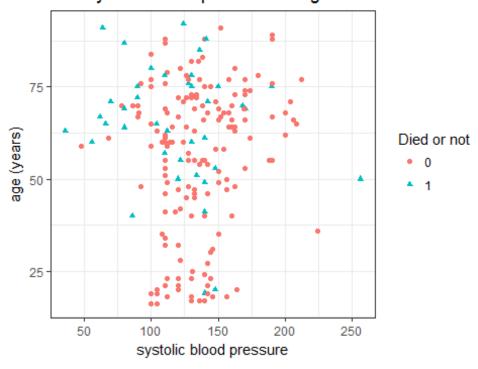
```
##
## male female
## white 108 67
## black 10 5
## other 6 4
```

4e. count by consciousness

5. graphical summaries

5a. scatterplot of sbp against age

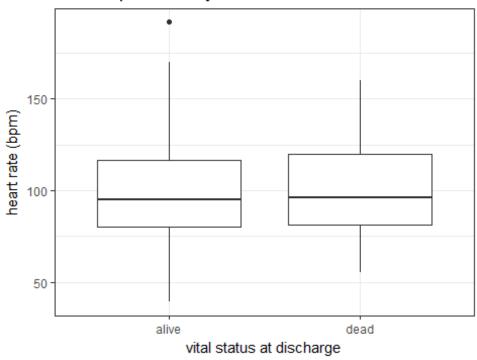
5a - systolic blood pressure vs age



5b. boxplots of hr, grouped by consciousness

```
ggplot(icu) +
  theme_bw() +
  geom_boxplot(aes(group = as.factor(died), x=as.factor(died), y= hr)) +
  xlab("vital status at discharge") +
  ylab("heart rate (bpm)") +
  ggtitle("5b - boxplot of hr by consciousness") +
  scale_x_discrete(labels = c("alive", "dead"))
```

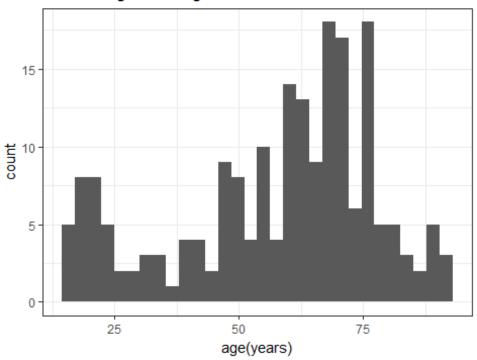
5b - boxplot of hr by consciousness



5c. histograms of age

```
ggplot(icu) +
  theme_bw() +
  geom_histogram(aes(x=age)) +
  xlab("age(years)") +
  ylab("count") +
  ggtitle("5c - histogram of age")
```

5c - histogram of age



5d. densit estimates of hr, grouped by consciousness

```
ggplot(icu) +
  theme_bw() +
  geom_line(aes(x=hr, col=as.factor(conscious), group=conscious),
stat="density") +
  xlab("Age (in years)") +
  ylab("Density") +
  scale_color_discrete("conscious") +
  ggtitle("5d - density estimate of hr, grouped by consciousness") #from
notes
```

5d - density estimate of hr, grouped by consciousnes

