

# ECON883\_HW1

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## Introduction

1. Paper: Peer effects, teacher incentives, and the impact of tracking: Evidence from a randomized evaluation in Kenya. *American economic review*.

2. Abstract of the paper:

To the extent that students benefit from high-achieving peers, tracking will help strong students and hurt weak ones. However, all students may benefit if tracking allows teachers to better tailor their instruction level. Lower-achieving pupils are particularly likely to benefit from tracking when teachers have incentives to teach to the top of the distribution. We propose a simple model nesting these effects and test its implications in a randomized tracking experiment conducted with 121 primary schools in Kenya. While the direct effect of high-achieving peers is positive, tracking benefited lower-achieving pupils indirectly by allowing teachers to teach to their level.

3. Data source: <https://www.openicpsr.org/openicpsr/project/112446/version/V1/view>
4. This paper used regression discontinuity (RD) design to test whether students at the median are better off being assigned to the top section (measured by percentile of student).
5. Model:

$$y_{ij} = \delta B_{ij} + \lambda_1 P_{ij} + \lambda_2 P_{ij}^2 + \lambda_3 P_{ij}^3 + X_{ij}\beta + \epsilon_{ij}$$

where  $P_{ij}$  is the percentile of the child on the baseline distribution in his or her school,  $y_{ij}$  is the standardized test score,  $B_{ij}$  are control variables.

6. Variables:

1.  $y$  = standardized student's score
2.  $x$  = students' percentile in previous exams
3. control variables:
  - gender: dummy
  - age
  - teacher: dummy, whether the student was taught by civil-service teacher.

```

# remove variables from work space
rm(list = ls())

set.seed(2023) # for reproducibility

# LIBRARIES
library(haven) # for data loading
library(dplyr) # for easy data shaping

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
##   filter, lag

## The following objects are masked from 'package:base':
##
##   intersect, setdiff, setequal, union

library(ggplot2) # for plotting
library(binsreg) # for binscatter
library(rdrobust)
library(sandwich)

# kernel mean and local linear regression
source("/Users/shaoyutong/Library/Mobile Documents/com~apple~CloudDocs/ECON883/HW/HW1/plot_funs.R")
source("/Users/shaoyutong/Library/Mobile Documents/com~apple~CloudDocs/ECON883/HW/HW1/loc_lin.R")

path_figure_save <- "/Users/shaoyutong/Library/Mobile Documents/com~apple~CloudDocs/ECON883/HW/HW1/figure"

```

## Summary of major data set

`Student_test_data.dta` is the main data set in wide format (one observation per student).

It includes baseline characteristics of the students, their test scores at both the endline (fall 2006) and long-term follow-up (fall 2007) tests, and the “treatment” dummies – whether the school was sampled for “Tracking”, whether the student was assigned to the Contract Teacher, etc.

I simplify the data set by extracting variables of interest, including:

- $y$  = standardized student’s score
- $x$  = students’ percentile in previous exams
- control variables: gender, age, teacher

The scatter plot are as follows.

```
df1 <- read_dta('/Users/shaoyutong/Library/Mobile Documents/com~apple~CloudDocs/ECON883/HW/HW1/data/112')
df_pres <- read_dta('/Users/shaoyutong/Library/Mobile Documents/com~apple~CloudDocs/ECON883/HW/HW1/data/112')
```

```
# simplify data set
df_simp <- data.frame(SD_std_mark = df1$SDstream_std_mark,
                      MEAN_std_mark = df1$MEANstream_std_mark,
                      x = df1$percentile,
                      xx = df1$realpercentile,
                      total_score = df1$totalscore,
                      y = df1$std_mark,
                      gender = df1$girl,
                      age = df1$agetest,
                      teacher = df1$etpteacher
                      )
dat_use <- na.omit(df_simp) # remove missing values

head(dat_use)
```

