

dataTransformations

September 26, 2019

0.1 data transformations

The regression function we have explored take the form,

$$p(y|X) \sim N(X\beta, \sigma^2),$$

and this model makes some important assumptions. * Our response (y) is linearly related to X . * The observations (y_i, x_i) are independent from one another. * The conditional probability of our response y is normally distributed. * The same σ applies to all values of X . This is called homoskedasticity of errors.

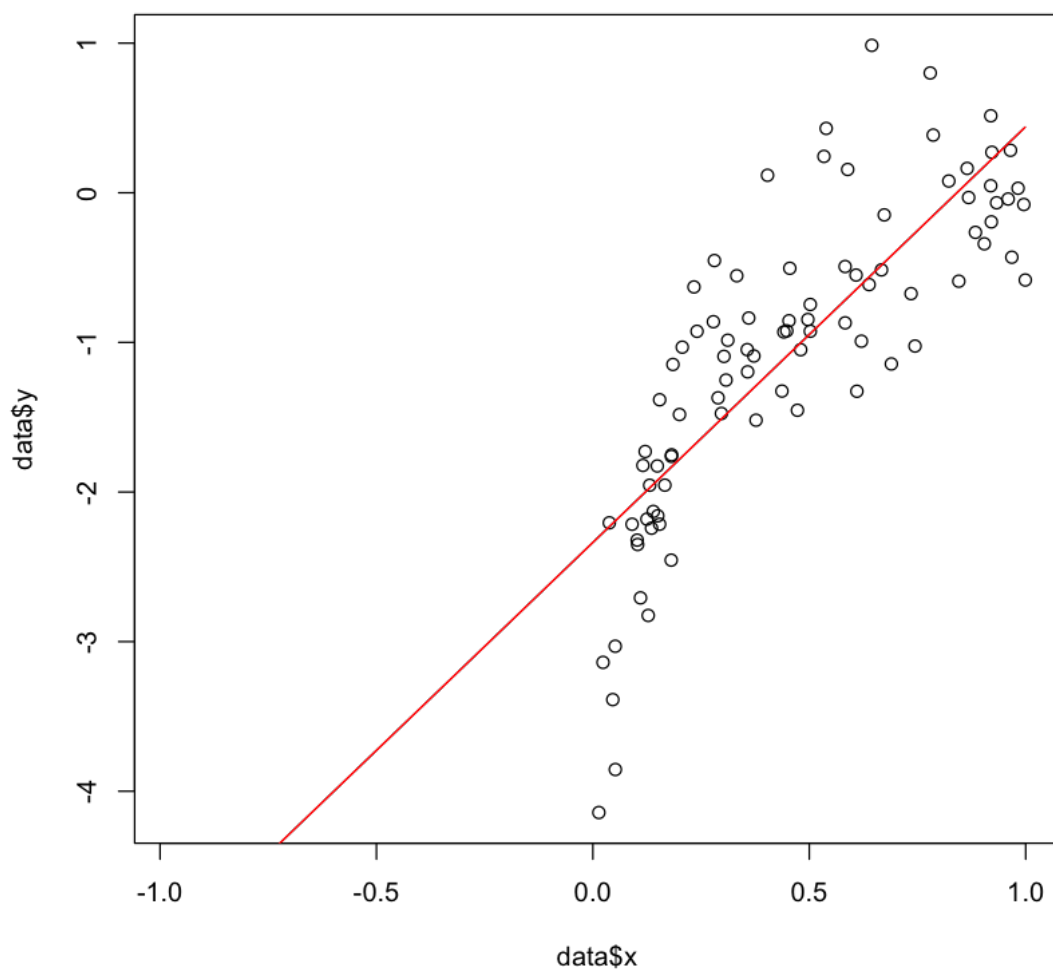
0.2 when these assumptions fail

When our collected data fails to fit all the assumptions of linear regression we have two options: (i) pick a different, more complicated model or (ii) transform our original data so that our transformed data meet the above linear regression assumptions. We'll spend quite a bit of time on (i). For now, lets spend time on (ii).

