Introduction to Machine Learning

Lab 3

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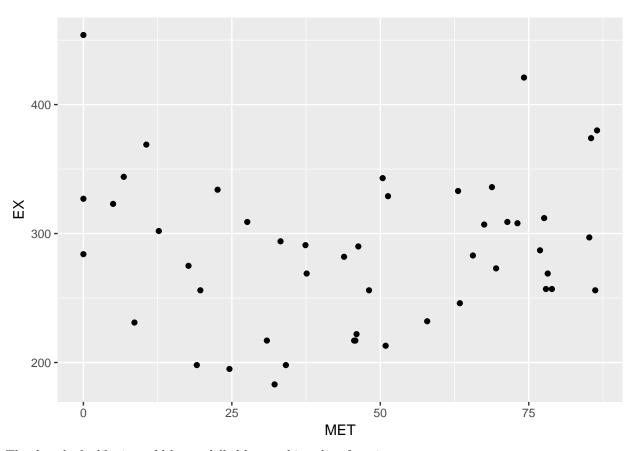
2016-11-24

Contents

Assignment 1	f 1
1	-
2	
3	
4	
5	
Assignment 2	2
1	-
2	
3	
4	
Appendix	
Code for As	ssignment 1
	ssignment 2

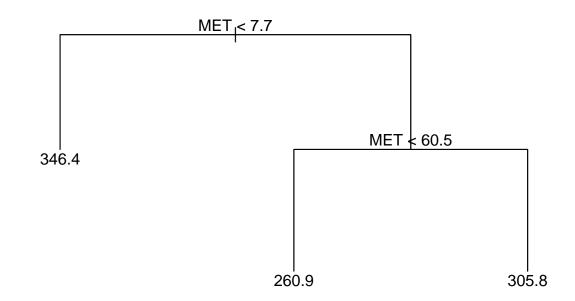
Assignment 1

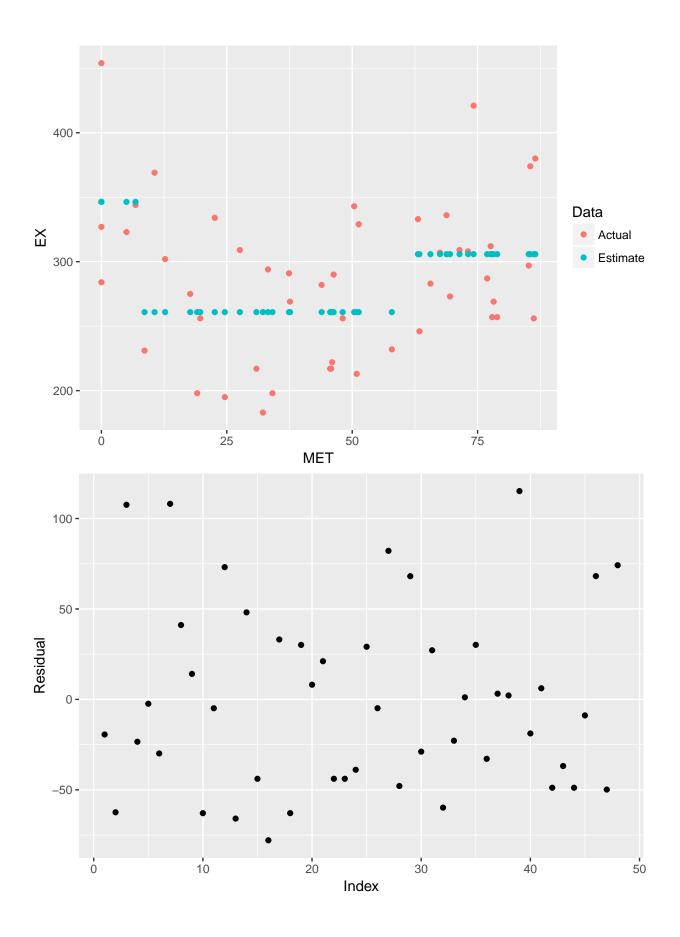
1



The data looks like it could be modelled by a cubic spline function.