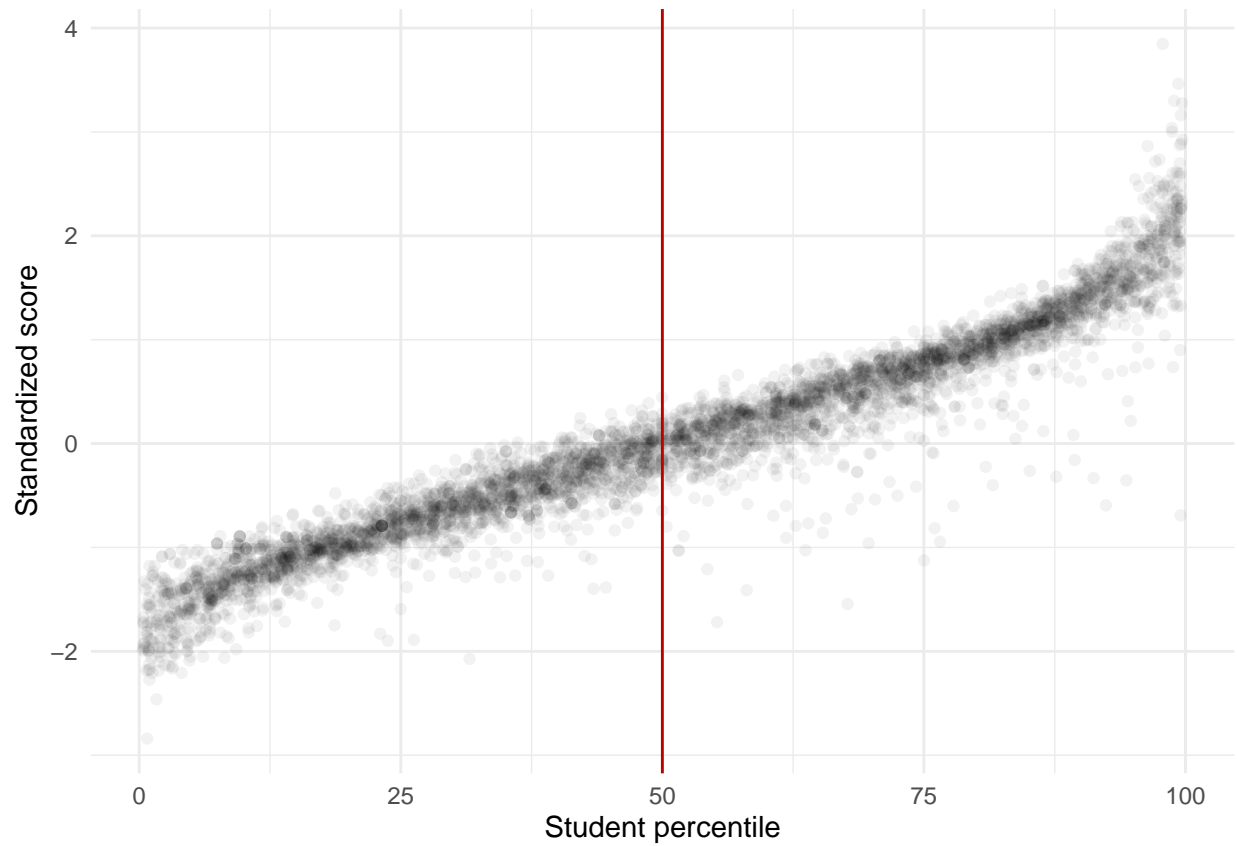
```
##   SD_std_mark MEAN_std_mark      x xx total_score      y gender age
## 3    0.6692453   -0.8192898  3.205132  4    2.9000001 -2.145025      1   8
## 4    0.6692453   -0.8192898  4.487181  5    0.2357143 -1.879484      0  14
## 5    0.6692453   -0.8192898  5.769229  6    9.8571424 -1.809605      0  11
## 6    0.6692453   -0.8192898  8.333338  9    6.5999999 -1.194667      0  10
## 7    0.6692453   -0.8192898 10.897434 11    5.1999998 -1.362377      0  10
## 8    0.6692453   -0.8192898 12.179488 13    9.7428570 -1.124788      1   9
##   teacher
## 3         0
## 4         0
## 5         0
## 6         0
## 7         0
## 8         0
```

```
summary(dat_use)
```

```
##   SD_std_mark      MEAN_std_mark      x      xx
## Min.   :0.1541   Min.   :-0.906560   Min.   : 0.3546   Min.   : 1.00
## 1st Qu.:0.5428   1st Qu.: -0.741663   1st Qu.:27.1792   1st Qu.: 28.00
## Median :0.7965   Median : 0.005767   Median :51.8182   Median : 52.00
## Mean   :0.7597   Mean   : 0.006725   Mean   :51.3407   Mean   : 51.83
## 3rd Qu.:0.9880   3rd Qu.: 0.737004   3rd Qu.:75.7501   3rd Qu.: 76.00
## Max.   :1.1925   Max.   : 0.889537   Max.   :99.7059   Max.   :100.00
##   total_score      y      gender      age
## Min.   : 0.000   Min.   :-2.84003   Min.   :0.0000   Min.   : 5.000
## 1st Qu.: 5.857   1st Qu.: -0.73795   1st Qu.:0.0000   1st Qu.: 8.000
## Median :11.286   Median :-0.01431   Median :0.0000   Median : 9.000
## Mean   :13.068   Mean   : 0.03059   Mean   :0.4913   Mean   : 9.318
## 3rd Qu.:18.861   3rd Qu.: 0.76442   3rd Qu.:1.0000   3rd Qu.:10.000
## Max.   :42.729   Max.   : 3.84869   Max.   :1.0000   Max.   :19.000
##   teacher
## Min.   :0.0000
```

```
## 1st Qu.:0.0000
## Median :1.0000
## Mean   :0.5026
## 3rd Qu.:1.0000
## Max.   :1.0000
```

```
# plot(x=df_simp$x, y = df_simp$y,
#       main = 'Standardized score vs student percentile',
#       xlab = 'percentile',
#       ylab = 'standardized score')
# x = df_simp$x
# y = df_simp$std_score
ggplot(mapping=aes(x=x,y=y)) +
  geom_point(data=dat_use, alpha= 0.05) +
  geom_vline(aes(xintercept=50), colour="#BB0000") +
  xlab('Student percentile') +
  ylab('Standardized score') +
  theme_mp()
```



