

Multiple linear regression

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Grading the professor

Many college courses conclude by giving students the opportunity to evaluate the course and the instructor anonymously. However, the use of these student evaluations as an indicator of course quality and teaching effectiveness is often criticized because these measures may reflect the influence of non-teaching related characteristics, such as the physical appearance of the instructor. The article titled, “Beauty in the classroom: instructors’ pulchritude and putative pedagogical productivity” (Hamermesh and Parker, 2005) found that instructors who are viewed to be better looking receive higher instructional ratings. (Daniel S. Hamermesh, Amy Parker, Beauty in the classroom: instructors pulchritude and putative pedagogical productivity, *Economics of Education Review*, Volume 24, Issue 4, August 2005, Pages 369-376, ISSN 0272-7757, 10.1016/j.econedurev.2004.07.013. <http://www.sciencedirect.com/science/article/pii/S0272775704001165>.)

In this lab we will analyze the data from this study in order to learn what goes into a positive professor evaluation.

The data

The data were gathered from end of semester student evaluations for a large sample of professors from the University of Texas at Austin. In addition, six students rated the professors’ physical appearance. (This is a slightly modified version of the original data set that was released as part of the replication data for *Data Analysis Using Regression and Multilevel/Hierarchical Models* (Gelman and Hill, 2007).) The result is a data frame where each row contains a different course and columns represent variables about the courses and professors.

```
load("more/evals.RData")
```

variable	description
score	average professor evaluation score: (1) very unsatisfactory - (5) excellent.
rank	rank of professor: teaching, tenure track, tenured.
ethnicity	ethnicity of professor: not minority, minority.
gender	gender of professor: female, male.

variable	description
language	language of school where professor received education: english or non-english.
age	age of professor.
cls_perc_eval	percent of students in class who completed evaluation.
cls_did_eval	number of students in class who completed evaluation.
cls_students	total number of students in class.
cls_level	class level: lower, upper.
cls_profs	number of professors teaching sections in course in sample: single, multiple.
cls_credits	number of credits of class: one credit (lab, PE, etc.), multi credit.
bty_f1lower	beauty rating of professor from lower level female: (1) lowest - (10) highest.
bty_f1upper	beauty rating of professor from upper level female: (1) lowest - (10) highest.
bty_f2upper	beauty rating of professor from second upper level female: (1) lowest - (10) highest.

variable	description
bty_m1lower	beauty rating of professor from lower level male: (1) lowest - (10) highest.
bty_m1upper	beauty rating of professor from upper level male: (1) lowest - (10) highest.
bty_m2upper	beauty rating of professor from second upper level male: (1) lowest - (10) highest.
bty_avg	average beauty rating of professor.
pic_outfit	outfit of professor in picture: not formal, formal.
pic_color	color of professor's picture: color, black & white.

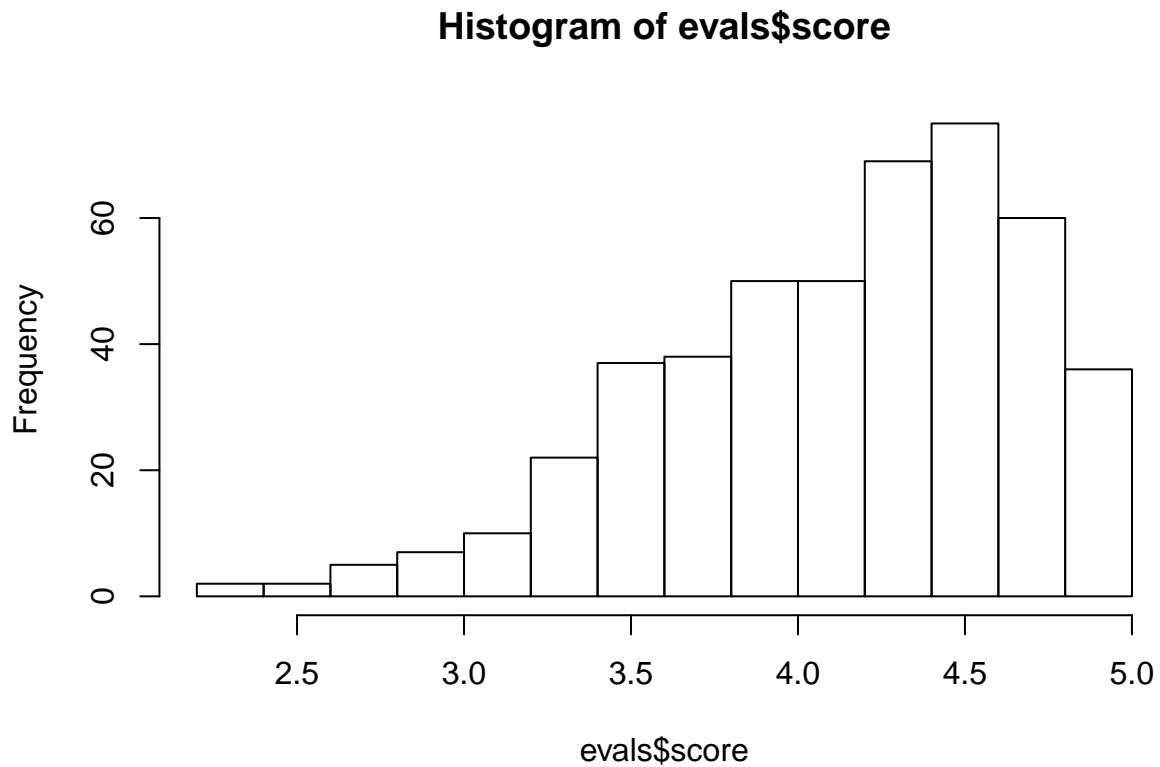
Exploring the data

1. Is this an observational study or an experiment? The original research question posed in the paper is whether beauty leads directly to the differences in course evaluations. Given the study design, is it possible to answer this question as it is phrased? If not, rephrase the question.

Answer : This is an observational study since there are no control and experimental groups. Since, this is only an observational study there cannot be causation between the explanatory and response variables. Instead there can only be a correlation. What we can say is the instructor's beauty has a positive (or negative) correlation to student course evaluation.

2. Describe the distribution of **score**. Is the distribution skewed? What does that tell you about how students rate courses? Is this what you expected to see? Why, or why not?

```
hist(evals$score)
```

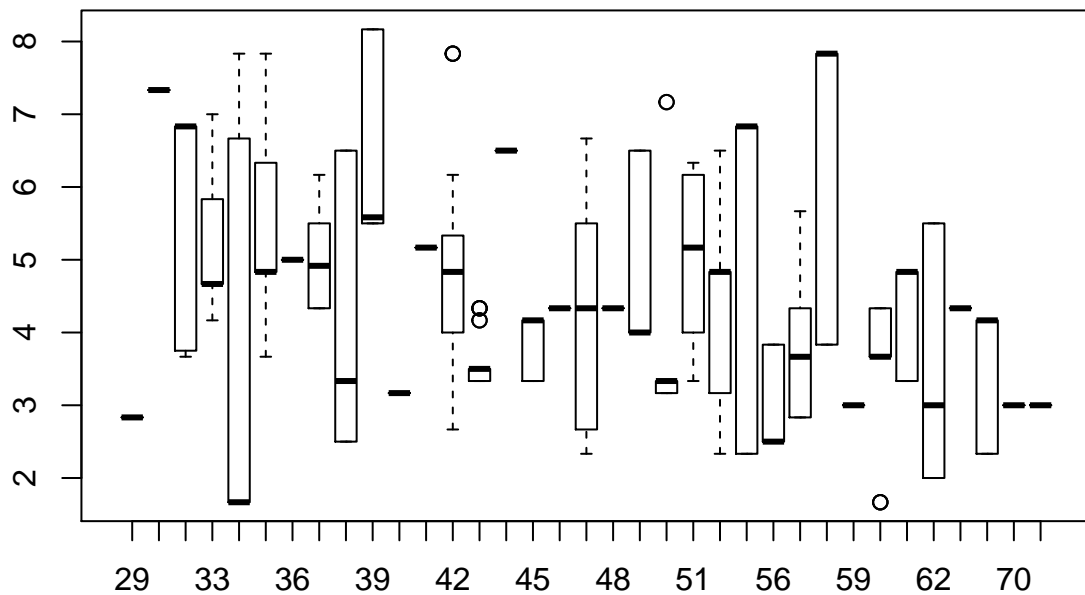


Answer : The evaluation scores are skewed to the left. Students have far more positive evaluations than negative evaluations for their teachers. This is not what I expected. We expected a normal distribution where most teachers would be rated as average and fewer teachers will be evaluated in the extremes - excellent or unsatisfactory.

Excluding score, select two other variables and describe the

3. Excluding `score`, select two other variables and describe their relationship using an appropriate visualization (scatterplot, side-by-side boxplots, or mosaic plot).

```
boxplot(evals$bty_avg ~ evals$age)
```

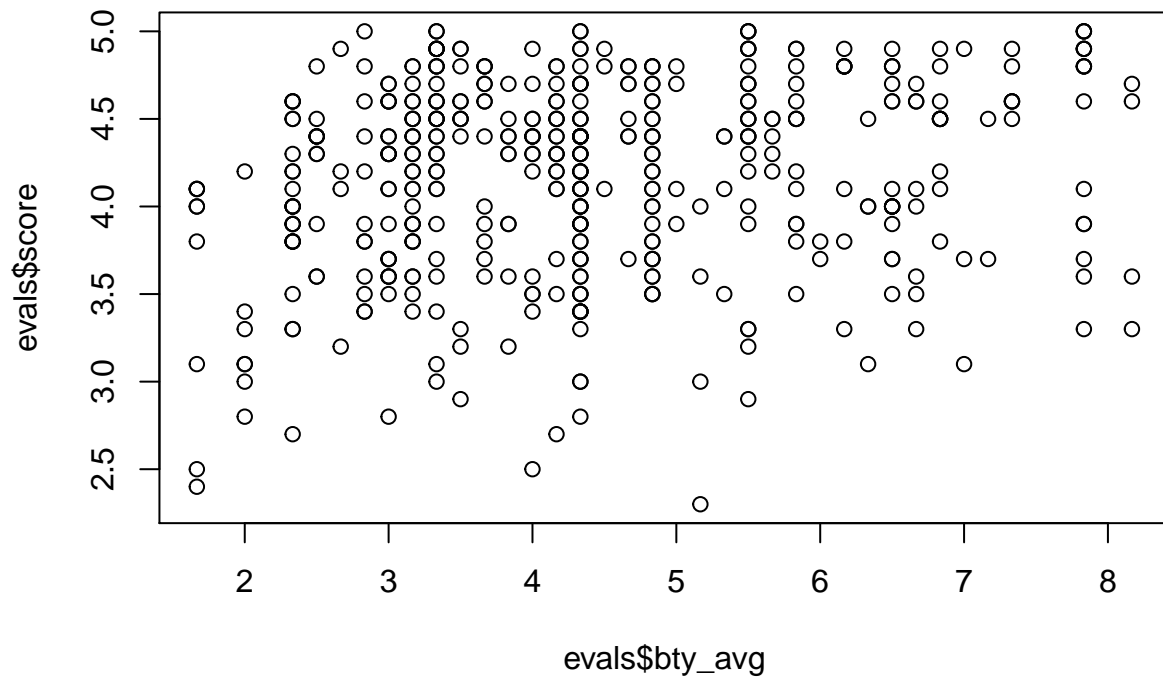


Answer : There doesn't seem to be a relationship between the teacher's age and beauty score. I would have expected an inverse relationship (younger teacher's would have higher beauty scores), but base on the median beauty scores per age group, this is not the case.