 $\mathbf{2}$

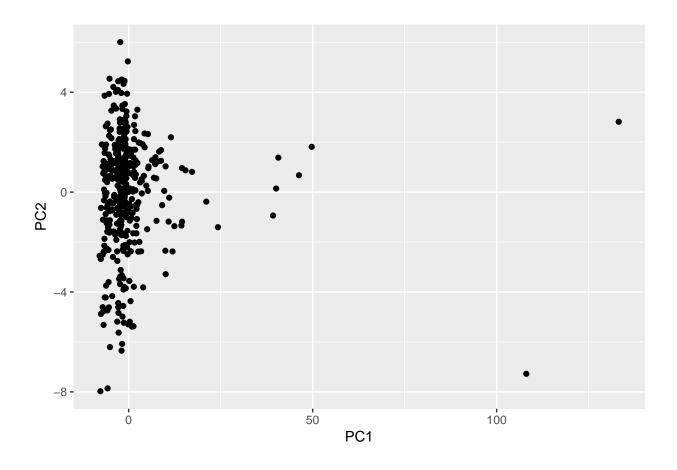


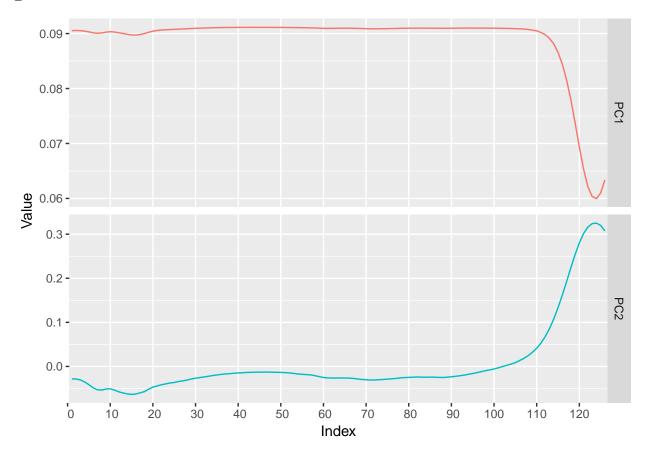


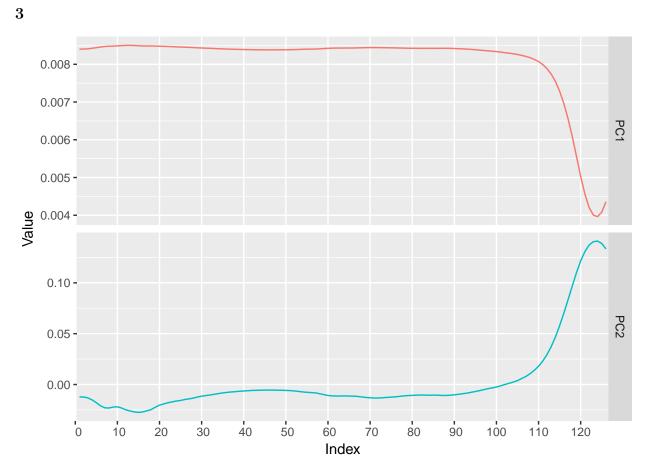
Assignment 2

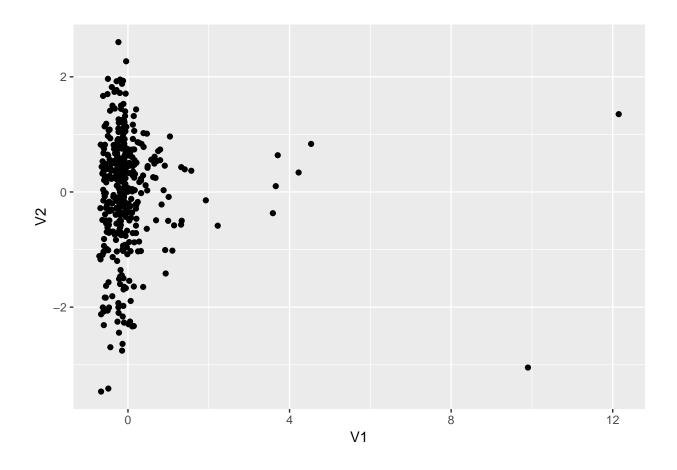
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   [121] "1.000" "1.000" "1.000" "1.000" "1.000" "1.000"
```

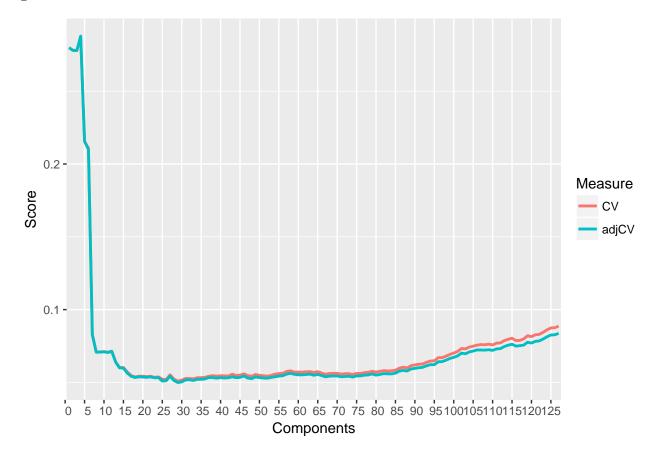












Appendix

Code for Assignment 1