## Single covariate

### Linear regression

I first run a basic linear regression on each side of the discontinuity. Note that the author used polynomial form, but I implemented a basic one for illustration.

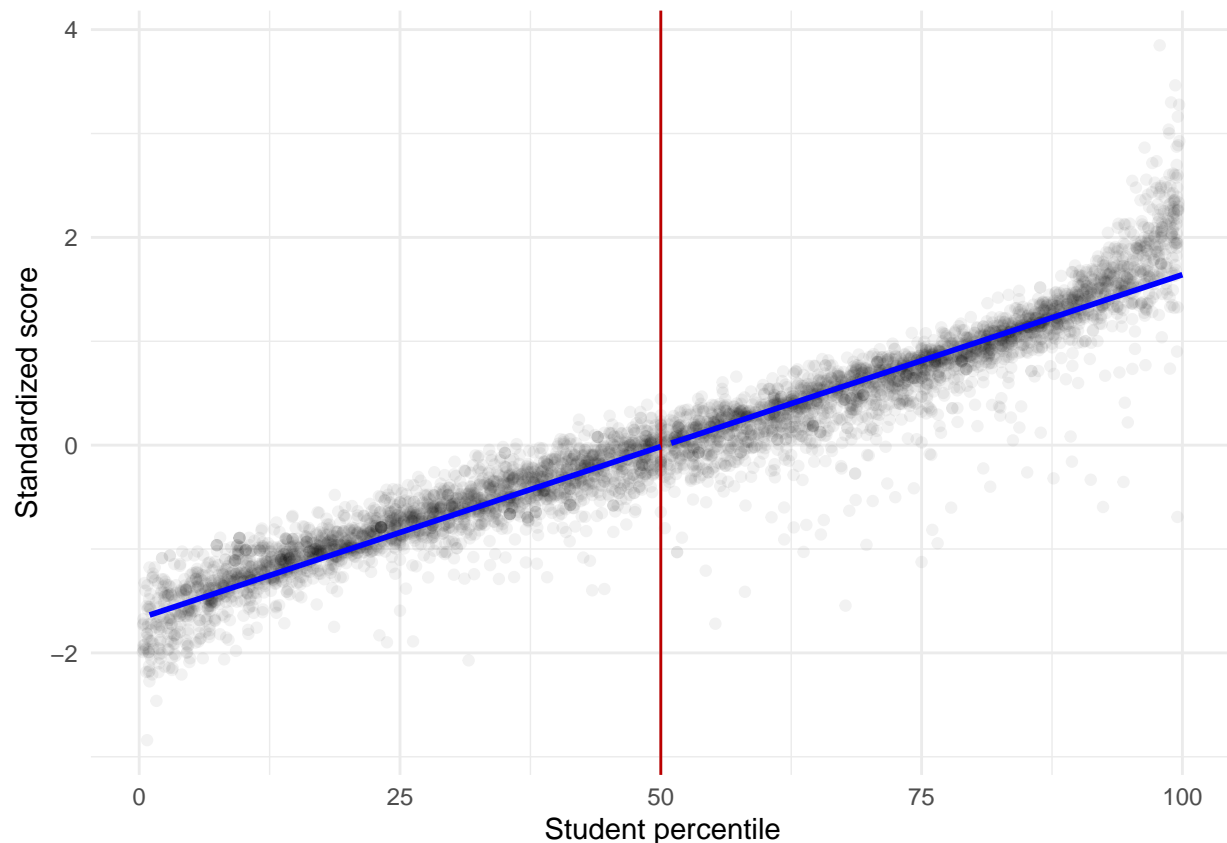
```
group1 <- subset(dat_use, x <= 50)
group2 <- subset(dat_use, x > 50)
lm_1 <- lm(formula = y ~ x, data=dat_use)
lm_2 <- lm(formula = y ~ x, data=dat_use)

# mean of first group
x_curve1 <- sort(unique(group1$xx))

y_curve1 <- predict(lm_1, newdata = data.frame(x=x_curve1))
dat_curve1 <- data.frame(y=y_curve1, x=x_curve1) # population mean curve (blue line)
# dat_curve1
# mean of second group
x_curve2 <- sort(unique(group2$xx))

y_curve2 <- predict(lm_2, newdata = data.frame(x=x_curve2))
dat_curve2 <- data.frame(y=y_curve2, x=x_curve2)

ggplot(mapping=aes(x=x,y=y)) +
  geom_point(data=dat_use, alpha= 0.05) +
  geom_vline(aes(xintercept=50), colour="#BB0000") +
  geom_line(data = dat_curve1, color='blue', linewidth=1.) +
  geom_line(data = dat_curve2, color='blue', linewidth=1.) +
  xlab('Student percentile') +
  ylab('Standardized score') +
  theme_mp()
```



## Kernel Regression

I then applied kernel regression using triangular kernel with bandwidth from 1 to 10. And plotted regression results for 4 of them.

```
xx <- sort(unique(dat_use$xx))
# xx
# xx <- seq(1, 100, 2)

yy <- kernel_mean(y=dat_use$y, x=dat_use$x, xx=xx, hs=1:10)

# yy
# create plots to illustrate
for (h in c(1,3,5,10)) {
  ggplot(mapping=aes(x=x,y=y)) +
    geom_point(data=dat_use, alpha=0.03) +
    geom_vline(aes(xintercept=50), colour="#BB0000") +
    geom_line(data = dat_curve1, color='blue', linewidth=1.) +
    geom_line(data = dat_curve2, color='blue', linewidth=1.) +
    geom_point(data=data.frame(x=xx,y=yy[,h]), color="orange", size=1) +
    xlab('Student Percentile') +
    ylab('Standardized score') +
    ggtitle(paste("h", "=", h, " ", "triangular kernel")) +
    theme_mp()
}
```

```

saveplot(paste0(path_figure_save,"triangular_h",h))
}