Simple linear regression

The fundamental phenomenon suggested by the study is that better looking teachers are evaluated more favorably. Let's create a scatterplot to see if this appears to be the case:

```
plot(evals$score ~ evals$bty_avg)
```



Before we draw conclusions about the trend, compare the number of observations in the data frame with the approximate number of points on the scatterplot. Is anything awry?

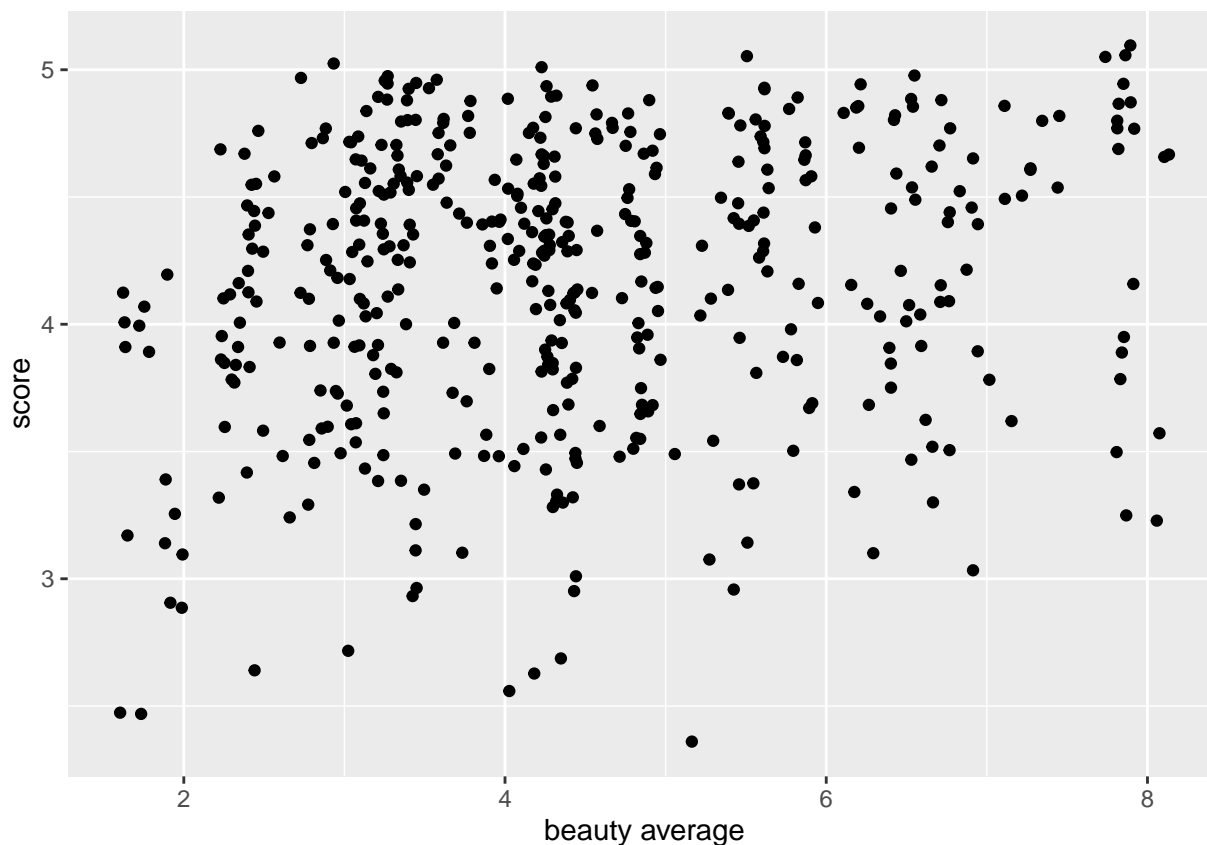
```
nrow(evals)
```

```
## [1] 463
```

Answer : There seem to be more observations than the approximate number of points on the scatterplot.

4. Replot the scatterplot, but this time use the function `jitter()` on the y - or the x -coordinate. (Use `?jitter` to learn more.) What was misleading about the initial scatterplot?

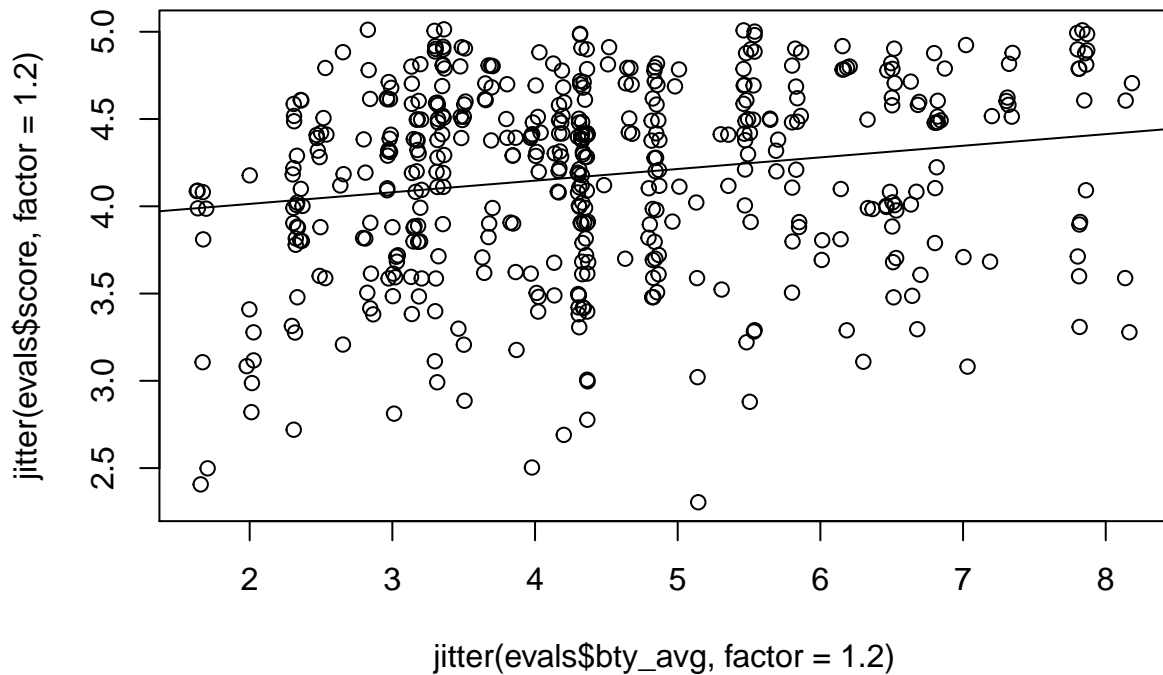
```
library(ggplot2)
ggplot(evals, aes(bty_avg, score)) + geom_point(position = position_jitter(w = 0.3, h = 0.3)) + ylab("score")
```



Answer : It wasn't able to show the relationship between beauty average and score for the teacher because of multiple ties (overlapping scores) that is just represented by a single circle on the scatterplot.

5. Let's see if the apparent trend in the plot is something more than natural variation. Fit a linear model called `m_bty` to predict average professor score by average beauty rating and add the line to your plot using `abline(m_bty)`. Write out the equation for the linear model and interpret the slope. Is average beauty score a statistically significant predictor? Does it appear to be a practically significant predictor?

```
m_bty <- lm(evals$score ~ evals$bty_avg)
plot(jitter(evals$score, factor=1.2) ~ jitter(evals$bty_avg, factor=1.2))
abline(m_bty)
```



```
cor(evalscore, evalsbty_avg)
```

```
## [1] 0.1871424
```

```
summary(m_bty)
```

```
##
## Call:
## lm(formula = evalscore ~ evalsbty_avg)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.9246 -0.3690  0.1420  0.3977  0.9309
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   3.88034    0.07614   50.96 < 2e-16 ***
## evalsbty_avg  0.06664    0.01629    4.09 5.08e-05 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.5348 on 461 degrees of freedom
## Multiple R-squared:  0.03502,    Adjusted R-squared:  0.03293
## F-statistic: 16.73 on 1 and 461 DF,  p-value: 5.083e-05
```


Answer :

Yes, `bty_avg` is a statistically significant predictor of evaluation score with p-value close of 0. It may not be a practically significant predictor of evaluation score though since for every 1 point increase in `bty_ave`, the model only predicts an increase of 0.06664 which barely changes the evaluation score.

6. Use residual plots to evaluate whether the conditions of least squares regression are reasonable. Provide plots and comments for each one (see the Simple Regression Lab for a reminder of how to make these).

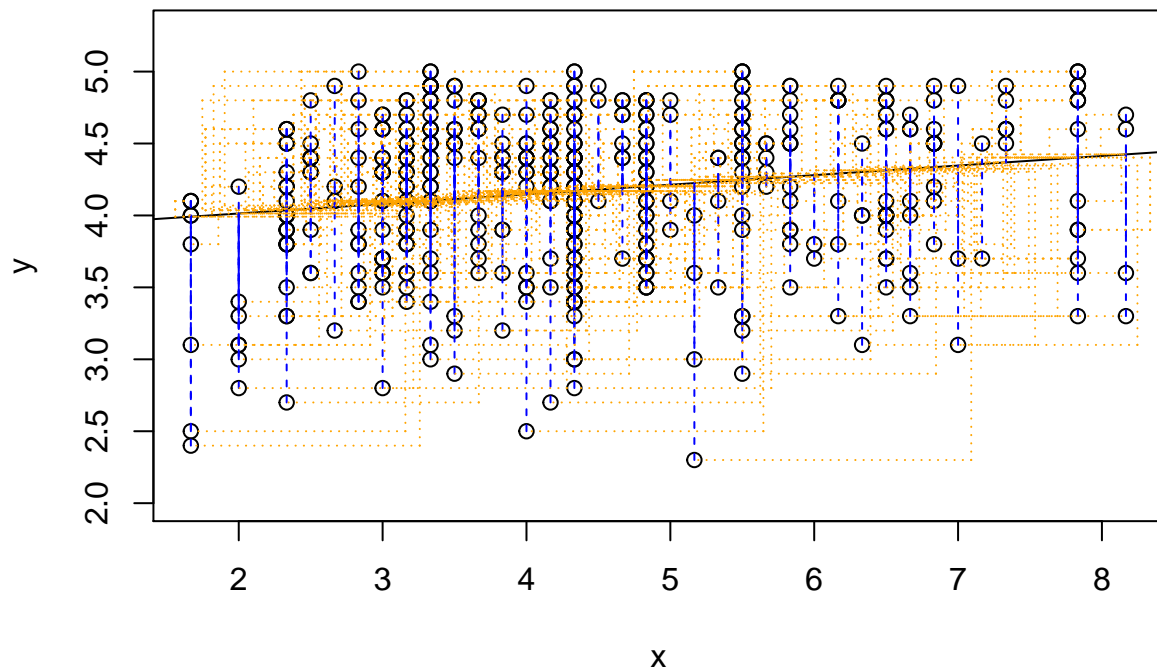
```
plot_ss <- function(x, y, showSquares = FALSE, leastSquares = FALSE){
  plot(y~x, asp = 1) # xlab = paste(substitute(x)), ylab = paste(substitute(y)))

  if(leastSquares){
    m1 <- lm(y~x)
    y.hat <- m1$fit
  } else{
    cat("Click two points to make a line.")
    pt1 <- locator(1)
    points(pt1$x, pt1$y, pch = 4)
    pt2 <- locator(1)
    points(pt2$x, pt2$y, pch = 4)
    pts <- data.frame("x" = c(pt1$x, pt2$x), "y" = c(pt1$y, pt2$y))
    m1 <- lm(y ~ x, data = pts)
    y.hat <- predict(m1, newdata = data.frame(x))
  }
  r <- y - y.hat
  abline(m1)

  oSide <- x - r
  LLim <- par()$usr[1]
  RLim <- par()$usr[2]
  oSide[oSide < LLim | oSide > RLim] <- c(x + r)[oSide < LLim | oSide > RLim] # move boxes to avoid mar

  n <- length(y.hat)
  for(i in 1:n){
    lines(rep(x[i], 2), c(y[i], y.hat[i]), lty = 2, col = "blue")
    if(showSquares){
      lines(rep(oSide[i], 2), c(y[i], y.hat[i]), lty = 3, col = "orange")
      lines(c(oSide[i], x[i]), rep(y.hat[i], 2), lty = 3, col = "orange")
      lines(c(oSide[i], x[i]), rep(y[i], 2), lty = 3, col = "orange")
    }
  }

  SS <- round(sum(r^2), 3)
  cat("\r                                     ")
  print(m1)
  cat("Sum of Squares: ", SS)
}
plot_ss(x = evals$bty_avg, y = evals$score, showSquares = TRUE)
```



```
## Click two points to make a line.
```

