Final project

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```
rm(list = ls())
library(ggplot2)
library(ggmosaic)
library(dplyr)
library(lmerTest)
library(lme4)
library(lmeresampler)
library(chi)
library(VGAM)
```

```
library(jsonlite)
df <- fromJSON("https://query.data.world/s/204phxd36r6xkjug2gcnnrgzbr4bla")</pre>
```

1. The data is found at data.world (https://data.world/durhamnc/north-carolina-school-performance-data/workspace/file?filename=north-carolina-school-performance-data_1.csv) and it is about North Carolina school performance. There are 8 variables with 2425 observation chosen (each observation is a school). For this project I will ignore the type of school:

```
district: the district of the school school: the name of the school growthstatus: the academic growth status of the school (NotMet, Met,Exceeded) pctpassingseogtest: percentage of students passing EOG Math & English Test pctfreereduced: percentage of students who receive free or reduced lunch hasstringprogram: whether the school has a string program (0,1) schoolscore: school performance grade score grade: school grade (A-F)
```

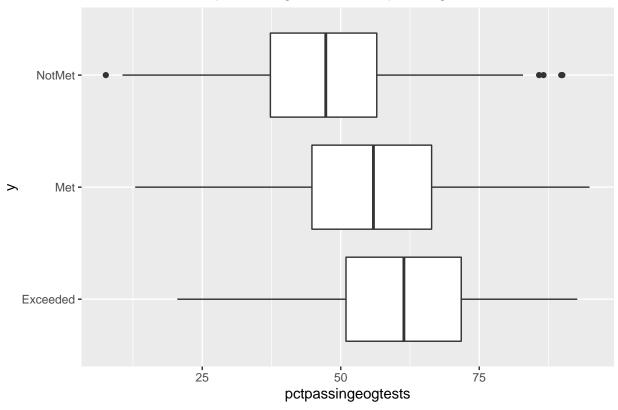
The questions of interest here are does school score, percentage of students passing eog test and percentage of students of get free or reduced lunch significant in determine growth status of the schools in North Carolina and does district and growth status affect school score?

2. EDA GLM:

```
df <- df%>% filter(growthstatus !="N/A")%>%
  mutate(y=growthstatus)

ggplot(data=df, aes(x=pctpassingeogtests,y=y, na.rm = TRUE))+
  geom_boxplot(na.rm = TRUE)+ ggtitle("Growth status vs percentage of student passing")
```

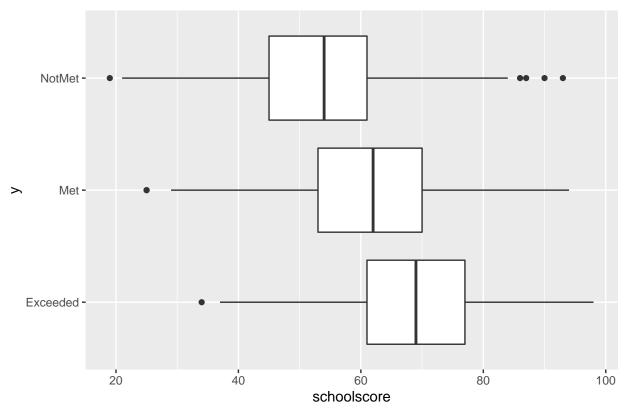




From the plot, it seems like percentage of passing EOG test does not have a significant effect of growth status of school. But, some school at lower percentage do not meet the growth status. But overall, the mean percentage are different for the 3 growth status

```
ggplot(data=df, aes(x=schoolscore,y=y, na.rm = TRUE))+
geom_boxplot(na.rm = TRUE)+ ggtitle("Growth status vs school score")
```

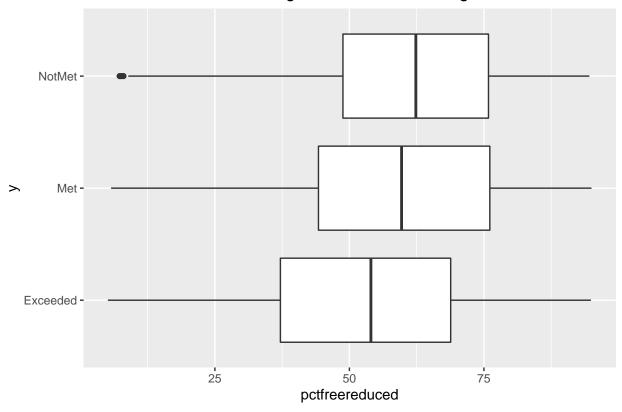
Growth status vs school score



From the plot, schools that have higher score tend to exceeded the growth status although there are some that don't met. But it does seem like school with lower school score have not yet met the growth status