```

## Local Linear Regression

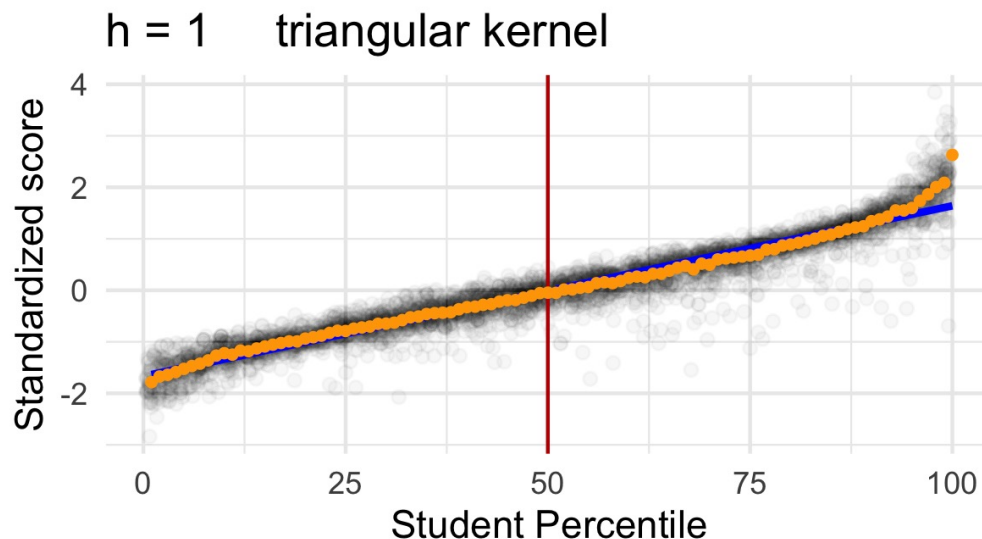
```

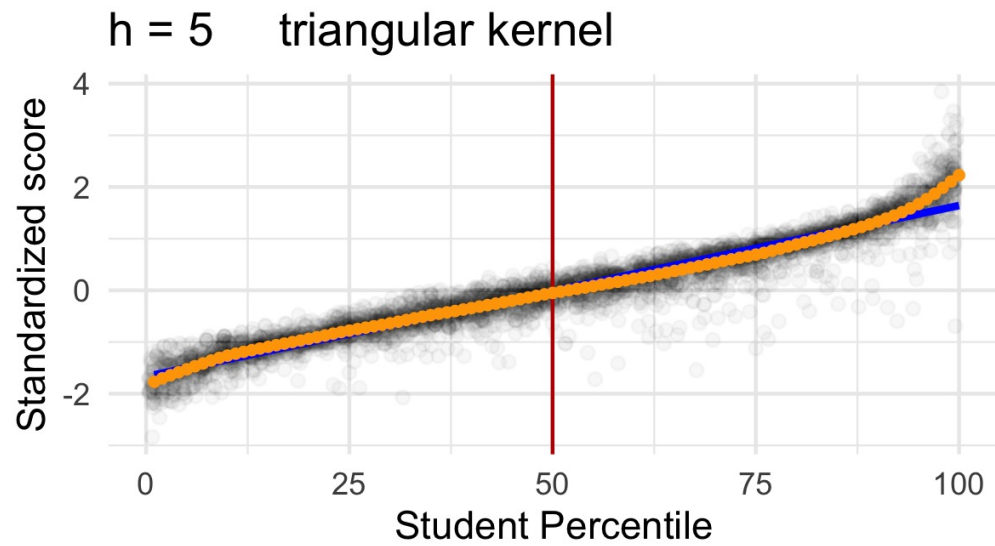
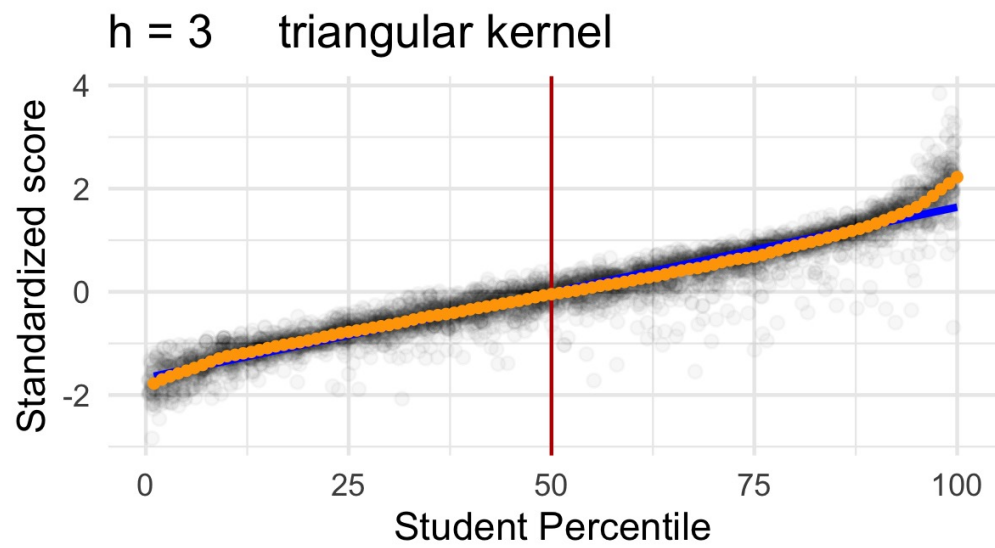
kern <- "triangular"
xx <- sort(unique(dat_use$xx)) # evaluation points
reg_loclin <- loclin_reg(y=dat_use$y, x=dat_use$x, xx=xx, hs=1:10, kernel=kern)

yy <- reg_loclin$yy
# yy
# create plots to illustrate
saveplot <- function(filename,
                        plot=last_plot(),
                        width=4, height=2.25, units="in") {
  ggsave(filename=paste0(filename,".jpg"), plot=plot, width=width, height=height, units=units)
}

