## Usage Lab: Red Cells (Ketju)

```
library(forecast)
library(ggplot2)
library(gridExtra)
library(knitr)
library(readxl)
library(plyr)
library(lubridate)
library(numbers)
source("src/evalhelp.R")
```

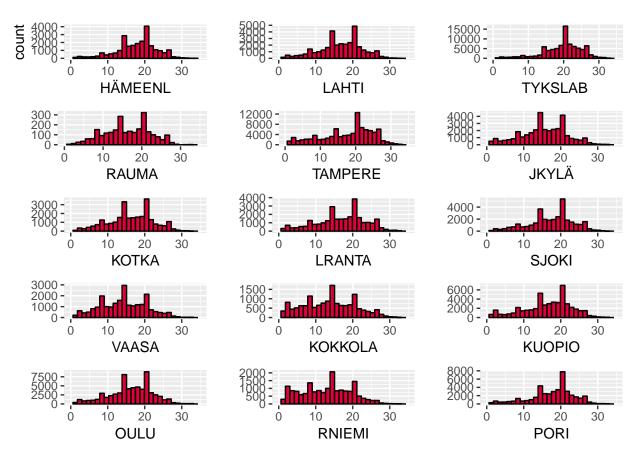
#### Intro

Currently under development. In this notebook we study how to use hospital blood product usage data to create demand predictions.

### Create original datasets that should remain immutable throughout labbing

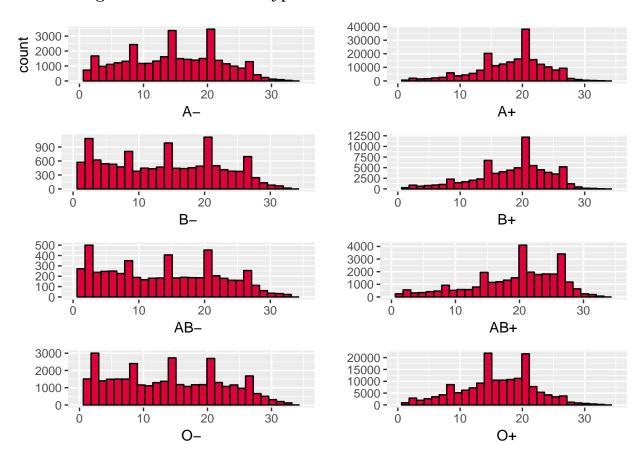
```
# Load data
# All deliveries
deliv <- read_excel("./data/ketju_data.xlsx", sheet = "Ketju-punasolutoimitukset 2014-")[, c('Päivämäär
colnames(deliv) <- c("time", "deliveries") # Change column names</pre>
deliv$time <- as.Date(deliv$time)</pre>
# Ketju usage 2014 -->
# I'm using read.csv() instead of read_excel() here, because this sheet contains some fields that kills
ketju <- read.csv("./data/ketju_data.csv", header = TRUE, sep = ",", colClasses=c("NULL", NA, "NULL", ")</pre>
colnames(ketju) <- c("hospital", "type", "time", "exp", "pcs") # Change column names</pre>
# Ensure compliant time format with lubridate
ketju$time <- mdy(ketju$time)</pre>
ketju$exp <- mdy(ketju$exp) # This will produce an error "failed to parse" for fields that aren't date
# Arrange by time
ketju <- arrange(ketju, time)</pre>
# Find usage
usage <- aggregate(ketju$pcs, by = list(ketju$time), sum); colnames(usage) <- c("time", "pcs")
```

### Histograms of how fresh blood is used across hospitals



The x-axis can be read as "days until an used product would've expired", so a larger number corresponds to a fresher product. We see a similar pattern across all hospitals: products are used in a manner that resembles a "weekly cycle". Most products are used 21 days from expiration, then 14 days, then 7 days...

### Same histogram but with blood types



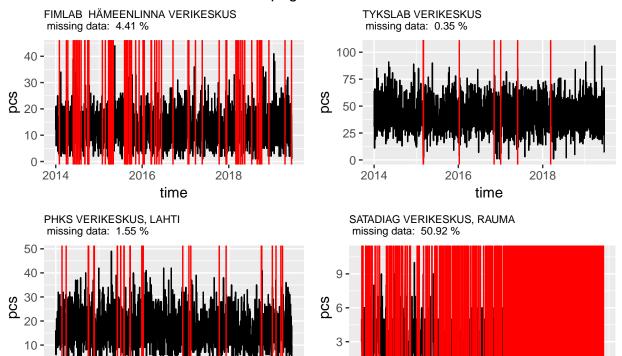
Here we observe the same weekly pattern, but the shape of + products is somewhat different from - products. Older products are used more with - products.