0.2.1 Exploratory Data Analysis

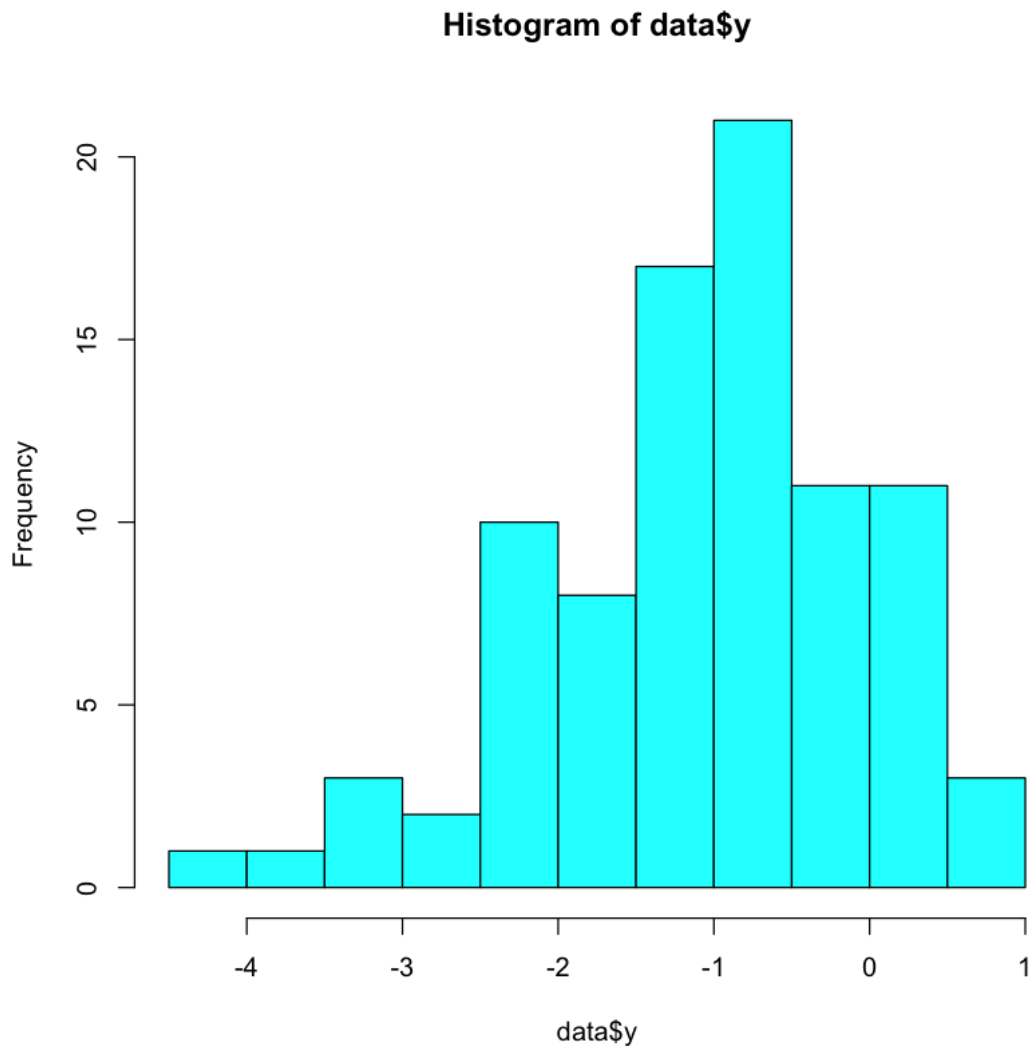
X and Y data in hand, we can plot the relationship between these variables and a linear regression fit.

```
[69]: data <- read.csv("./dataSet1.csv")
      plot(data$x,data$y)
      model1 <- lm(y~x,data=data)
      lines(data$x,predict.lm(model1,data.frame(x=data$x)),col='red')
```



The linear regression fits well for x values great than $1/2$ but it looks like the relationship between x and y is non-linear. Before we begin transforming our variables, let look at a histogram of y .

```
[70]: hist(data$y,10,col='cyan')
```