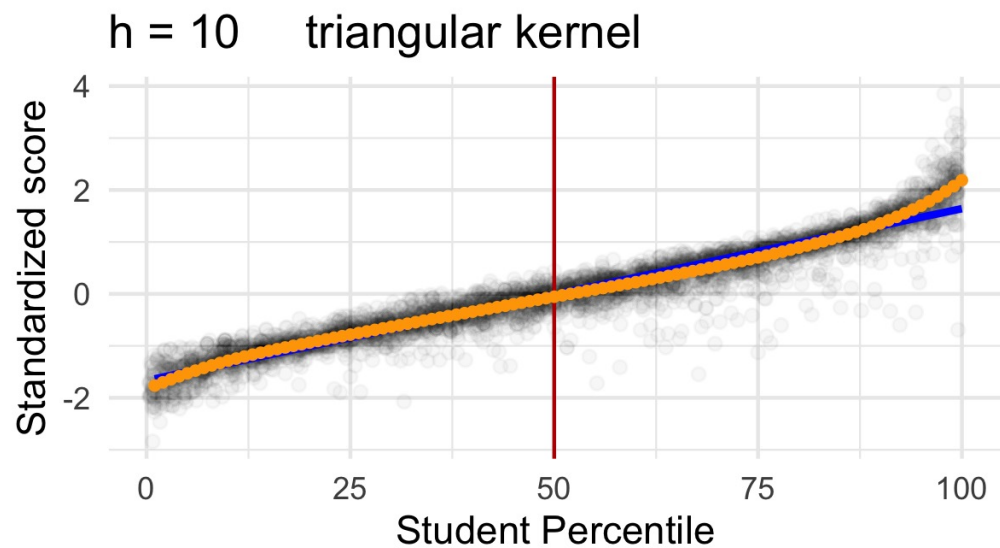
for (h in c(1,3,5,10)) {
  ggplot(mapping=aes(x=x,y=y)) +
    geom_point(data=dat_use, alpha=0.03) +
    geom_vline(aes(xintercept=50), colour="#BB0000") +
    geom_line(data = dat_curve1, color='blue', linewidth=1.) +
    geom_line(data = dat_curve2, color='blue', linewidth=1.) +
    geom_point(data=data.frame(x=xx,y=yy[,h]), color="orange", size=1) +
    xlab('Student Percentile') +
    ylab('Standardized score') +
    ggtitle(paste("h", "=", h, " ", "triangular kernel")) +
    theme_mp()
  saveplot(paste0(path_figure_save,"loclin_",kern,"_h",h))
}