#### Series of usage

```
# Create a convenience vector for hospital tags
hospitals <-c("FIMLAB HÄMEENLINNA VERIKESKUS", "PHKS VERIKESKUS, LAHTI", "TYKSLAB VERIKESKUS", "SATADI.
              "FIMLAB TAMPERE VERIKESKUS", "FIMLAB VERIKESKUS, JYVÄSKYLÄ", "KYMKS VERIKESKUS, KOTKA", "
              "SEINÄJOEN KS VERIKESKUS", "VAASAN KS VERIKESKUS", "NORDLAB KOKKOLA VERIKESKUS", "ISLAB K
              "NORDLAB OULU VERIKESKUS", "NORDLAB ROVANIEMI VERIKESKUS", "SATADIAG VERIKESKUS, PORI")
plots <- list()
i = 0
for(hospital in hospitals){
   i <- i + 1
  hospital.data <- ketju[ketju$hospital == hospital, ]
  hospital.usage <- aggregate(hospital.data$pcs, by = list(hospital.data$time), sum)
  colnames(hospital.usage) <- c("time", "pcs")</pre>
  temp <- make_whole(hospital.usage)</pre>
  hospital.whole <- temp[[1]]
  hospital.missing <- temp[[2]]
  # Plot
```

```
hospital.plot <- ggplot() +
    geom_line(data = hospital.whole, aes(x = time, y = pcs)) +
    geom_vline(xintercept = hospital.missing, color = "red") +
    xlab("time") +
    ggtitle(paste(hospital, "\n missing data: ", round(length(hospital.missing)/length(hospital.whole$p theme(plot.title = element_text(size = 8))
    plots[[i]] <- hospital.plot
}
ml <- marrangeGrob(plots, nrow=2, ncol=2)
ml</pre>
```

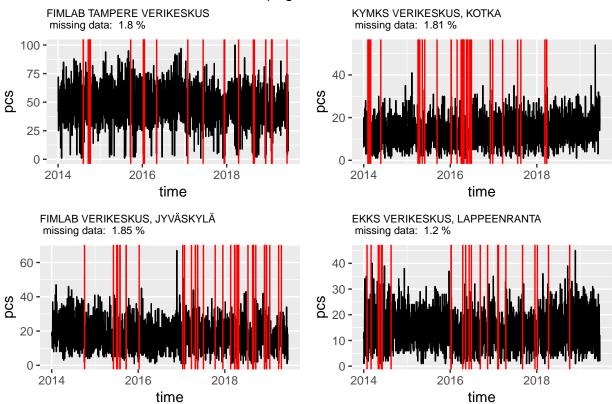
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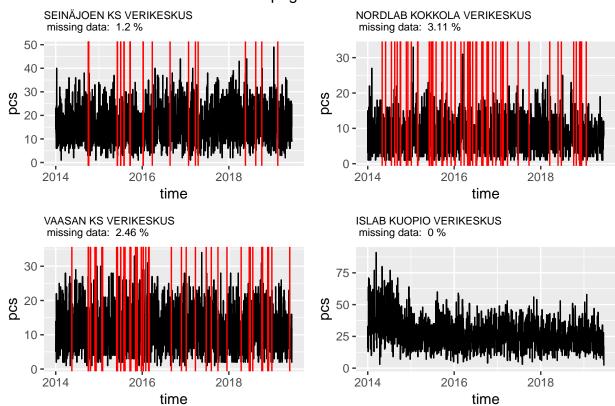
time

time

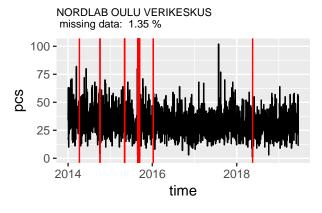
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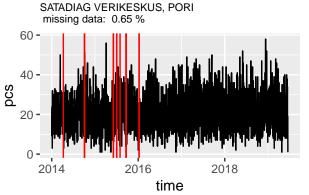


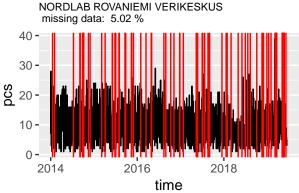
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These are daily red cell product usage data from each hospital. A red line means a missing data point. The missing data point might just be a zero, which is most probable with RAUMA. Let's limit our explorations to 2019 for now. This probably means we'll have to exclude RAUMA.

