# Hw-3

Jacob Lee

5 June 2017

### Homework 3

### Question 1

Our companies monthly revenues could be modeled via an exponential smoothing model. There is both a relationship to the previous level, the revenue trends and the seasonality (time of year) that determine the level of incomings.

### Question 2

For this question I perfomed an analysis on average monthly data and on the daily data. I selected what I thought appropriate levels of alpha, beta and gamma given the levels and then plotted a one year forecast.

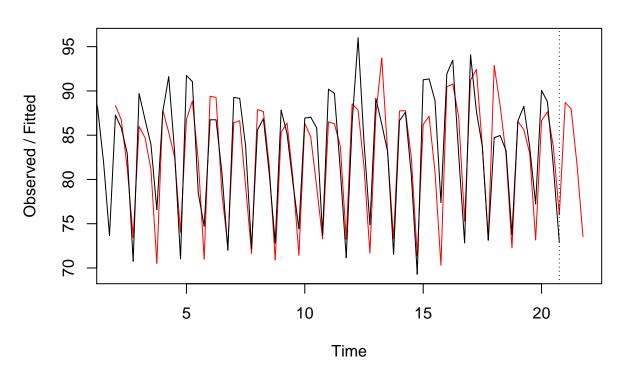
```
library('smooth')
```

```
## This is package "smooth", v1.9.9
temps = read.table('temps.txt', header = TRUE)
row.names(temps) <- temps$DAY</pre>
temps <- temps[2:21]
Monthly <- data.frame(M1=as.numeric(character()),</pre>
                        M2=as.numeric(character()),
                        M3=as.numeric(character()),
                        M4=as.numeric(character()))
mnths <- vector()</pre>
for(i in 1:20) {
  M1 <- mean(temps[1:31,i])
  M2 <- mean(temps[32:62,i])
  M3 <- mean(temps[63:92,i])
  M4 <- mean(temps[93:123,i])
  mnths \leftarrow c(mnths, M1, M2, M3, M4)
  Monthly <- rbind(Monthly, i=c(M1, M2, M3, M4))
}
monthly.ts <- ts(mnths, frequency = 4)
monthly.hw <- HoltWinters(monthly.ts, alpha=0.05, beta=0.1, gamma = 0.7)
preds <- predict(monthly.hw, n.ahead=(1*4))</pre>
series <- as.vector(temps[,1])</pre>
for(i in 2:20) {
  series<-c(series,temps[,i])</pre>
temps.ts <-ts(series, frequency = 123)</pre>
temps.hw <- HoltWinters(temps.ts, alpha = 0.2, beta = 0.0005, gamma = 0.5)
preds.days <- predict(temps.hw, n.ahead=(1*123))</pre>
```

Given we now have two reasonable models we can look at the forecast to determine if the end of summer is coming later.

plot(monthly.hw,preds)

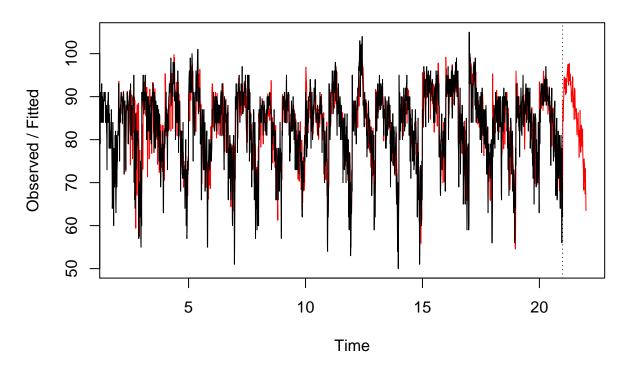
## **Holt-Winters filtering**



If summer is ending later we would expect in our forecast to see a increasing temperatures in the 3rd and 4th months of our period. This is not observed.

plot(temps.hw,preds.days)

### **Holt-Winters filtering**



This doesn't seem to me to be the best means of determining how long summer is, but lets compare our forecast year to previous results. If we define the end of summer to be the peak then summer peaks on the 35th day in our period. Previously we have seen the weather turn down after 20 days into the period, or as late as the 59th day in the period. Given this we cannot conclude summers are ending sooner.