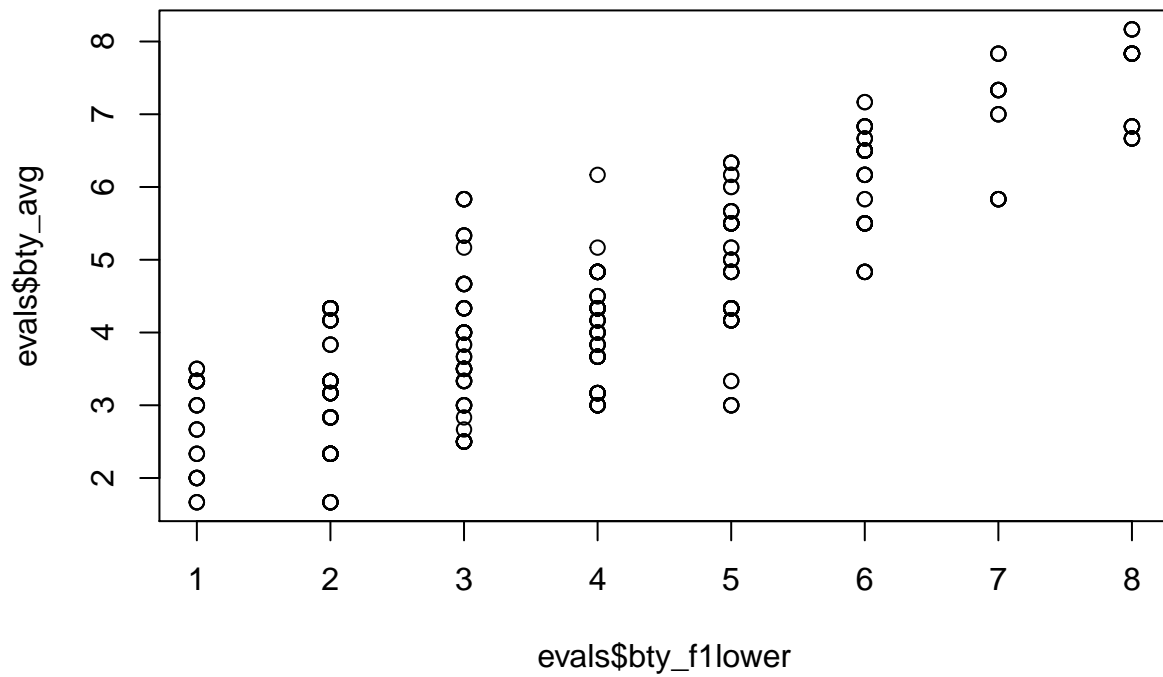
```
## Call:
## lm(formula = y ~ x, data = pts)
##
## Coefficients:
## (Intercept)          x
##      3.88034      0.06664
##
## Sum of Squares:  131.868
```

There are too many outliers and the distribution is not normal.

Multiple linear regression

The data set contains several variables on the beauty score of the professor: individual ratings from each of the six students who were asked to score the physical appearance of the professors and the average of these six scores. Let's take a look at the relationship between one of these scores and the average beauty score.

```
plot(evals$bty_avg ~ evals$bty_follower)
```

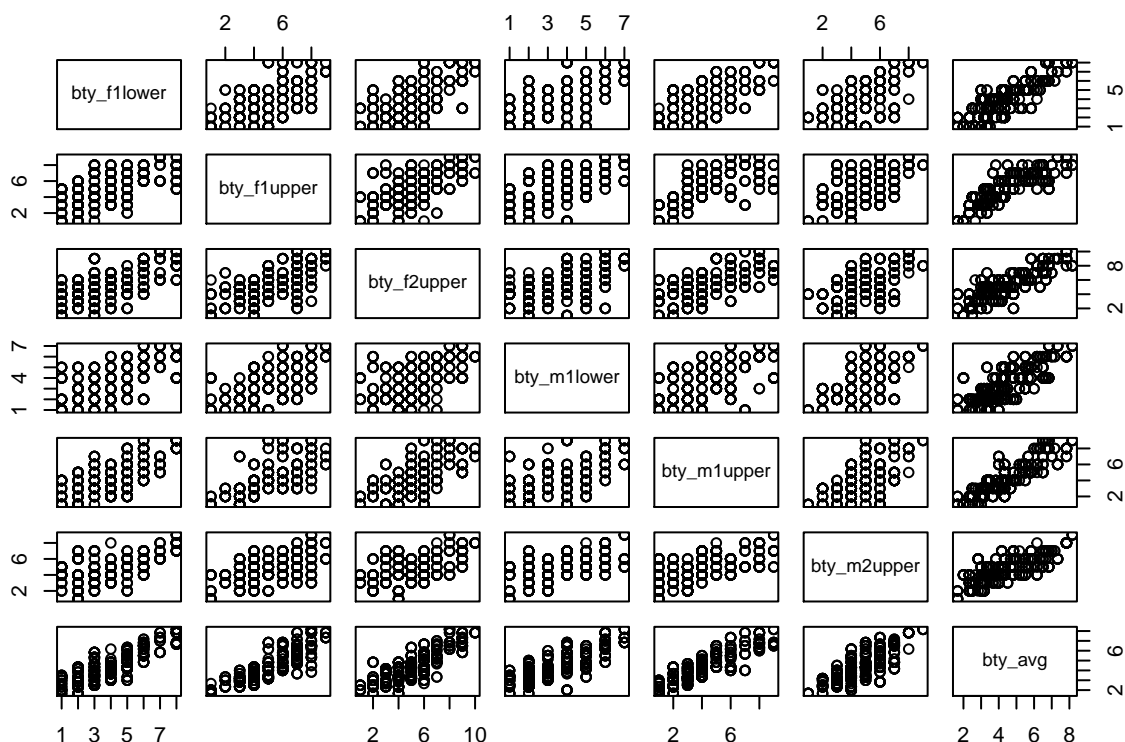


```
cor(evals$bty_avg, evals$bty_f1lower)
```

```
## [1] 0.8439112
```

As expected the relationship is quite strong - after all, the average score is calculated using the individual scores. We can actually take a look at the relationships between all beauty variables (columns 13 through 19) using the following command:

```
plot(evals[,13:19])
```



These variables are collinear (correlated), and adding more than one of these variables to the model would not add much value to the model. In this application and with these highly-correlated predictors, it is reasonable to use the average beauty score as the single representative of these variables.

In order to see if beauty is still a significant predictor of professor score after we've accounted for the gender of the professor, we can add the gender term into the model.

```
m_bty_gen <- lm(score ~ bty_avg + gender, data = evals)
summary(m_bty_gen)
```

```
##
## Call:
## lm(formula = score ~ bty_avg + gender, data = evals)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.8305 -0.3625  0.1055  0.4213  0.9314
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)   3.74734    0.08466  44.266 < 2e-16 ***
## bty_avg        0.07416    0.01625   4.563 6.48e-06 ***
## gendermale    0.17239    0.05022   3.433 0.000652 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
```

```
## Residual standard error: 0.5287 on 460 degrees of freedom
## Multiple R-squared:  0.05912,    Adjusted R-squared:  0.05503
## F-statistic: 14.45 on 2 and 460 DF,  p-value: 8.177e-07
```

```
names(m_bty_gen)
```

```
## [1] "coefficients" "residuals"      "effects"      "rank"
## [5] "fitted.values" "assign"         "qr"          "df.residual"
## [9] "contrasts"     "xlevels"        "call"         "terms"
## [13] "model"
```

7. P-values and parameter estimates should only be trusted if the conditions for the regression are reasonable. Verify that the conditions for this model are reasonable using diagnostic plots. **Answer :**
8. the residuals of the model are nearly normal

```
library(StMoSim)
```

```
## Warning: package 'StMoSim' was built under R version 3.3.2

## Loading required package: RcppParallel

## Warning: package 'RcppParallel' was built under R version 3.3.2

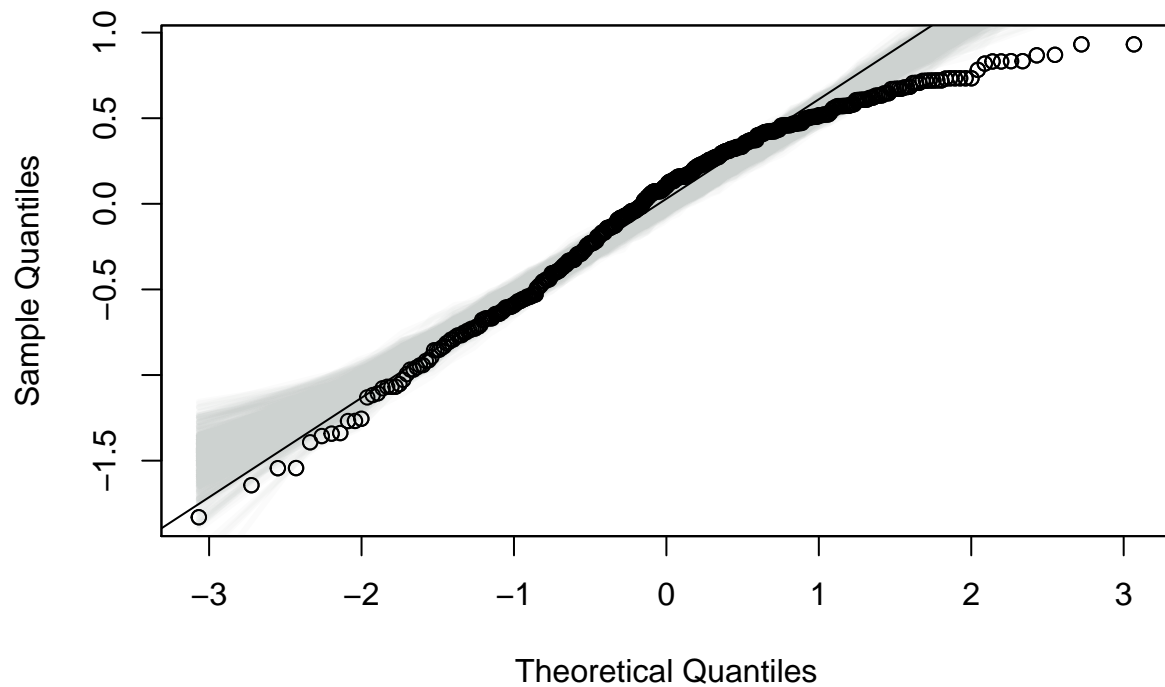
## Loading required package: Rcpp

##
## Attaching package: 'Rcpp'

## The following object is masked from 'package:RcppParallel':
##
##      LdFlags
```

```
qqnormSim(m_bty_gen$residuals)
```