```
ggplot(data=df, aes(x=pctfreereduced,y=y, na.rm = TRUE))+
geom_boxplot(na.rm = TRUE)+ ggtitle("Growth status vs Percentage of students receiving free or reduced.")
```

Growth status vs Percentage of students receiving free or reduced lunc



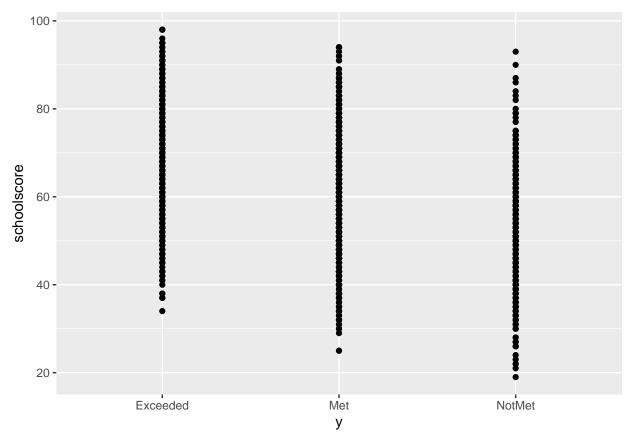
From the plot, although the mean are different, overall it does not seem like percentage of student receiving free or reduced lunch has any effect on growth status.

Multilevel: EDA for multilevel data has to take into account the differently-sized experimental units; in this case, we have districts (level 2 experiment unit) and school (level 1 experimental unit).

Level 1: Since level 1 is school, I assess individual growth status. The plot b

```
ggplot(data=df, aes(x=y,y=schoolscore)) + geom_point(binaxis="y")
```

Warning: Ignoring unknown parameters: binaxis



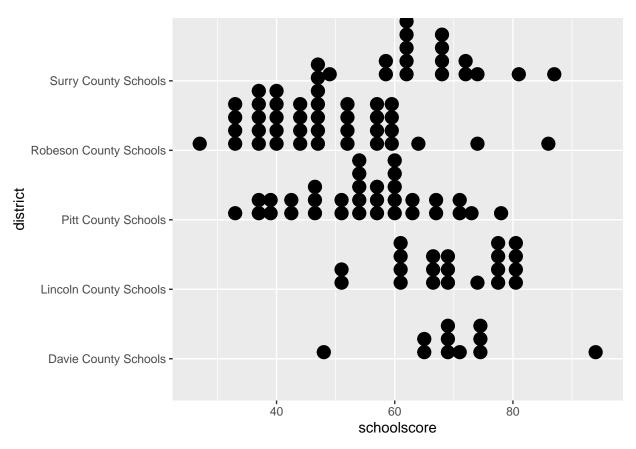
Since level 1 is school, I assess individual growth status. The plot shows that schools that has not yet met the growth status has lower school score and schools that has exceeded the growth status has higher school score.

Level 2:

```
id1 <-df%>% filter(district == "Surry County Schools")
id2 <-df%>% filter(district == "Robeson County Schools")
id3 <-df%>% filter(district == "Davie County Schools")
id4 <-df%>% filter(district == "Pitt County Schools")
id5 <-df%>% filter(district == "Lincoln County Schools")

df2<- bind_rows(id1,id2,id3,id4,id5)
ggplot(data=df2, aes(x=district, y=schoolscore)) + geom_dotplot(binaxis = "y") + coord_flip()</pre>
```

'stat_bindot()' using 'bins = 30'. Pick better value with 'binwidth'.



Level 2 is district and since there are 117 districts, I will look at 5 districts. From the plot, it can clearly be seen that every district has different school score. However, the districts have different number of school

3. Analysis and Conclusion

GLM: Since growth status of school has 3 categorical variables, multinomial model is chosen. the proportional-odds model is considered and it has 5 parameters (2 intercepts, 3 slopes. The non proportional-odds model has 8 parameter (2 intercepts, 6 slopes). The degree of freedom is 8-5=3 and the p-value that is calculated is extremely low so we can reject the proportional-odds model. Therefore, school score, percentage passing eog test and percentage of students who receive free or reduced lunch do contribute to school growth status because the variables all have low p-value. But the model chosen would be the non-proportional one.

```
fit<- vglm(y~schoolscore+pctpassingeogtests+pctfreereduced,cumulative(parallel=TRUE), data = df)
fit1<- vglm(y~schoolscore+pctpassingeogtests+pctfreereduced,cumulative(parallel=FALSE), data = df)
L0 <- logLik(fit)
L1 <- logLik(fit1)
1-pchisq(2*(L1-L0),3)</pre>
```

[1] 5.541058e-07

The interaction between the explanatory variables are not considered in the model because the log-likehood is ${\rm N/A}$

```
fit2<- vglm(y~(schoolscore+pctpassingeogtests+pctfreereduced)^2,cumulative(parallel=TRUE), data = df)
##
## Call:
## vglm(formula = y ~ (schoolscore + pctpassingeogtests + pctfreereduced)^2,
       family = cumulative(parallel = TRUE), data = df)
##
##
## Coefficients:
##
                        (Intercept):1
                                                            (Intercept):2
##
                        -2.923138e+01
                                                            -2.667953e+01
##
                          schoolscore
                                                       pctpassingeogtests
##
                         4.219643e-01
                                                             1.080006e-01
##
                       pctfreereduced
                                          schoolscore:pctpassingeogtests
##
                         1.860048e-01
                                                            -2.090057e-03
##
          schoolscore:pctfreereduced pctpassingeogtests:pctfreereduced
                        -9.912245e-04
##
                                                            -1.278731e-03
##
## Degrees of Freedom: 4614 Total; 4606 Residual
## Residual deviance: 4023.471
## Log-likelihood: -2011.735
Multilevel: The p-value here is 0.7219. We fail to reject the null hypothesis. The extra term (district:growth)
is unnecessary. The reduced model is preferred
model <- lmer(schoolscore~district + growthstatus + district:growthstatus + (1|school), REML = T,data =
## fixed-effect model matrix is rank deficient so dropping 30 columns / coefficients
model1<-lmer(schoolscore~district + growthstatus+ (1 | school), REML = T, data = df)
anova(model, model1)
## refitting model(s) with ML (instead of REML)
## Data: df
## Models:
## model1: schoolscore ~ district + growthstatus + (1 | school)
## model: schoolscore ~ district + growthstatus + district:growthstatus +
## model:
               (1 | school)
##
                       BIC logLik deviance Chisq Df Pr(>Chisq)
                  AIC
          npar
## model1 121 18565 19265 -9161.6
                                        18323
           323 18780 20648 -9066.8
                                        18134 189.76 202
                                                              0.7219
To confirm whether district or growthstatus alone would be a better fit to the model, the following anova
test is carried. Both the pvalue for term district alone and growthstatus alone are < 2.2e-16. Which means
that we can reject the null hypothesis. model 1 is preferred.
model2 <- lmer(schoolscore~district + (1 | school), REML = T, data = df)</pre>
## boundary (singular) fit: see ?isSingular
```

```
anova(model1,model2)
## refitting model(s) with ML (instead of REML)
## Data: df
## Models:
## model2: schoolscore ~ district + (1 | school)
## model1: schoolscore ~ district + growthstatus + (1 | school)
         npar AIC BIC logLik deviance Chisq Df Pr(>Chisq)
## model2 119 19015 19703 -9388.4
                                   18777
## model1 121 18565 19265 -9161.6
                                  18323 453.54 2 < 2.2e-16 ***
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
model3 <- lmer(schoolscore~growthstatus + (1 |school), REML = T, data = df)</pre>
## boundary (singular) fit: see ?isSingular
anova(model1,model3)
## refitting model(s) with ML (instead of REML)
## Data: df
## Models:
## model3: schoolscore ~ growthstatus + (1 | school)
## model1: schoolscore ~ district + growthstatus + (1 | school)
         npar AIC BIC logLik deviance Chisq Df Pr(>Chisq)
## model3
           5 18952 18981 -9471.0
                                  18942
## model1 121 18565 19265 -9161.6
                                    18323 618.66 116 < 2.2e-16 ***
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
```