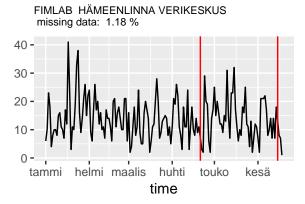
```
hospitals <- c("FIMLAB HÄMEENLINNA VERIKESKUS", "PHKS VERIKESKUS, LAHTI", "TYKSLAB VERIKESKUS",
              "FIMLAB TAMPERE VERIKESKUS", "FIMLAB VERIKESKUS, JYVÄSKYLÄ", "KYMKS VERIKESKUS, KOTKA", "
              "SEINÄJOEN KS VERIKESKUS", "VAASAN KS VERIKESKUS", "NORDLAB KOKKOLA VERIKESKUS", "ISLAB K
              "NORDLAB OULU VERIKESKUS", "NORDLAB ROVANIEMI VERIKESKUS", "SATADIAG VERIKESKUS, PORI")
plots <- list()
i = 0
for(hospital in hospitals){
   i <- i + 1
  hospital.data <- ketju[ketju$hospital == hospital, ]
  hospital.usage <- aggregate(hospital.data$pcs, by = list(hospital.data$time), sum)
  colnames(hospital.usage) <- c("time", "pcs")</pre>
  hospital.usage <- hospital.usage[hospital.usage$time >= as.Date("2019-01-01"), ]
  temp <- make_whole(hospital.usage)</pre>
  hospital.whole <- temp[[1]]
  hospital.missing <- temp[[2]]
  # Plot
  hospital.plot <- ggplot() +
    geom_line(data = hospital.whole, aes(x = time, y = pcs)) +
    geom_vline(xintercept = hospital.missing, color = "red") +
    ggtitle(paste(hospital, "\n missing data: ", round(length(hospital.missing)/length(hospital.whole$p
```

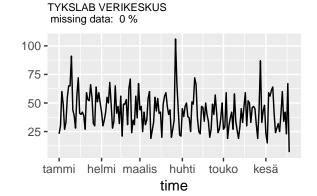
```
theme(plot.title = element_text(size = 8)) +
   ylab("")

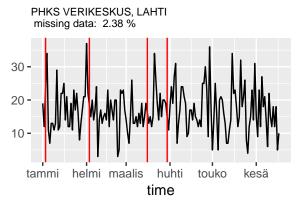
plots[[i]] <- hospital.plot
}

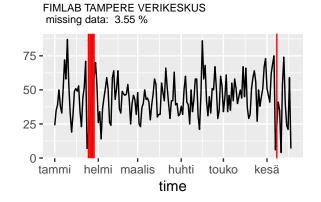
ml <- marrangeGrob(plots, nrow=2, ncol=2)
ml</pre>
```

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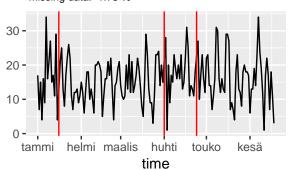




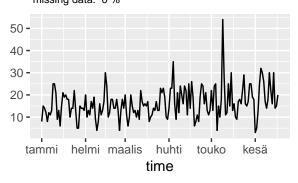


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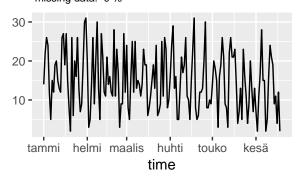
FIMLAB VERIKESKUS, JYVÄSKYLÄ missing data: 1.78 %



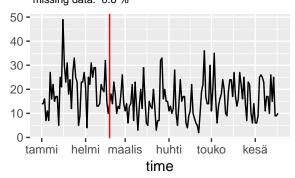
KYMKS VERIKESKUS, KOTKA missing data: 0 %