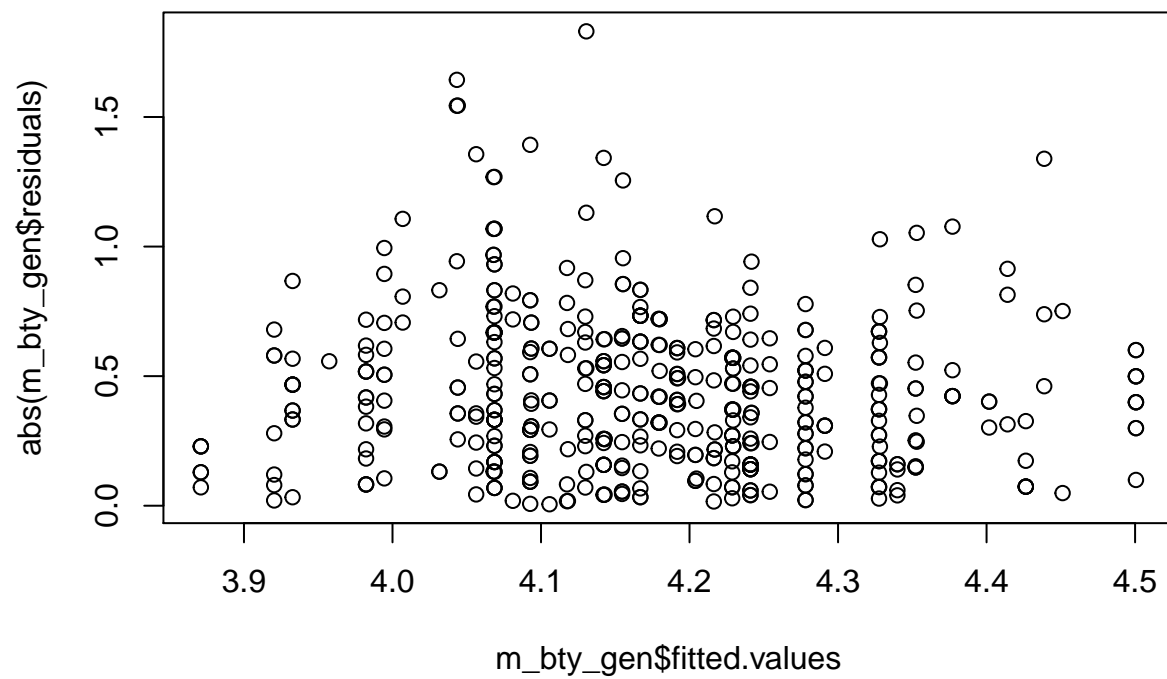
Normal Q-Q Plot – SIM



The residuals of the model is not normal as residual values for the the higher quantiles are less than what a normal distribution would predict

2. Absolute values of residuals against fitted values. (the variability of the residuals is nearly constant)

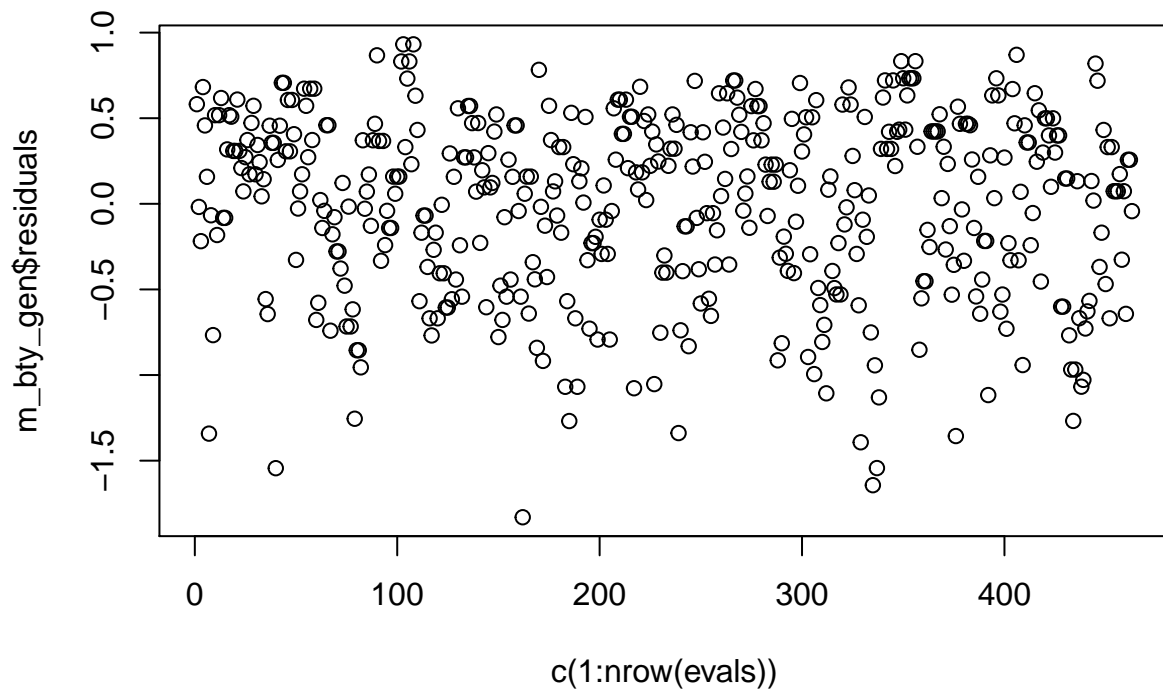
```
plot(abs(m_bty_gen$residuals) ~ m_bty_gen$fitted.values)
```



There some outliers although overall, most of the residual values are close to the fitted values.

3. The residuals are independent

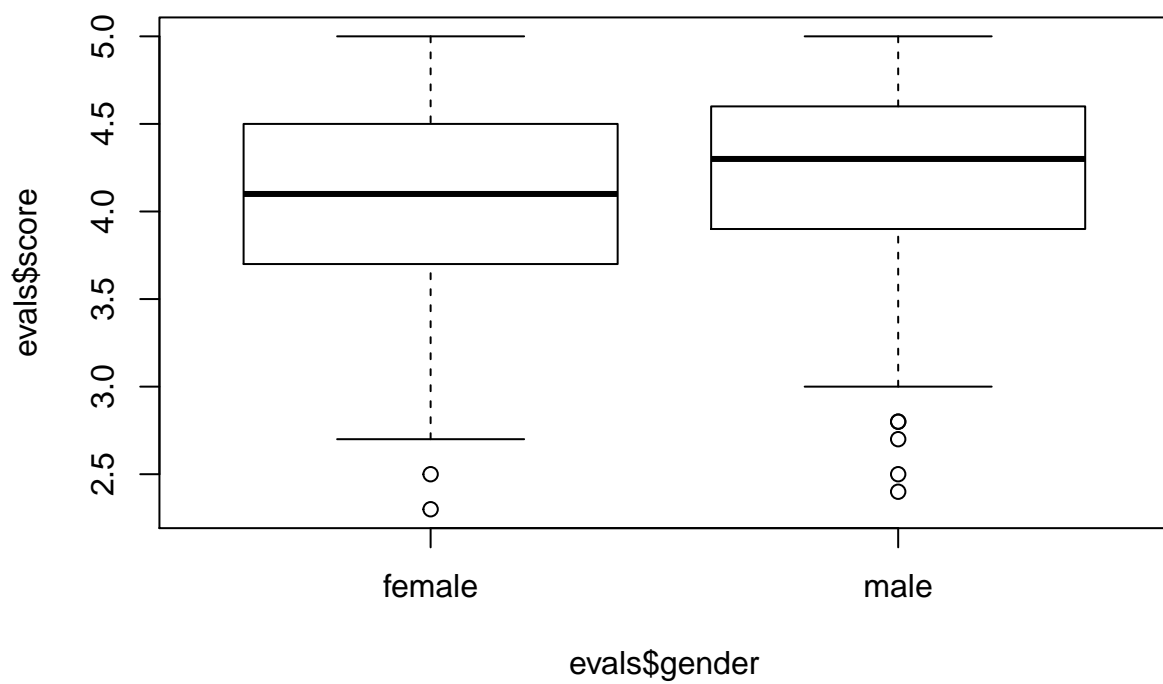
```
plot(m_bty_gen$residuals ~ c(1:nrow(evals)))
```



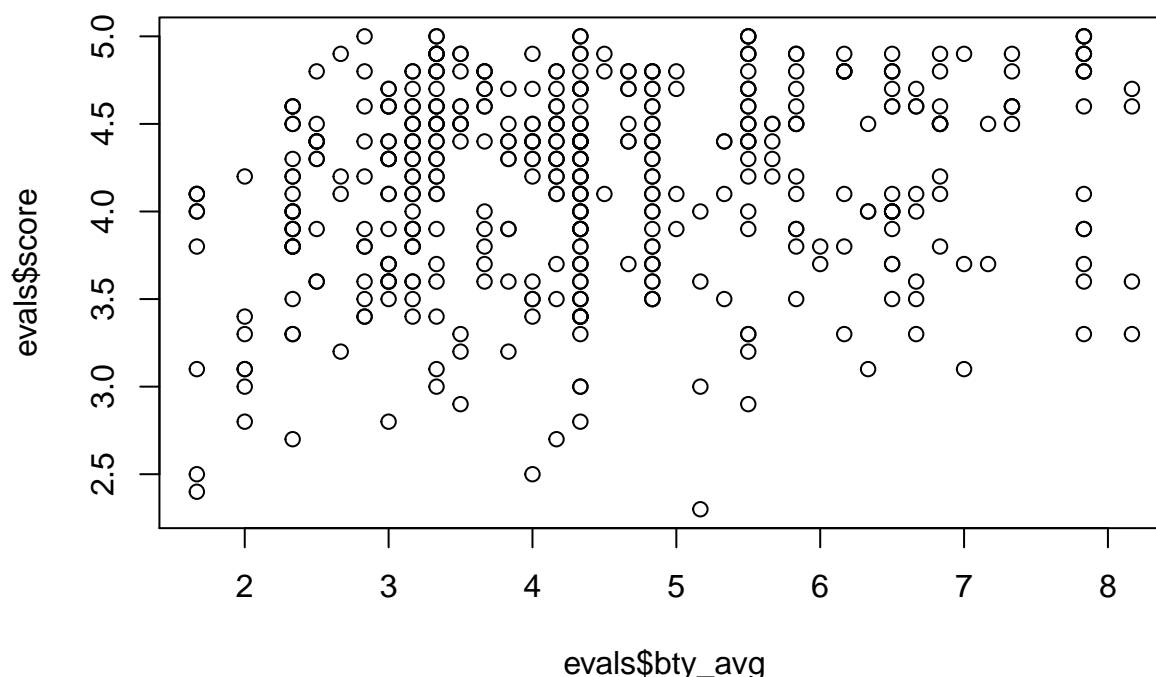
Yes, this condition is met. The residuals based on the sequence when it was gathered shows that they were randomly gathered.

4. each variable is linearly related to the outcome.

```
plot(evals$score ~ evals$gender)
```

```
plot(evals$score ~ evals$gender)
```



There is a linear relationship between gender and evaluation score. The median scores and variability for both males and females are similar in terms of evaluation scores. As was established in the previous exercises, there is a linear relationship between beauty average and teaching evaluation score.

8. Is `bty_avg` still a significant predictor of `score`? Has the addition of `gender` to the model changed the parameter estimate for `bty_avg`? **Answer :** Yes it is, gender made beauty average even more significant as the p-value computed is even smaller now compared to a model where beauty average was the sole variable.