0.3 Checking linearity (transforming X)

Our y values seem to exhibit a roughly Normal shape. Transforming the y covariate may move the distribution of y further away from normal. We can look at different transformation of x first.

```
[60]: par(mfrow=c(2,2))

transformedX = log(data$x)
plot(transformedX,data$y)
model <- lm(y~log(x),data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
```

```

lines(transformedX
      ,predictions
      ,col='red')

transformedX = (data$x)^0.5
plot(data$x^0.5,data$y)
model <- lm(y~I(x^0.5),data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
lines(transformedX
      ,predictions
      ,col='red')

transformedX = (data$x)^0.333
plot(data$x^0.333,data$y)
model <- lm(y~I(x^0.333),data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
lines(transformedX
      ,predictions
      ,col='red')

transformedX = (data$x)^0.25
plot(data$x^0.25,data$y)
model <- lm(y~I(x^0.25),data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
lines(transformedX
      ,predictions
      ,col='red')

```

Warning message in log(data\$x):

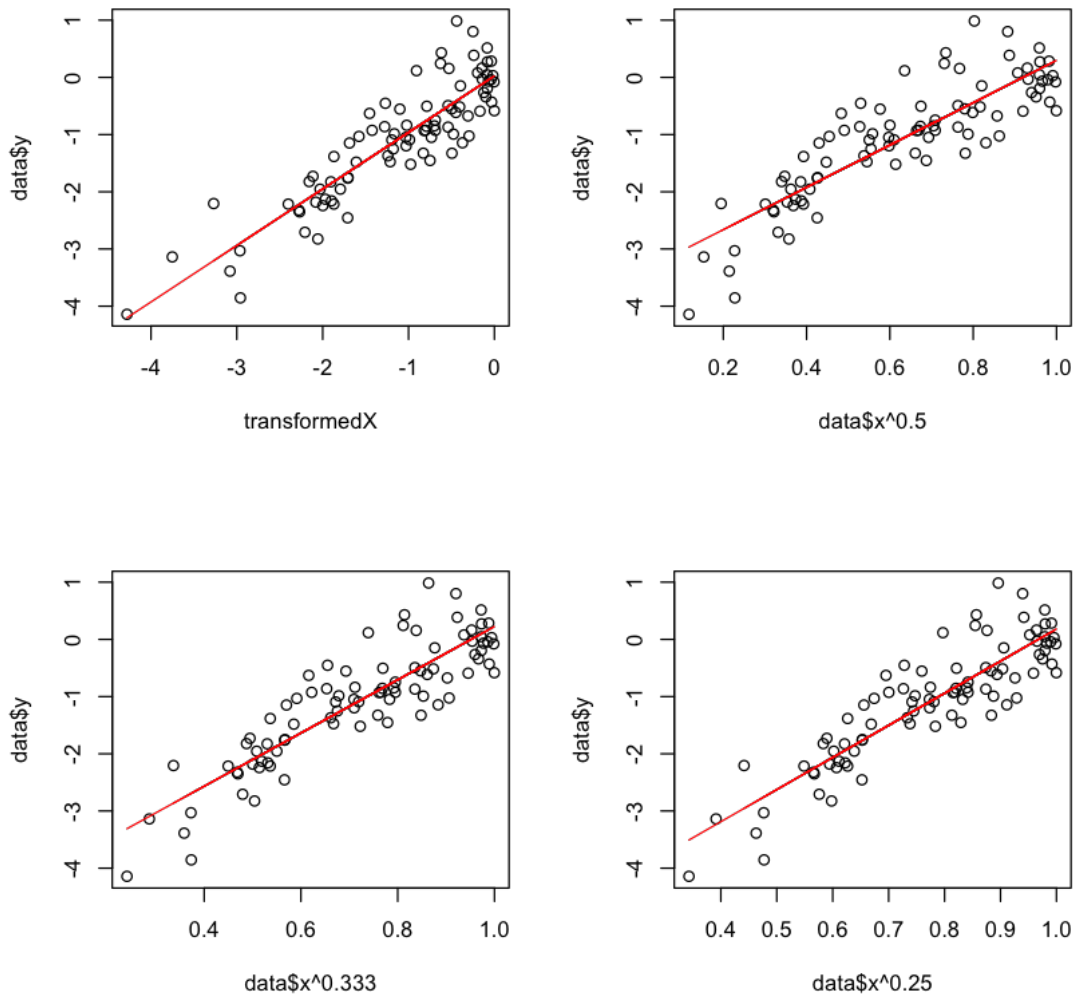
â€œNaNs producedâ€œ

Warning message in log(x):