### EKKS VERIKESKUS, LAPPEENRANTA missing data: 0 %

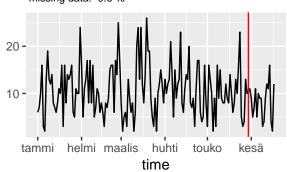


# SEINÄJOEN KS VERIKESKUS missing data: 0.6 %

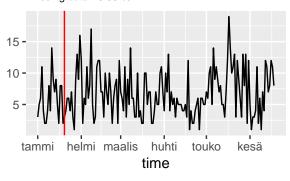


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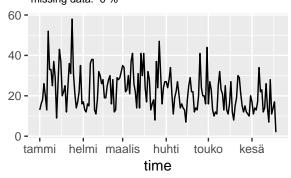
VAASAN KS VERIKESKUS missing data: 0.6 %



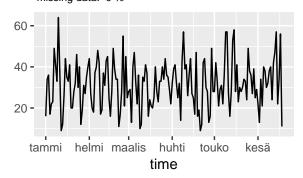
NORDLAB KOKKOLA VERIKESKUS missing data: 0.59 %



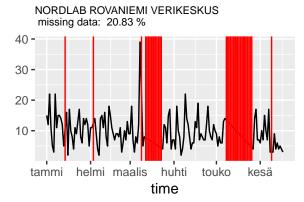
### ISLAB KUOPIO VERIKESKUS missing data: 0 %

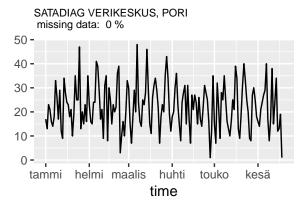


## NORDLAB OULU VERIKESKUS missing data: 0 %



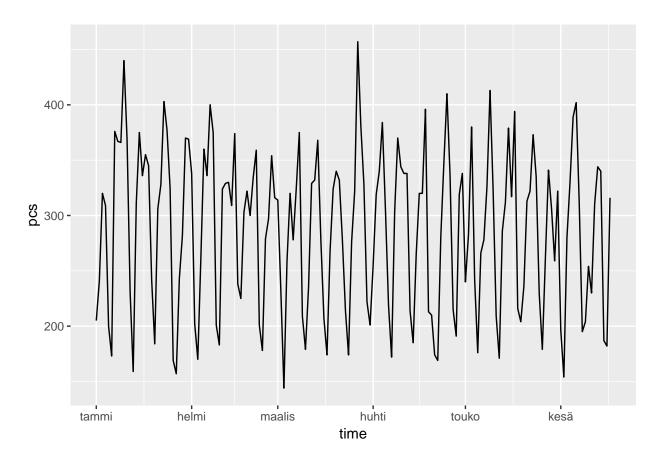
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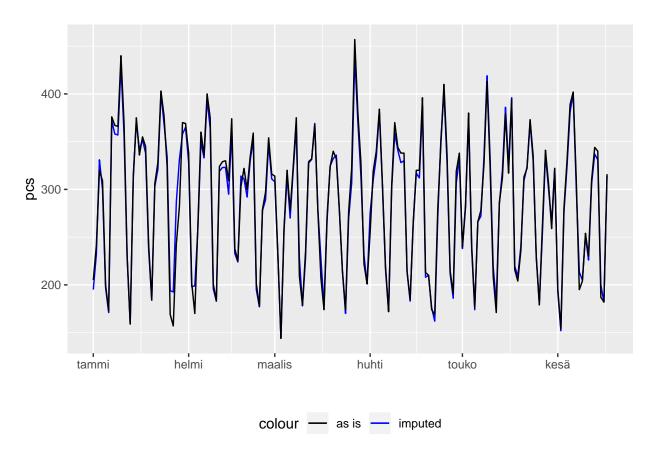


There is a "clear" weekly seasonality in most hospitals, and it is more pronounced where a lot of blood is used.

### Total usage across all hospitals (in 2019, without imputation)



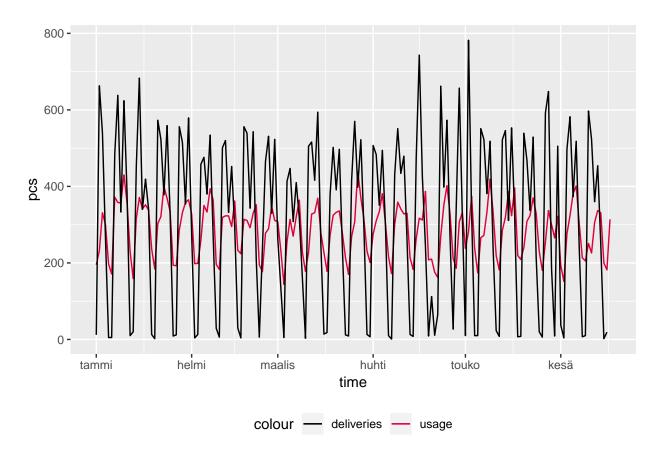
### Difference in total usage with and without imputation



The imputed series does not differ significantly from the raw series, so we can probably use either.