### Question 3

We can use regression to predict house prices in the area. Features we may use are: \* average block size \* median income \* crime rate \* distance to cbd \* number of decent cafes

### Question 4

I will build a model with linear regression and simply normalise all data points

```
crime = read.table('uscrime.txt', header = TRUE)
library('caret')

## Loading required package: lattice

## Loading required package: ggplot2

preprocessParams <- preProcess(crime[1:15], method=c("scale","center"))

transX <- predict(preprocessParams, crime[1:15])

transCrime <- crime

transCrime[1:15] <- predict(preprocessParams, crime[1:15])

# linear regression

model.lm <- lm(Crime~., data=transCrime)</pre>
```

We can see a problem with this result, we would not expect to see a result of 155 on an example with a population to 150, this is a large population with an estimate below any we have seen in the actual dataset. We will need to try again, this time we will 1) do a log transform on the data - we identified this as possibly explaining the shape of the data in the previous example 2) remove highly correlated features and 3) perform cross-validation on our data.

```
# Log transformation
logCrime<-crime
logCrime[3:15] <-log(logCrime[3:15])</pre>
logCrime[1] <-log(logCrime[1])</pre>
# Evaluate Algorithms: Feature Selection
# remove correlated attributes
# find attributes that are highly corrected
set.seed(42)
cutoff <- 0.70
correlations <- cor(logCrime[1:15])</pre>
highlyCorrelated <- findCorrelation(correlations, cutoff=cutoff)</pre>
# create a new dataset without highly corrected features
dataset_features <- logCrime[,-highlyCorrelated]</pre>
# Run algorithms using 10-fold cross validation
control <- trainControl(method="repeatedcv", number=10, repeats=3)</pre>
metric <- "RMSE"
# lm
set.seed(42)
fit.lm <- train(Crime~., data=dataset_features,</pre>
                 method="lm", metric=metric, preProc=c("center", "scale"), trControl=control)
# Compare algorithms
summary(fit.lm)
##
## Call:
## lm(formula = .outcome ~ ., data = dat)
##
## Residuals:
##
       Min
                 1Q Median
                                  30
                                         Max
## -407.43 -128.99
                       6.72
                              99.48 387.44
##
```

```
##
               Estimate Std. Error t value Pr(>|t|)
                                     26.538 < 2e-16 ***
## (Intercept) 905.085
                             34.105
## M
                 92.253
                             53.055
                                      1.739
                                               0.0906 .
## So
                 72.334
                             76.389
                                      0.947
                                               0.3500
## Po2
                             55.573
                246.720
                                      4.440 8.21e-05 ***
## LF
                 84.904
                             57.346
                                      1.481
                                               0.1474
## M.F
                 68.676
                             59.080
                                      1.162
                                               0.2527
## Pop
                 -8.784
                             59.296
                                     -0.148
                                               0.8831
## NW
                 96.017
                             64.554
                                      1.487
                                               0.1456
## U2
                 63.083
                             47.356
                                      1.332
                                              0.1912
                                               0.0701
## Prob
               -124.041
                             66.450
                                     -1.867
## Time
                -24.631
                             60.777
                                    -0.405
                                               0.6877
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 233.8 on 36 degrees of freedom
## Multiple R-squared: 0.714, Adjusted R-squared: 0.6345
## F-statistic: 8.986 on 10 and 36 DF, p-value: 3.4e-07
fit.lm$finalModel
##
## Call:
## lm(formula = .outcome ~ ., data = dat)
## Coefficients:
##
   (Intercept)
                           Μ
                                       So
                                                    Po2
                                                                   I.F
##
       905.085
                      92.253
                                   72.334
                                                246.720
                                                               84.904
##
           M.F
                                                     U2
                         Pop
                                       NW
                                                                 Prob
##
        68.676
                      -8.784
                                   96.017
                                                 63.083
                                                            -124.041
##
          Time
##
       -24.631
So we have built two models with the coefficients and goodness of fit as above. Now we can plug in the values
features <- as.data.frame.numeric(c(14.0,0,10.0,12.0,15.5,0.640,94.0,150,1.1,0.120,3.6,3200,20.1,0.04,3
```

```
features <- t(features)</pre>
cn <- colnames(crime)[1:15]</pre>
colnames(features)<-cn</pre>
features[1] <- log(features[1])</pre>
features[3:15] <- log(features[3:15])</pre>
features (-highlyCorrelated)
features<-t(as.data.frame(features, row.names = colnames(dataset_features[1:10])))
result <- predict(fit.lm,features)</pre>
result
```

## features ## 1299.663

## Coefficients:

Here we end up with a estimate of 1300, from our descriptive analysis this is a lot close to what we would

The next step in developing this model would be to look at removing values with a very low p-value. Note that we only have 3 values + the intercept passing a signifigance test at alpha = 0.1.