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## this code reads a bivariate dataset on two variables X and Y
## performs simple linear regression of Y on X
## using Kendall-Theil (KT) regression line
## Also, computes the test statistic T to perform
## KT test for significance of regression
## does NOT compute p value for significance of regression
## this code asumes there are no ties in X values
## if this assumption is not true, this code will result in error
## read the data file that includes observations from two variables in
two-column format
dataset = as.matrix(read.table("folderpath/filename.extension", header = T or
 F))
## Input the positions of covariate column and response column
## as understood from the statement of the problem
covariate_column_position = ## 1 or 2 depending on which column is covariate
response_column_position = ## 1 or 2 depending on which column is response
## Accordingly, define the original (unsorted) X and Y values)
X_original = dataset[ ,covariate_column_position]
Y_original = dataset[ ,response_column_position]
## now pair them
Pairs_original = cbind(X_original,Y_original)
## calculate sample size
n = length(X original)
print(paste("Sample size = n = ",n,sep=""))
## Now sort the pairs according to values of X ONLY from smallest to largest
Pairs_sorted_using_X_original_only = Pairs_original[order(X_original), ]
## now create the pairwise slopes
## extract X and Y from sorted pairs
X = Pairs_sorted_using_X_original_only[ ,1]
Y = Pairs_sorted_using_X_original_only[ ,2]
## initiate the empty matrix of pairwise slopes
slope_matrix = matrix(" ",n,n)
## fill in only the positions above the diagonal line
for (i in 1:(n-1)) ## i-th row
    for (j in (i+1):n) ## j-th column
        ##j > i implies (i,j) position is above the diagonal
        slope matrix[i,j] = round((Y[j] - Y[i])/(X[j] - X[i]),2)
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}
}
## want to use the sorted pairs as row headers as well as column headers of
## the matrix of pairwise slopes
## initiate empty name fields for columns
colnames(slope_matrix) = array("",n)
## allocate sorted pairs as column names
for (i in 1:n) colnames(slope_matrix)[i] = paste("(",X[i],",",Y[i],")",sep="")
## do the same for row names as well
rownames(slope matrix) = colnames(slope matrix)
## Now print the matrix of pairwise slopes
print("----")
print("Matrix of pairwise slopes")
print("----")
print(noquote(slope matrix))
print("----")
## Now, to estimate the slope b, extract the values of
##pairwise slopes from above matrix
collection_of_pairwise_slopes =
as.numeric(slope matrix[!is.na(as.numeric(slope matrix))])
## calculate the median of pairwise slopes
b_hat = median(collection_of_pairwise_slopes)
print(paste("Estimated slope = b_hat = ",b_hat,sep=""))
## now, to estimate intercept a
## calculate a1,a2,...,an, as in classnote
a\_vector = round(Y - c(b\_hat)*X,2)
## compute the median of a1,a2,...,an
a_hat = median(a_vector)
print(paste("Estimated intercept = a_hat = ",a_hat,sep=""))
## plot the estimated regression line and the data points
x11()
## plot the observations first
plot(X,Y,type="p",col="RED",xlab
=colnames(dataset)[covariate_column_position],
ylab=colnames(dataset)[response_column_position], main="")
## now draw a straight line with intercept a_hat and slope b_hat
abline(a=a_hat,b=b_hat,col="BLUE")
## prediction for a new observation
## specify the covariate value for the new observation as X new
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X_new =
# Predict the median of response value for the new observation
## using formula for estimated regression line
median_Y_new_prediction = a_hat + b_hat*X_new
print(paste("Predicted median of the distribution of Y for the new observation
 = ",median_Y_new_prediction,sep=""))
## Now, test for significance of regression
## the test statistic is difference between
## number of positive pairwise slopes and
## number of negative pairwise slopes
Number of positive pairwise slopes =
 length(which(collection_of_pairwise_slopes>0))
Number_of_negative_pairwise_slopes =
 length(which(collection of pairwise slopes<0))</pre>
T = Number_of_positive_pairwise_slopes - Number_of_negative_pairwise_slopes
print(paste("Test Statistic = T = ",T,sep=""))
## now, compute p value using appropriate probability tables
## as mentioned in class note
```