### Deliveries and usage 2019

```
ggplot() +
  geom_line(data = total.usage[total.usage$time >= "2019-01-01", ], aes(x = time, y = pcs, colour = "us
  geom_line(data = deliv[deliv$time >= "2019-01-01", ], aes(x = time, y = deliveries, colour = "deliver
  scale_colour_manual(values = c("black", "#DF013A")) +
  theme(legend.position = "bottom")
```



```
combined <- data.frame(usage = head(total.usage[total.usage$time >= "2018-01-01", ]$pcs, 521), deliveri
# Same week avg diff
diff0 <- 0
for(i in seq(from = 7, to = 469, by = 7)){
  diff0 <- diff0 + (sum(combined$usage[i:(i+6)]) - sum(combined$deliveries[i:(i+6)]))</pre>
}
avg0 \leftarrow diff0/67
# Next week sum
diff1 <- 0
for(i in seq(from = 7, to = 469, by = 7)){
 diff1 <- diff1 + (sum(combined$usage[i:(i+6)]) - sum(combined$deliveries[(i+7):(i+13)]))</pre>
}
avg1 <- diff1/67
# Second week sum
diff2 <- 0
for(i in seq(from = 7, to = 469, by = 7)){
  diff2 <- diff2 + (sum(combined$usage[i:(i+6)]) - sum(combined$deliveries[(i+14):(i+20)]))</pre>
}
avg2 \leftarrow diff2/67
# Third week sum
diff3 <- 0
for(i in seq(from = 7, to = 469, by = 7)){
```

Same week difference seems to be the smallest, but only by a very small margin. The difference constitutes a 10.79 % error, which is slightly higher than what we want. Let's go to the daily level and see what happens.

## Third week average diff: -213.69

```
# Same day avg diff
ddiff0 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
  ddiff0 <- ddiff0 + (combined$usage[i] - combined$deliveries[i])</pre>
}
davg0 <- ddiff0/462
# Next day diff
ddiff1 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
  ddiff1 <- ddiff1 + (combined$usage[i] - combined$deliveries[i+1])</pre>
}
davg1 \leftarrow ddiff1/462
# Second day diff
ddiff2 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
  ddiff2 <- ddiff2 + (combined$usage[i] - combined$deliveries[i+2])</pre>
}
davg2 \leftarrow ddiff2/462
# Third day diff
ddiff3 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
  ddiff3 <- ddiff3 + (combined$usage[i] - combined$deliveries[i+3])</pre>
}
davg3 <- ddiff3/462
# Fourth day diff
ddiff4 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
  ddiff4 <- ddiff4 + (combined$usage[i] - combined$deliveries[i+4])</pre>
davg4 \leftarrow ddiff4/462
# Fifth day diff
```

```
ddiff5 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
 ddiff5 <- ddiff5 + (combined$usage[i] - combined$deliveries[i+5])</pre>
davg5 <- ddiff5/462
# Sixth day diff
ddiff6 <- 0
for(i in seq(from = 7, to = 469, by = 1)){
 ddiff6 <- ddiff6 + (combined$usage[i] - combined$deliveries[i+6])</pre>
davg6 <- ddiff6/462
cat(paste("Same day average diff: ", round(davg0, digits = 2), "\n",
          "Next day average diff: ", round(davg1, digits = 2), "\n",
          "Second day average diff: ", round(davg2, digits = 2), "\n",
          "Third day average diff: ", round(davg3, digits = 2), "\n",
          "Fourth day average diff: ", round(davg4, digits = 2), "\n",
          "Fifth day average diff: ", round(davg5, digits = 2), "\n",
          "Sixth day average diff: ", round(davg6, digits = 2)
## Same day average diff: -30.52
```

## Same day average diff: -30.52

## Next day average diff: -31.74

## Second day average diff: -31.07

## Third day average diff: -29.97

## Fourth day average diff: -30.63

## Fifth day average diff: -30.25

## Sixth day average diff: -29.77

Same +10 % difference.

#### Forecasting usage

	$_{\mathrm{cMAPE}}$	MAPE	RMSE
DynReg	26.29518	17.23969	61.10245