â€œNaNs producedâ€œ

Warning message in log(x):

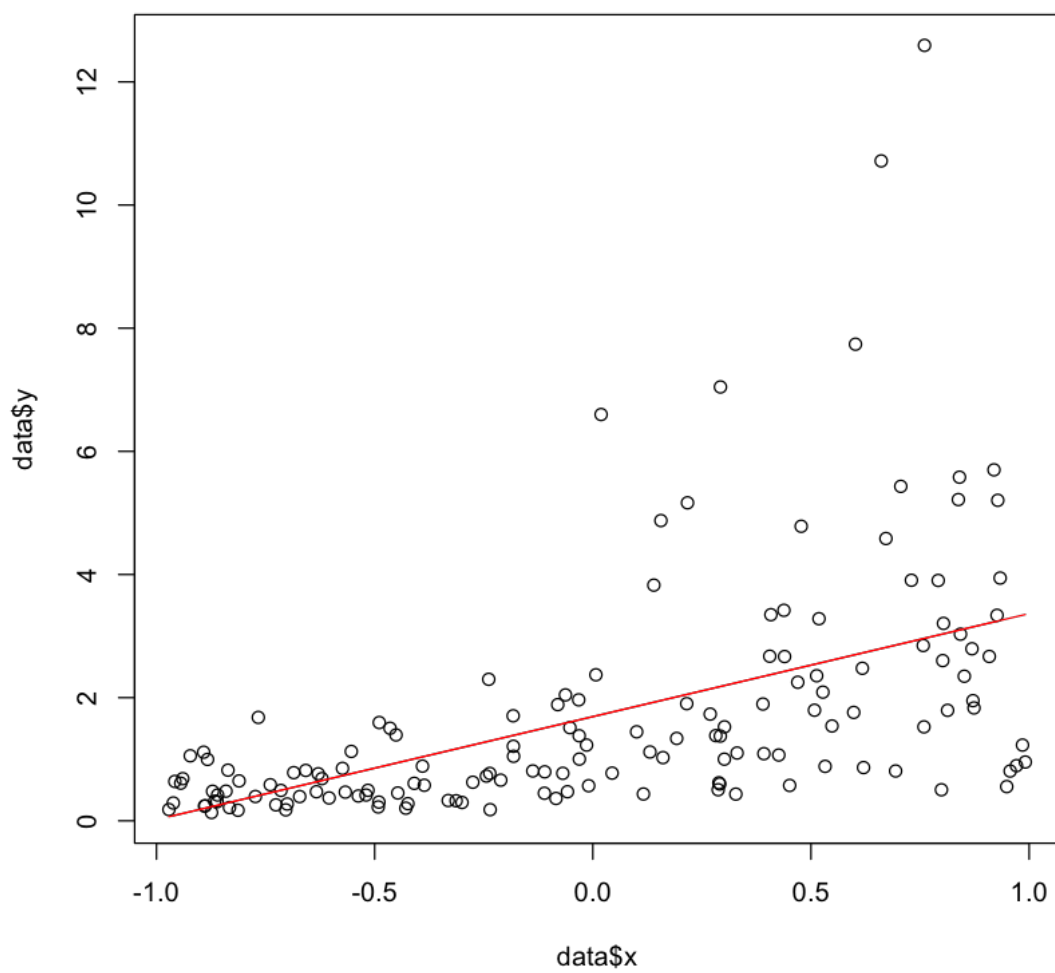
â€œNaNs producedâ€œ



Common transformations to try applying to x are: log transform, exp transform, and x raised to some power.