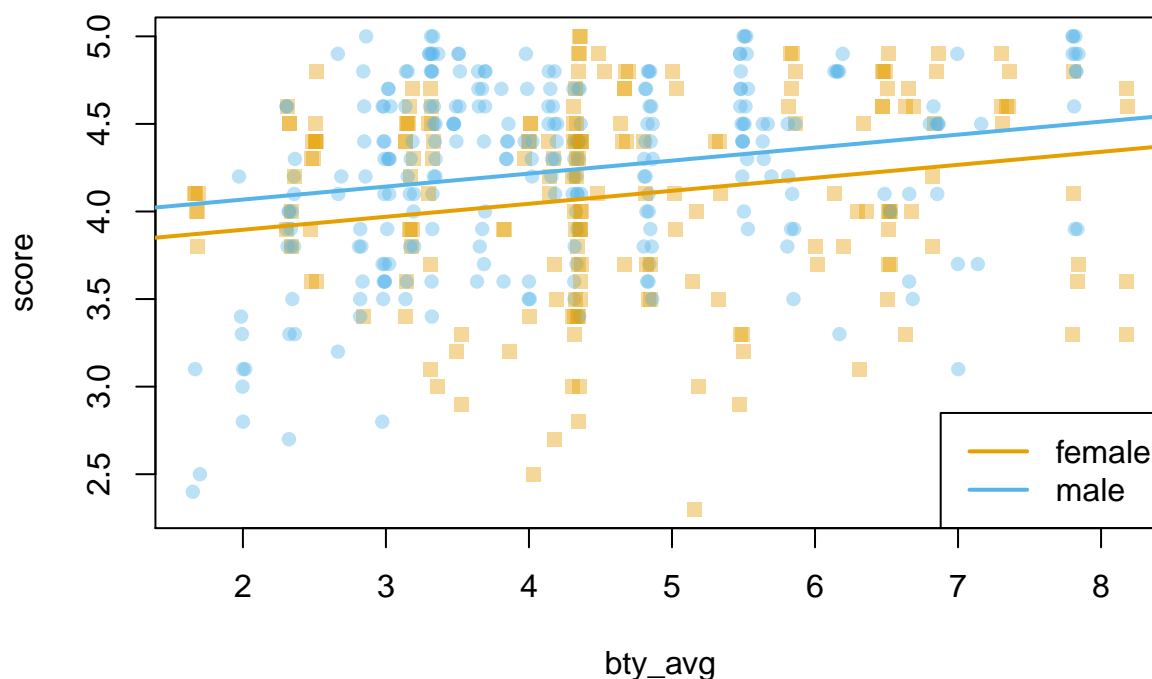
Note that the estimate for `gender` is now called `gendermale`. You'll see this name change whenever you introduce a categorical variable. The reason is that R recodes `gender` from having the values of `female` and `male` to being an indicator variable called `gendermale` that takes a value of 0 for females and a value of 1 for males. (Such variables are often referred to as “dummy” variables.)

As a result, for females, the parameter estimate is multiplied by zero, leaving the intercept and slope form familiar from simple regression.

$$\begin{aligned}\widehat{score} &= \hat{\beta}_0 + \hat{\beta}_1 \times bty_avg + \hat{\beta}_2 \times (0) \\ &= \hat{\beta}_0 + \hat{\beta}_1 \times bty_avg\end{aligned}$$

We can plot this line and the line corresponding to males with the following custom function.

```
multiLines(m_bty_gen)
```



9. What is the equation of the line corresponding to males? (*Hint:* For males, the parameter estimate is multiplied by 1.) For two professors who received the same beauty rating, which gender tends to have the higher course evaluation score?

$$\begin{aligned}\widehat{score} &= \hat{\beta}_0 + \hat{\beta}_1 \times bty_avg + \hat{\beta}_2 \times (1) \\ &= \hat{\beta}_0 + \hat{\beta}_1 \times bty_avg + \hat{\beta}_2\end{aligned}$$

Answer : Males tend to have a higher course evaluation score than females for professors who get the same rating.

The decision to call the indicator variable `gendermale` instead of `genderfemale` has no deeper meaning. R simply codes the category that comes first alphabetically as a 0. (You can change the reference level of a categorical variable, which is the level that is coded as a 0, using the `relevel` function. Use `?relevel` to learn more.)

10. Create a new model called `m_bty_rank` with `gender` removed and `rank` added in. How does R appear to handle categorical variables that have more than two levels? Note that the rank variable has three levels: `teaching`, `tenure track`, `tenured`.

```
m_bty_rank <- lm(score ~ bty_avg + rank, data = evals)
summary(m_bty_rank)
```

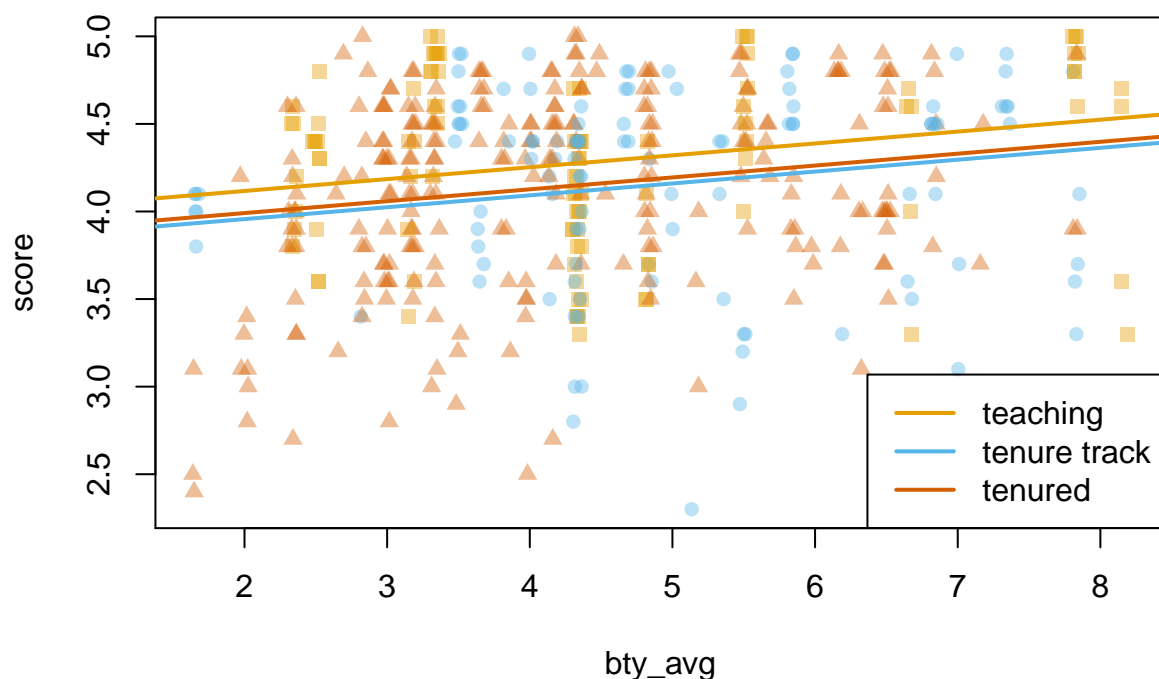
```
##
## Call:
```

```
## lm(formula = score ~ bty_avg + rank, data = evals)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.8713 -0.3642  0.1489  0.4103  0.9525
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    3.98155    0.09078  43.860 < 2e-16 ***
## bty_avg         0.06783    0.01655   4.098 4.92e-05 ***
## ranktenure track -0.16070    0.07395  -2.173  0.0303 *
## ranktenured     -0.12623    0.06266  -2.014  0.0445 *
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.5328 on 459 degrees of freedom
## Multiple R-squared:  0.04652,    Adjusted R-squared:  0.04029
## F-statistic: 7.465 on 3 and 459 DF,  p-value: 6.88e-05
```

```
names(m_bty_rank)
```

```
## [1] "coefficients" "residuals"      "effects"         "rank"
## [5] "fitted.values" "assign"          "qr"              "df.residual"
## [9] "contrasts"     "xlevels"        "call"            "terms"
## [13] "model"
```

```
multiLines(m_bty_rank)
```



The interpretation of the coefficients in multiple regression is slightly different from that of simple regression. The estimate for `bty_avg` reflects how much higher a group of professors is expected to score if they have a beauty rating that is one point higher *while holding all other variables constant*. In this case, that translates into considering only professors of the same rank with `bty_avg` scores that are one point apart.

The search for the best model

We will start with a full model that predicts professor score based on rank, ethnicity, gender, language of the university where they got their degree, age, proportion of students that filled out evaluations, class size, course level, number of professors, number of credits, average beauty rating, outfit, and picture color.

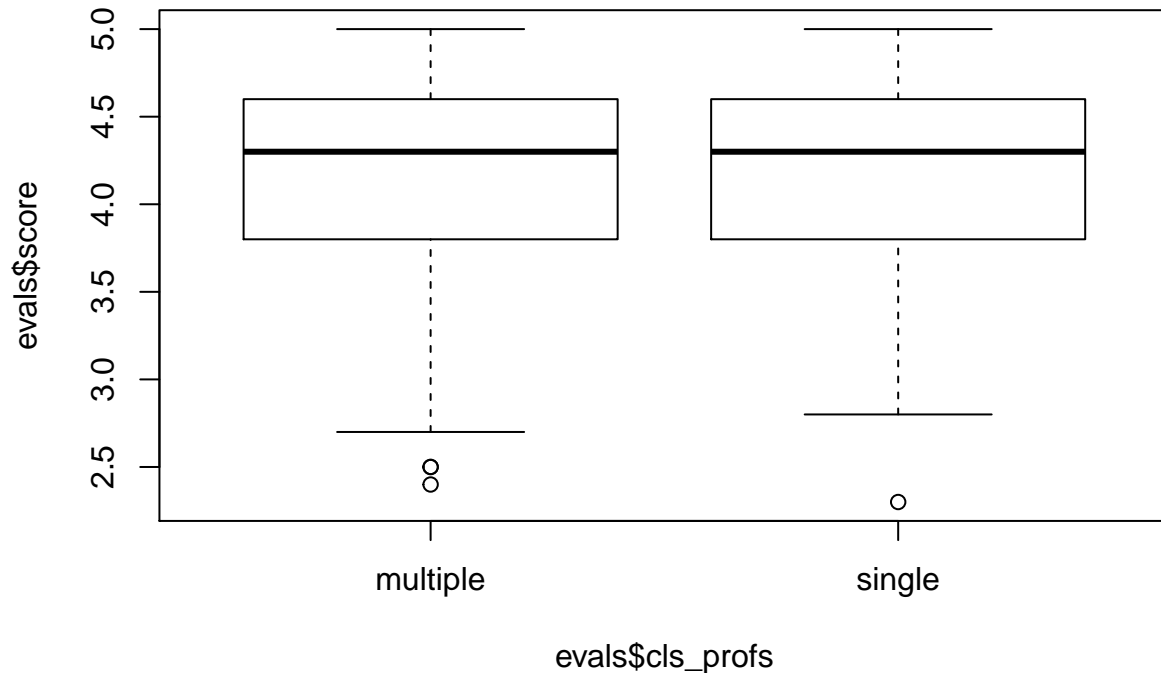
11. Which variable would you expect to have the highest p-value in this model? Why? *Hint:* Think about which variable would you expect to not have any association with the professor score. **Answer** : “number of professors” `cls_profs` as the variable to have the least association with the professor’s evaluation score.

Let’s run the model...

```
m_full <- lm(score ~ rank + ethnicity + gender + language + age + cls_perc_eval
+ cls_students + cls_level + cls_profs + cls_credits + bty_avg
+ pic_outfit + pic_color, data = evals)
summary(m_full)
```

12. Check your suspicions from the previous exercise. Include the model output in your response. **Answer** : Yes it is “number professors” that has the least association to “scores”. It has the highest p-value in the model.

```
plot(evals$score ~ evals$cls_profs)
```



13. Interpret the coefficient associated with the ethnicity variable. **Answer :** The ethnicity p-value of about 0.11 means that it has a weak relationship to scores and may be dropped as part of the model.
14. Drop the variable with the highest p-value and re-fit the model. Did the coefficients and significance of the other explanatory variables change? (One of the things that makes multiple regression interesting is that coefficient estimates depend on the other variables that are included in the model.) If not, what does this say about whether or not the dropped variable was collinear with the other explanatory variables?

```
m_back <- lm(score ~ rank + ethnicity + gender + language + age + cls_perc_eval +
  cls_students + cls_level + cls_credits + bty_avg + pic_outfit + pic_color,
  data = evals)
summary(m_back)
```

```
##
## Call:
## lm(formula = score ~ rank + ethnicity + gender + language + age +
##   cls_perc_eval + cls_students + cls_level + cls_credits +
##   bty_avg + pic_outfit + pic_color, data = evals)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
```

```
## -1.7836 -0.3257 0.0859 0.3513 0.9551
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      4.0872523   0.2888562   14.150 < 2e-16 ***
## ranktenure track  -0.1476746   0.0819824   -1.801 0.072327 .
## ranktenured       -0.0973829   0.0662614   -1.470 0.142349
## ethnicitynot minority 0.1274458   0.0772887    1.649 0.099856 .
## gendermale        0.2101231   0.0516873    4.065 5.66e-05 ***
## languagenon-english -0.2282894   0.1111305   -2.054 0.040530 *
## age              -0.0089992   0.0031326   -2.873 0.004262 **
## cls_perc_eval      0.0052888   0.0015317    3.453 0.000607 ***
## cls_students       0.0004687   0.0003737    1.254 0.210384
## cls_levelupper     0.0606374   0.0575010    1.055 0.292200
## cls_creditsone credit 0.5061196   0.1149163    4.404 1.33e-05 ***
## bty_avg            0.0398629   0.0174780    2.281 0.023032 *
## pic_outfitnot formal -0.1083227   0.0721711   -1.501 0.134080
## pic_colorcolor     -0.2190527   0.0711469   -3.079 0.002205 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4974 on 449 degrees of freedom
## Multiple R-squared:  0.187, Adjusted R-squared:  0.1634
## F-statistic: 7.943 on 13 and 449 DF, p-value: 2.336e-14
```

Yes. There was a slight change in the coefficients and significance of the other explanatory variables when `cls_profs` was removed. All the values are now slightly lower - meaning they are more significant now to the level than before

15. Using backward-selection and p-value as the selection criterion, determine the best model. You do not need to show all steps in your answer, just the output for the final model. Also, write out the linear model for predicting score based on the final model you settle on.

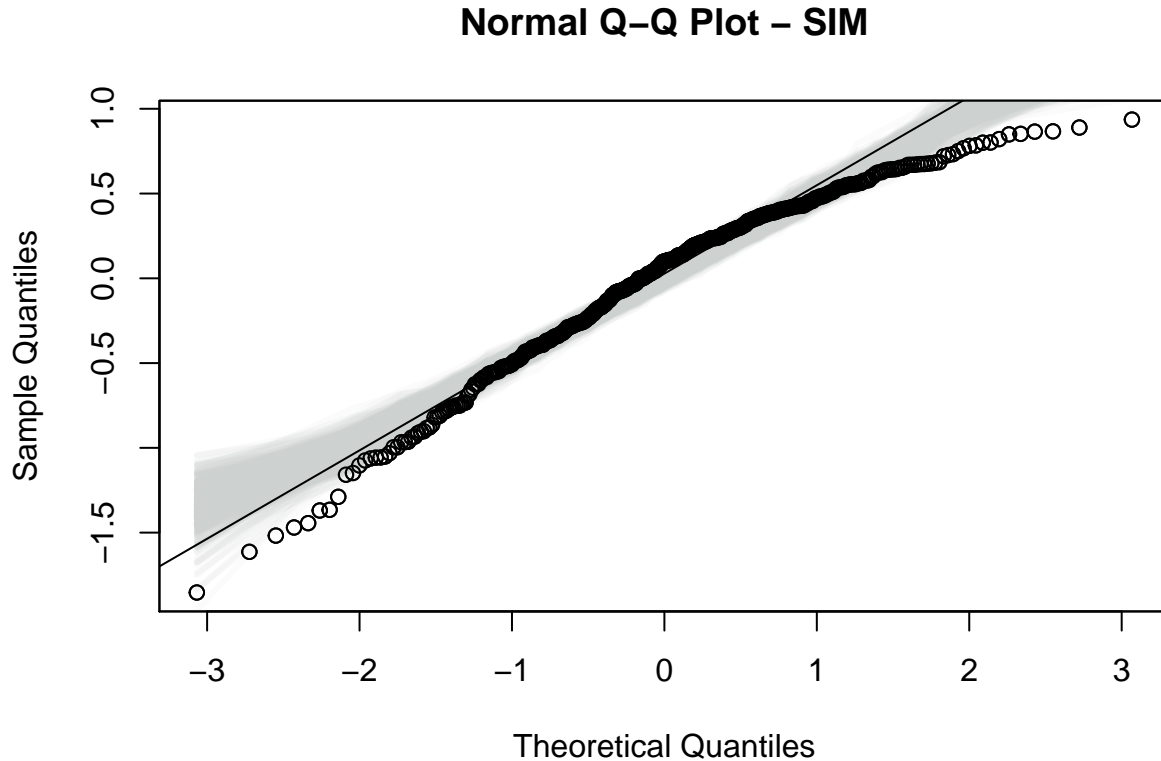
```
m_back2 <- lm(score ~ ethnicity + gender + language + age + cls_perc_eval +
  cls_credits + bty_avg + pic_color, data = evals)
summary(m_back2)

##
## Call:
## lm(formula = score ~ ethnicity + gender + language + age + cls_perc_eval +
##     cls_credits + bty_avg + pic_color, data = evals)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1.85320 -0.32394  0.09984  0.37930  0.93610
##
## Coefficients:
##               Estimate Std. Error t value Pr(>|t|)
## (Intercept)      3.771922   0.232053  16.255 < 2e-16 ***
## ethnicitynot minority 0.167872   0.075275   2.230 0.02623 *
## gendermale        0.207112   0.050135   4.131 4.30e-05 ***
## languagenon-english -0.206178   0.103639   -1.989 0.04726 *
## age              -0.006046   0.002612   -2.315 0.02108 *
```

```
## cls_perc_eval          0.004656    0.001435    3.244  0.00127 **
## cls_creditsone credit  0.505306    0.104119    4.853  1.67e-06 ***
## bty_avg                0.051069    0.016934    3.016  0.00271 **
## pic_colorcolor        -0.190579    0.067351   -2.830  0.00487 **
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.4992 on 454 degrees of freedom
## Multiple R-squared:  0.1722, Adjusted R-squared:  0.1576
## F-statistic: 11.8 on 8 and 454 DF, p-value: 2.58e-15
```

16. Verify that the conditions for this model are reasonable using diagnostic plots. **Answer :**
 17. The residuals of the model are nearly normal

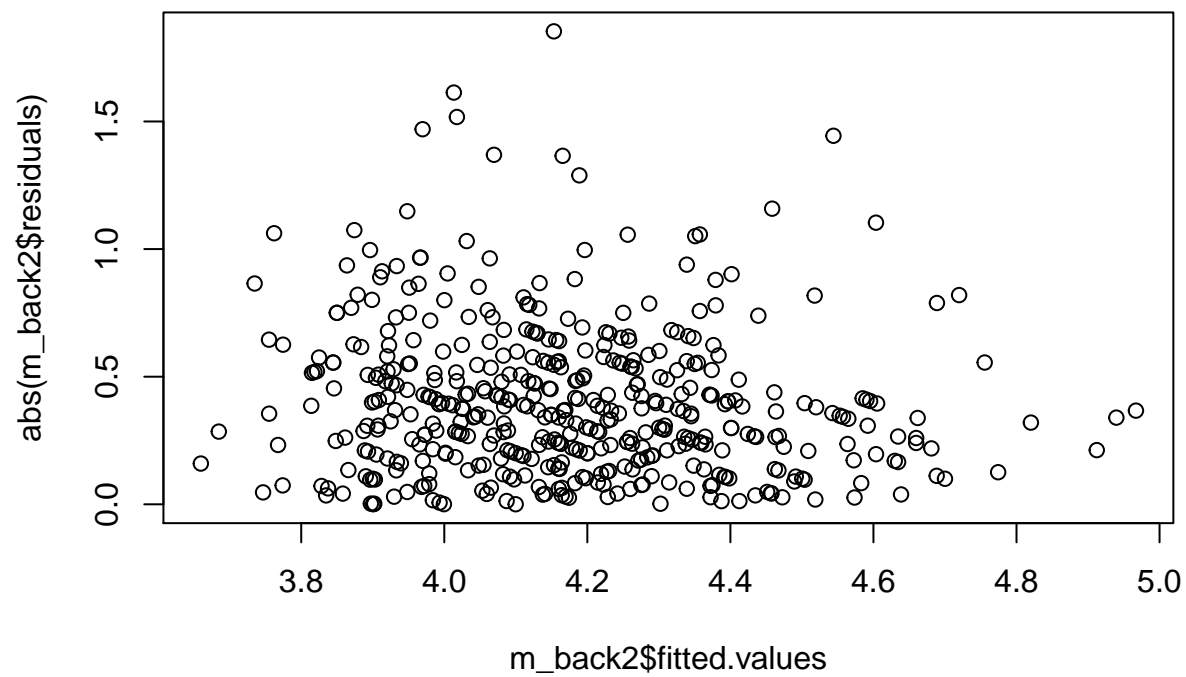
```
qqnormSim(m_back2$residuals)
```



The residuals of the model is not normal as residual values for the the higher and lower quantiles are less than what a normal distribution would predict

2. Absolute values of residuals against fitted values. (the variability of the residuals is nearly constant)

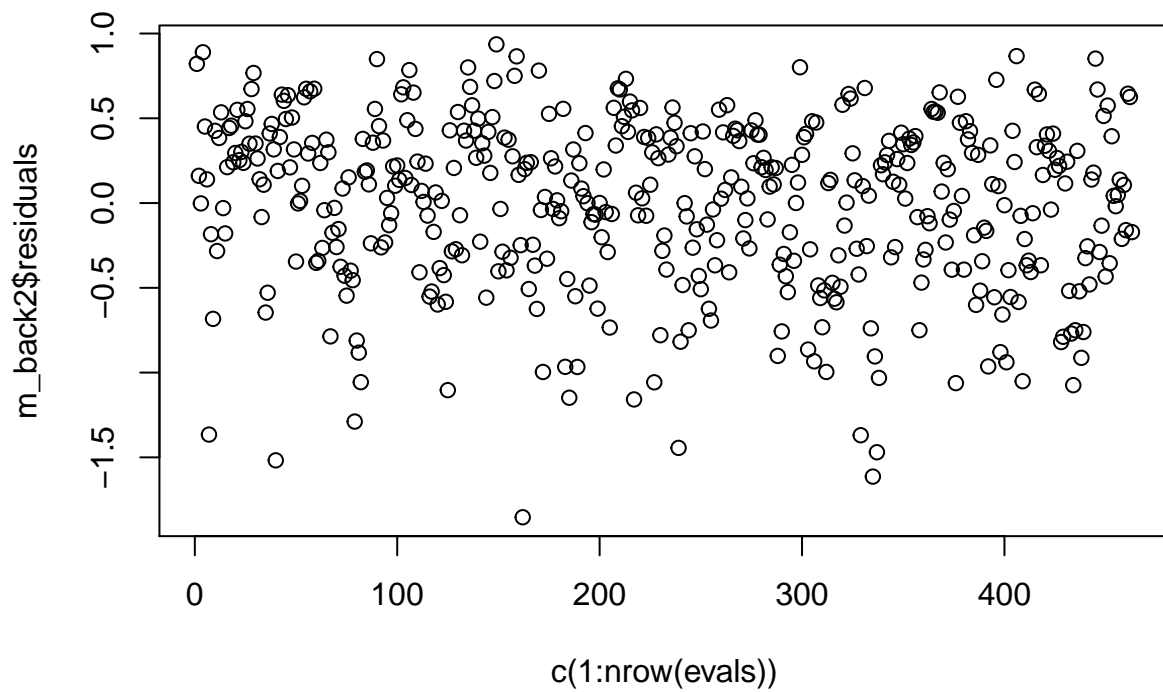
```
plot(abs(m_back2$residuals) ~ m_back2$fitted.values)
```

There some outliers although overall, most of the residual values are close to the fitted values.

3.The residuals are independent

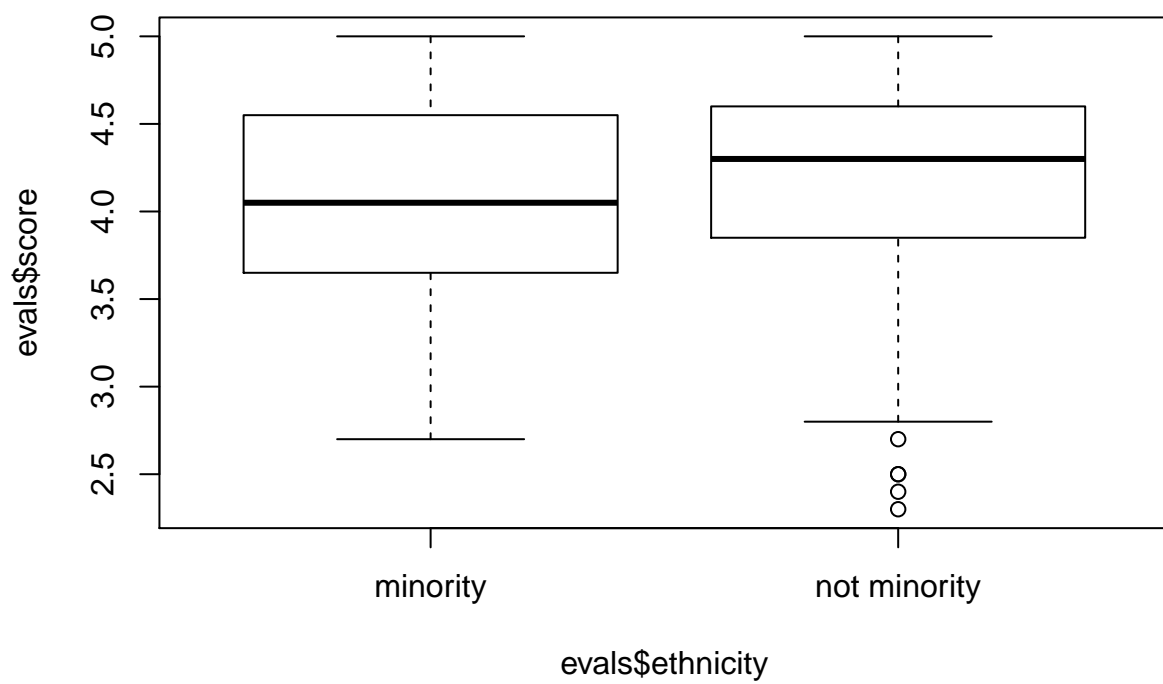
```
plot(m_back2$residuals ~ c(1:nrow(evals)))
```



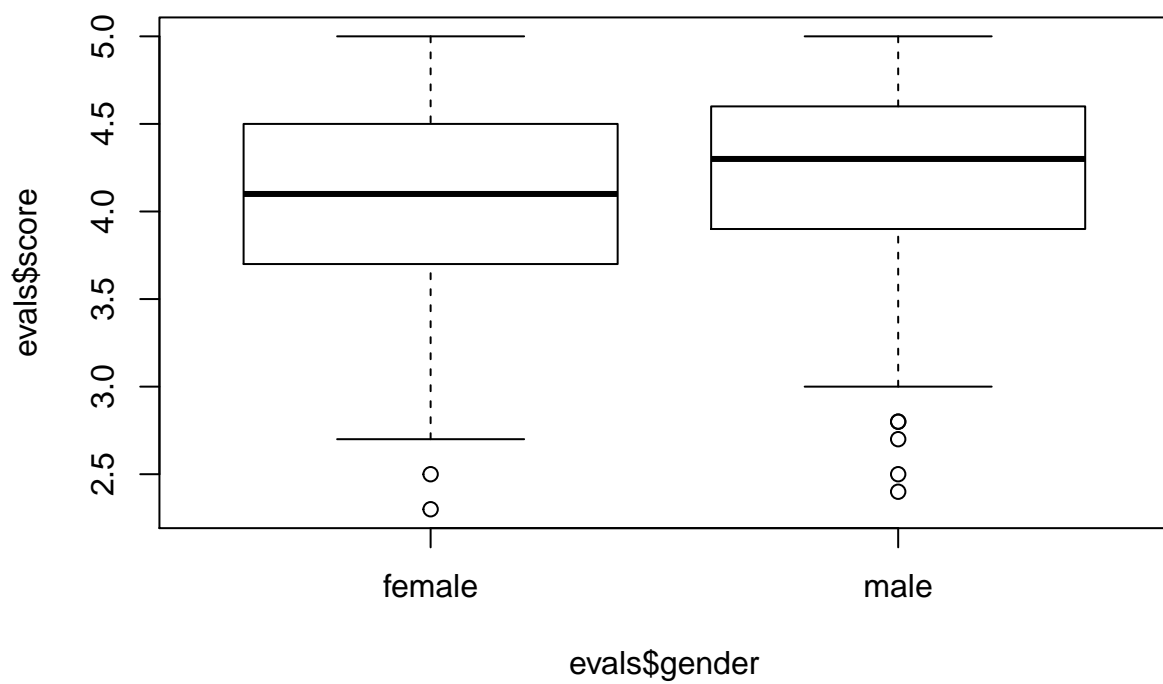
Yes, this condition is met. The residuals based on the sequence when it was gathered shows that they were randomly gathered.

4. Each variable is linearly related to the outcome.

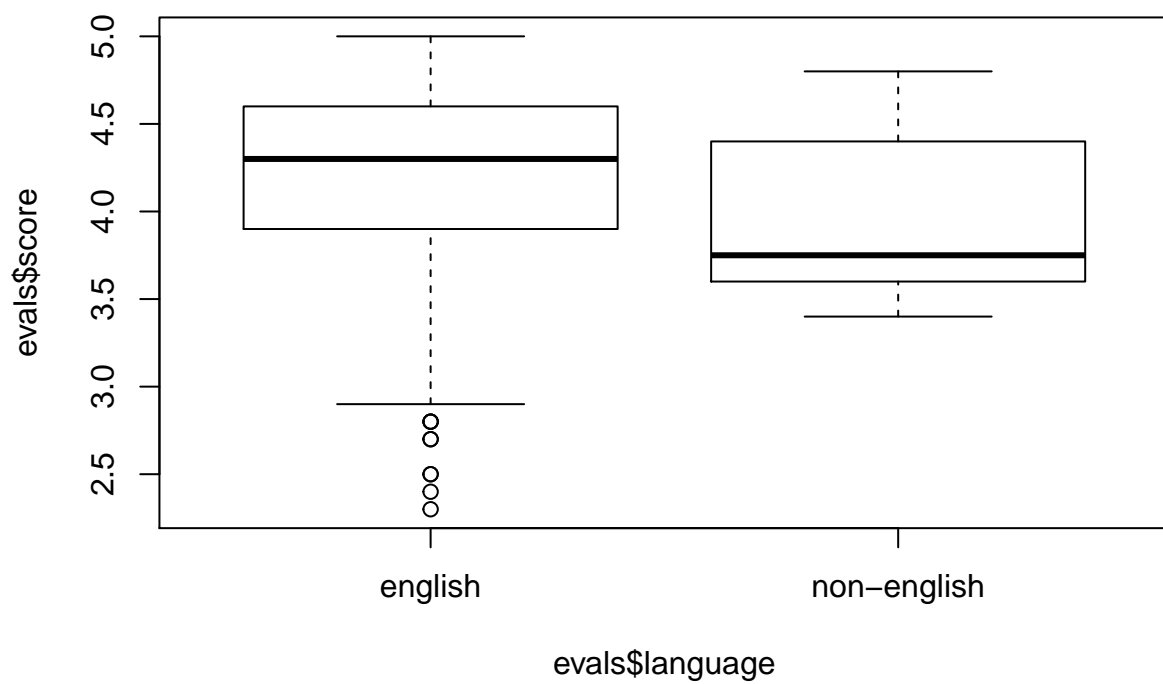
```
plot(evals$score ~ evals$ethnicity)
```



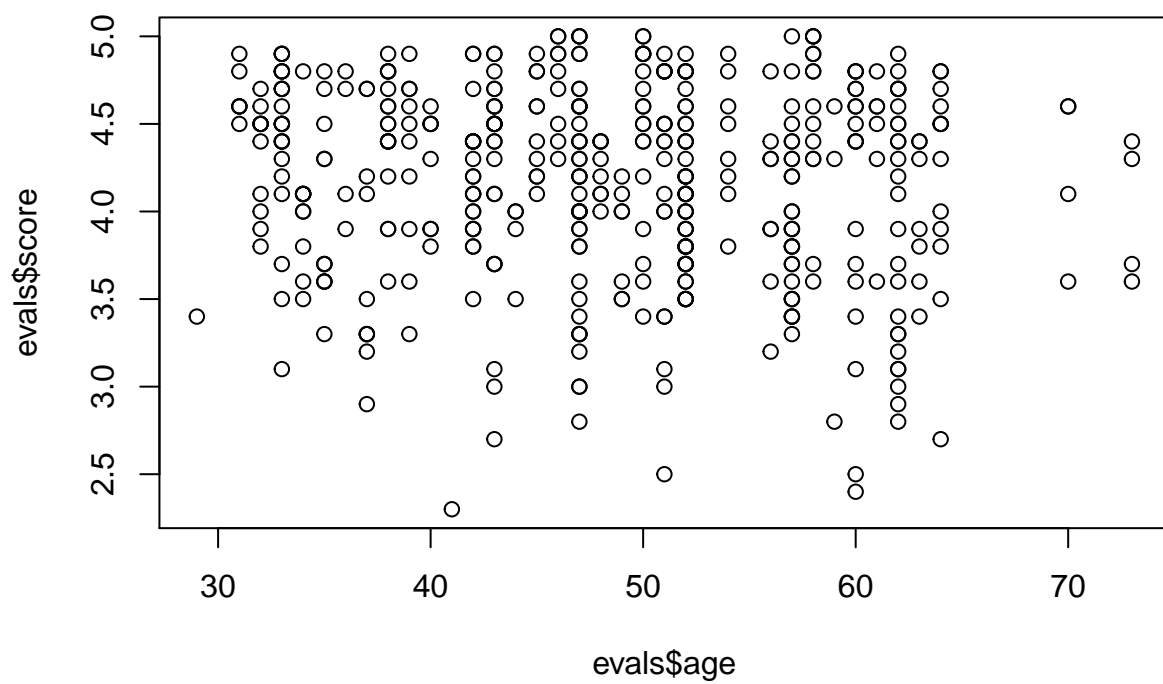
```
plot(evals$score ~ evals$gender)
```



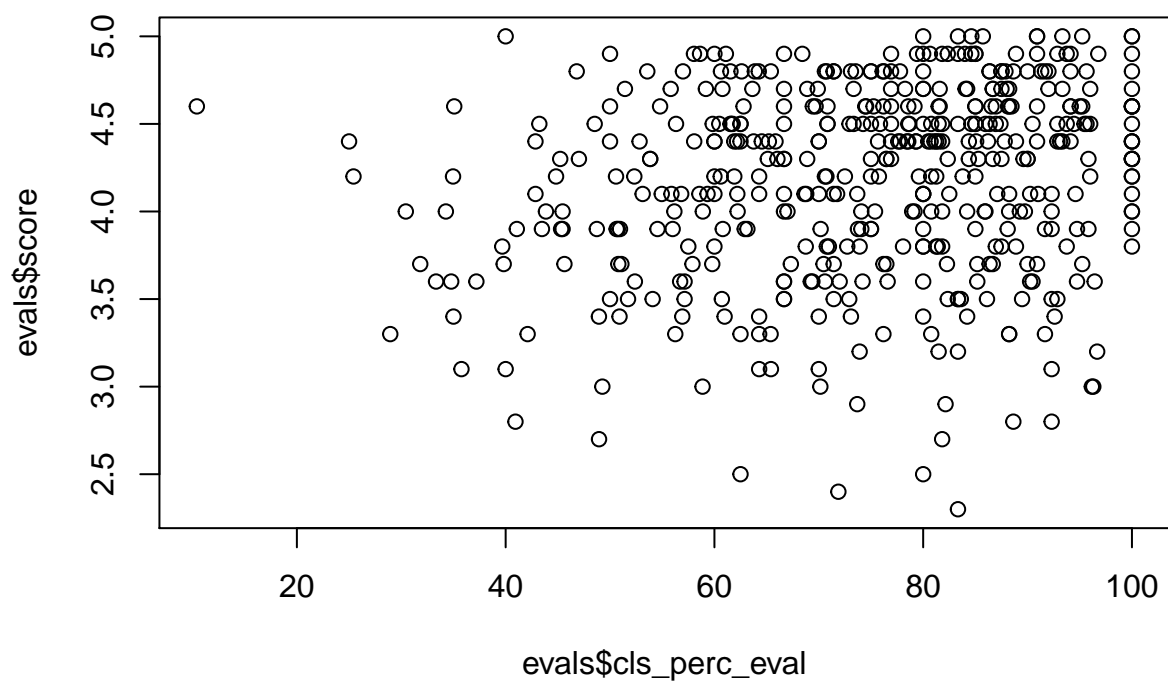
```
plot(evals$score ~ evals$language)
```



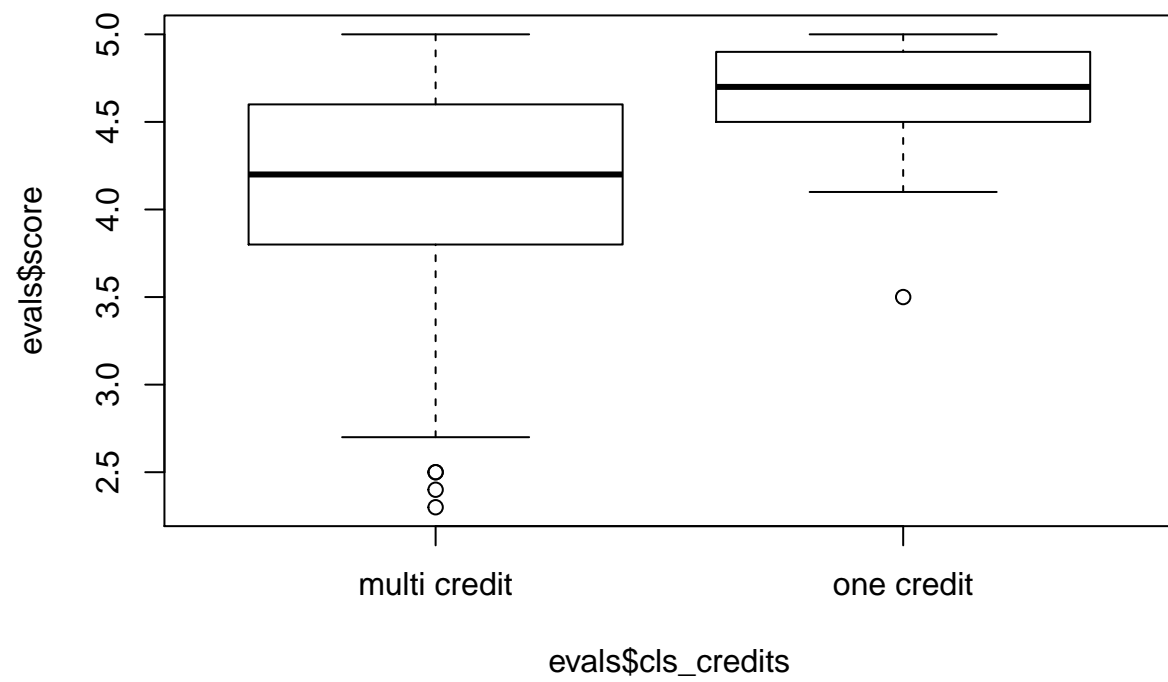
```
plot(evals$score ~ evals$age)
```



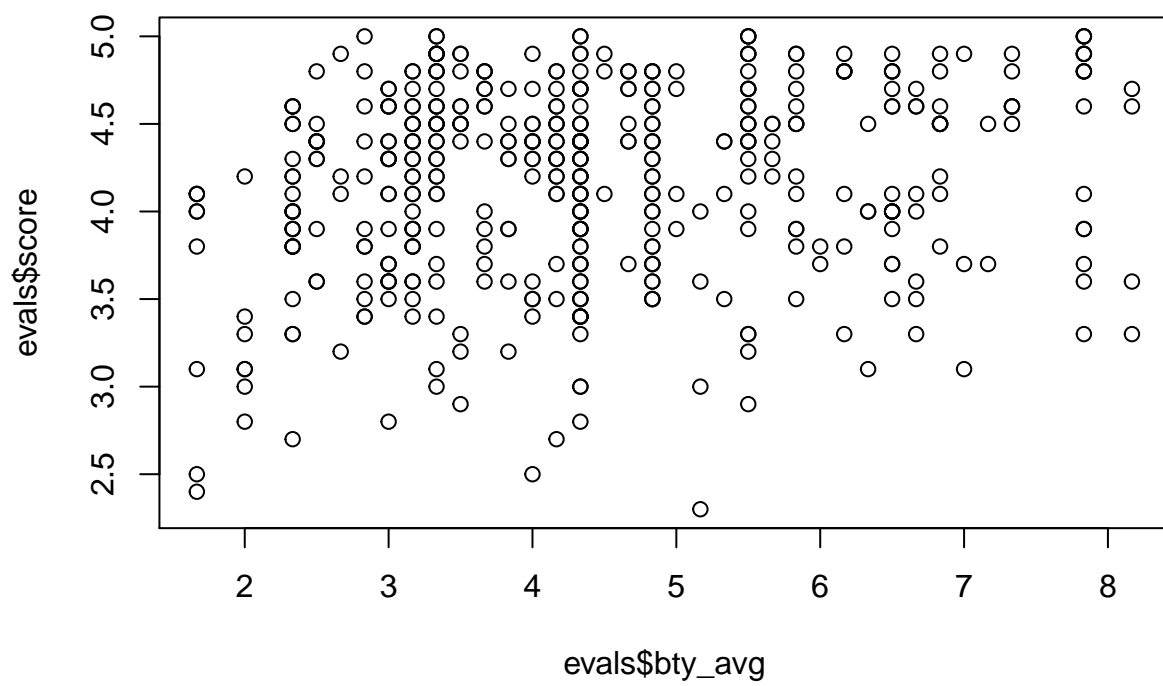
```
plot(evals$score ~ evals$cls_perc_eval)
```



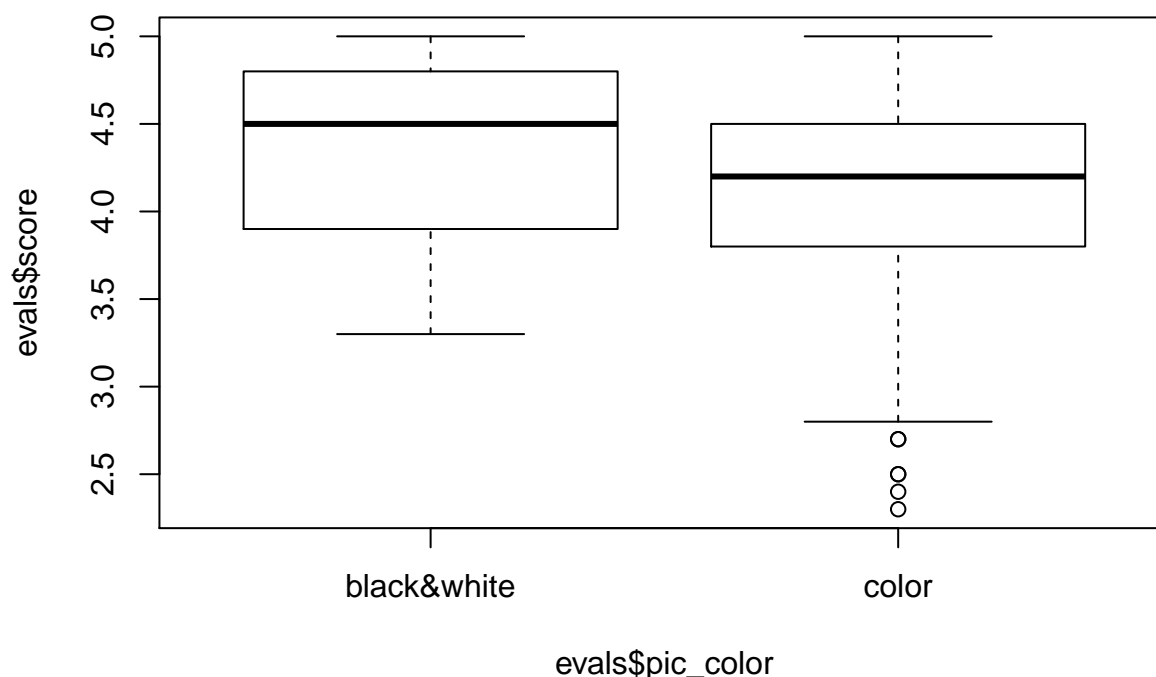
```
plot(evals$score ~ evals$cls_credits)
```



```
plot(evals$score ~ evals$bty_avg)
```

```
plot(evals$score ~ evals$pic_color)
```



The variables above are linearly related to the score - some more so than others.

17. The original paper describes how these data were gathered by taking a sample of professors from the University of Texas at Austin and including all courses that they have taught. Considering that each row represents a course, could this new information have an impact on any of the conditions of linear regression? **Answer :** No. Class courses are independent of each other so evaluation scores from one course is independent of the other even if the course is being taught by the same professor.
18. Based on your final model, describe the characteristics of a professor and course at University of Texas at Austin that would be associated with a high evaluation score. **Answer :** The professor is not a minority and male, must have graduated from an American (or English speaking) school and teaches a one credit course. He must also have a high beauty average score from the students and the professor's class photo should be in black and white. He must also be relatively young. And a good percentage of his class must have completed the evaluation.
19. Would you be comfortable generalizing your conclusions to apply to professors generally (at any university)? Why or why not? **Answer :** No. The sample size of 6 is too small. Also, some of the predictor variables are subjective and may vary with culture. Beauty, for one, is in the eye of the beholder. Picture preferences may also be culturally biased.

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