```
library(ggplot2)
library(tree)
library(reshape)
data <- read.csv2(".../data/State.csv", header=TRUE, sep=";")</pre>
data <- data[order(data$MET),]</pre>
ggplot(data) +
    geom_point(aes(x=MET, y=EX))
tree.control(nobs=nrow(data), minsize=8)
treefit <- tree(EX ~ MET, data=data)</pre>
set.seed(12345)
treefit.cv <- cv.tree(treefit, FUN=prune.tree, K=10)</pre>
optimal_leaf_count<- treefit.cv$size[which.min(treefit.cv$dev)]</pre>
optimal_tree <- prune.tree(treefit, best=optimal_leaf_count)</pre>
plot(optimal_tree)
text(optimal_tree, pretty=0)
predicted <- predict(optimal_tree, data)</pre>
plot_data <- data.frame(MET=data$MET, Actual=data$EX, Estimate=predicted)</pre>
plot_data <- melt(plot_data, id.vars="MET")</pre>
names(plot_data) <- c("MET", "Data", "EX")</pre>
ggplot(plot_data) +
    geom_point(aes(x=MET, y=EX, color=Data))
residuals <- resid(optimal_tree)</pre>
plot_data <- data.frame(x=1:length(residuals), y=residuals)</pre>
ggplot(plot_data) +
    xlab("Index") +
    ylab("Residual") +
    geom_point(aes(x=x, y=y))
```

Code for Assignment 2

```
library(ggplot2)
library(fastICA)
library(pls)
library(reshape2)

data <- read.csv2(".../data/NIRSpectra.csv")

X <- scale(data[, -ncol(data)])
y <- data[, ncol(data)]</pre>
```

```
traceplot <- function(n, pc1, pc2) {</pre>
    plot_data <- data.frame(x=1:n, PC1=pc1, PC2=pc2)</pre>
    plot_data <- melt(plot_data, id="x")</pre>
    names(plot_data) <- c("Index", "Component", "Value")</pre>
    xlimits \leftarrow seq(0, n, by=10)
    ggplot(plot_data) +
        geom_line(aes(x=Index, y=Value, color=Component), show.legend=FALSE) +
        scale_x_discrete(limits=xlimits) +
        facet_grid(Component ~ ., scales="free")
}
pca <- prcomp(X)</pre>
## Eigenvalues
lambda <- pca$sdev^2</pre>
variances <- lambda / sum(lambda)</pre>
var99_comp_count <- which.max(cumsum(variances * 100) > 99)
components <- as.data.frame(pca$x[, 1:var99_comp_count])</pre>
U <- pca$rotation
traceplot(nrow(U), U[, 1], U[, 2])
set.seed(12345)
ica <- fastICA(X, var99_comp_count, alg.typ = "parallel", fun = "logcosh", alpha = 1,
                method = "R", row.norm = FALSE, maxit = 200, tol = 1e-06, verbose = FALSE)
W_prime <- ica$K %*% ica$W
components <- as.data.frame(ica$S)</pre>
traceplot(nrow(W_prime), W_prime[, 1], W_prime[, 2])
ggplot(components) +
    geom_point(aes(x=V1, y=V2))
set.seed(12345)
pcrfit <- pcr(Viscosity ~ ., data=data, scale=TRUE)</pre>
cvpcrfit <- crossval(pcrfit, segments=10, segment.type="random")</pre>
cv_scores <- t(matrix(MSEP(cvpcrfit)$val, nrow=2))</pre>
plot_data <- data.frame(cbind(1:ncol(data), cv_scores))</pre>
colnames(plot_data) <- c("Components", "CV", "adjCV")</pre>
plot_data <- melt(plot_data, id="Components", variable_name="Measure")</pre>
names(plot_data)[ncol(plot_data)] <- "Score"</pre>
xlimits <- seq(0, ncol(data), by=5)</pre>
ggplot(plot_data) +
    geom_line(aes(x=Components, y=Score, color=Measure), size=1) +
    scale_x_discrete(limits=xlimits)
```