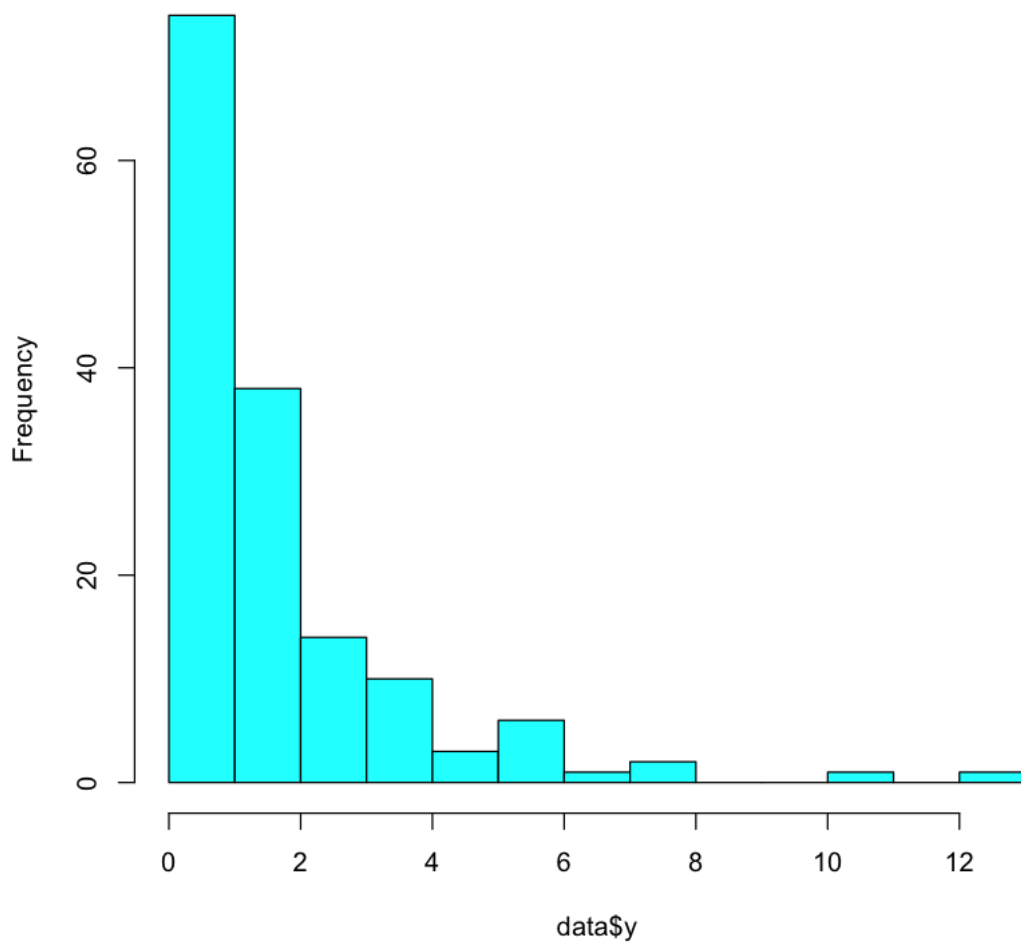
0.4 Checking Normality (transforming y)

```
[71]: data <- read.csv("../dataSet2.csv")
      plot(data$x,data$y)
      model1 <- lm(y~x,data=data)
      lines(data$x,predict.lm(model1,data.frame(x=data$x)),col='red')
```



```
[72]: hist(data$y,10,col='cyan')
```

Histogram of data\$y



```
[74]: par(mfrow=c(3,2))

transformedy = log(data$y)
plot(data$x,transformedy)
model <- lm(transformedy~x,data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
lines(data$x
      ,predictions
      ,col='red')

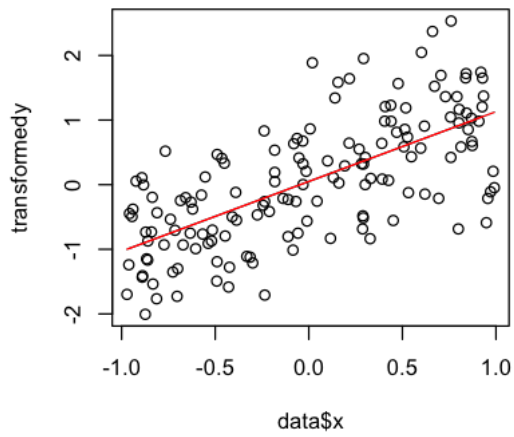
hist()
```

```
transformedy = log(data$y)
plot(data$x,transformedy)
model <- lm(transformedy~x,data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
lines(data$x
      ,predictions
      ,col='red')

transformedy = log(data$y)
plot(data$x,transformedy)
model <- lm(transformedy~x,data=data)

predictions <- predict.lm(model,data.frame(x=data$x))
lines(data$x
      ,predictions
      ,col='red')
```

0.5 The QQ-plot

[]: