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# R Code for HW7
                                         Sta108, Fall 2007, Utts
### Problem 9.25
Data=read.table("~/Documents/School/Sta108utts/APPENC01.txt")
#(b)
#Dataset used above (from file "APPENC01.txt") needs to be edited.
#1.Consider observations numbered: 57-113 in your dataset
dim (Data)
Data=Data [57:113,]
#2.Remove two predictors not needed for the analysis:
#these are categorical variables: "MedSchool", "Region" (Columns 8,9)
#3.remove ID column not needed for the analysis (Column 1)
Data=Data[, -c(1, 8, 9)]
#4.transform the response Y to: log10(Y)
Data$Stay=log10 (Data$Stay)
#Now, your dataset is clean to be used in the problem
Data
DataY=Data[,1] #separate response Y
DataX=Data[,-1] #separate predictors, X variables
pairs (DataX)
cor (DataX)
#(c) (NOTE THAT Y IS NOW THE LOG10 TRANSFORMED VERSION OF THE ORIGINAL Y, FROM #4 ABOVE)
library(leaps)
leaps(x=DataX, y=DataY,
     names=c("Age", "Risk", "Cultur", "Xray", "Beds", "Census", "Nurse", "Facility"),
     method="Cp")
#Three best models are chosen by 3 lowest Cp criterions
#To automatically print models in the increasing order of Cp criterion:
ModelSel = leaps(x=DataX, y=DataY,
      names=c("Age", "Risk", "Cultur", "Xray", "Beds", "Census", "Nurse", "Facility"),
      method="Cp")
ModelSel$which[ order( ModelSel$Cp ), ]
#To print Cp criterion in increasing order
sort( ModelSel$Cp )
#To plot Cp against p, and add reference line: Cp=p
plot( ModelSel$size, ModelSel$Cp, pch=19)
abline(0,1)
#Fit the best chosen model
Fit = lm(Stay ~ Age + Xray + Census, data=Data)
Fit
#Residual plot
plot(Fit$fitted.values, Fit$residuals, main="Residuals vs. Fitted Values", xlab="Fitted
Values", ylab="Residuals", pch=19)
abline(h=0)
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