```







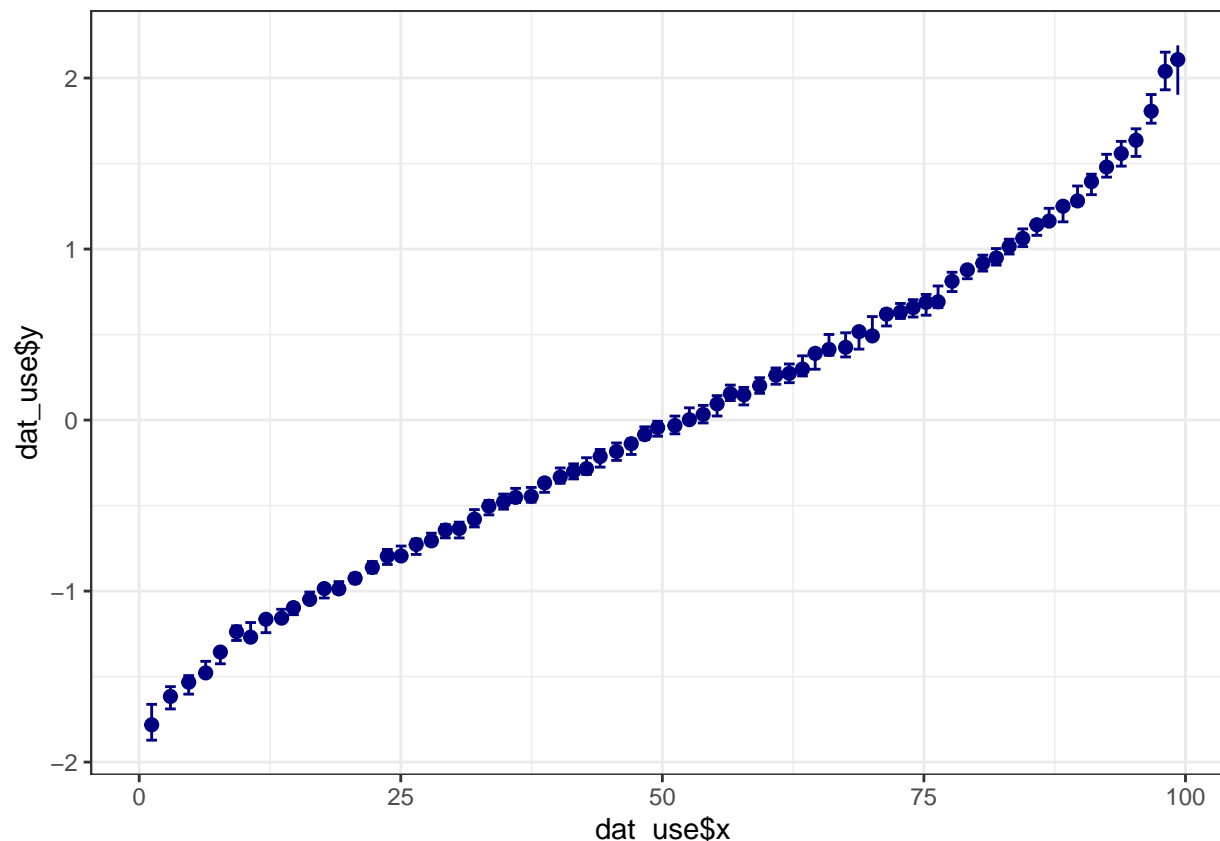


## Binscatter Regression

Next I implemented binscatter regression using R package. Results are as follows.

```
# binscatter  
breg <- binsreg(dat_use$y, dat_use$x, ci=c(3,3))
```

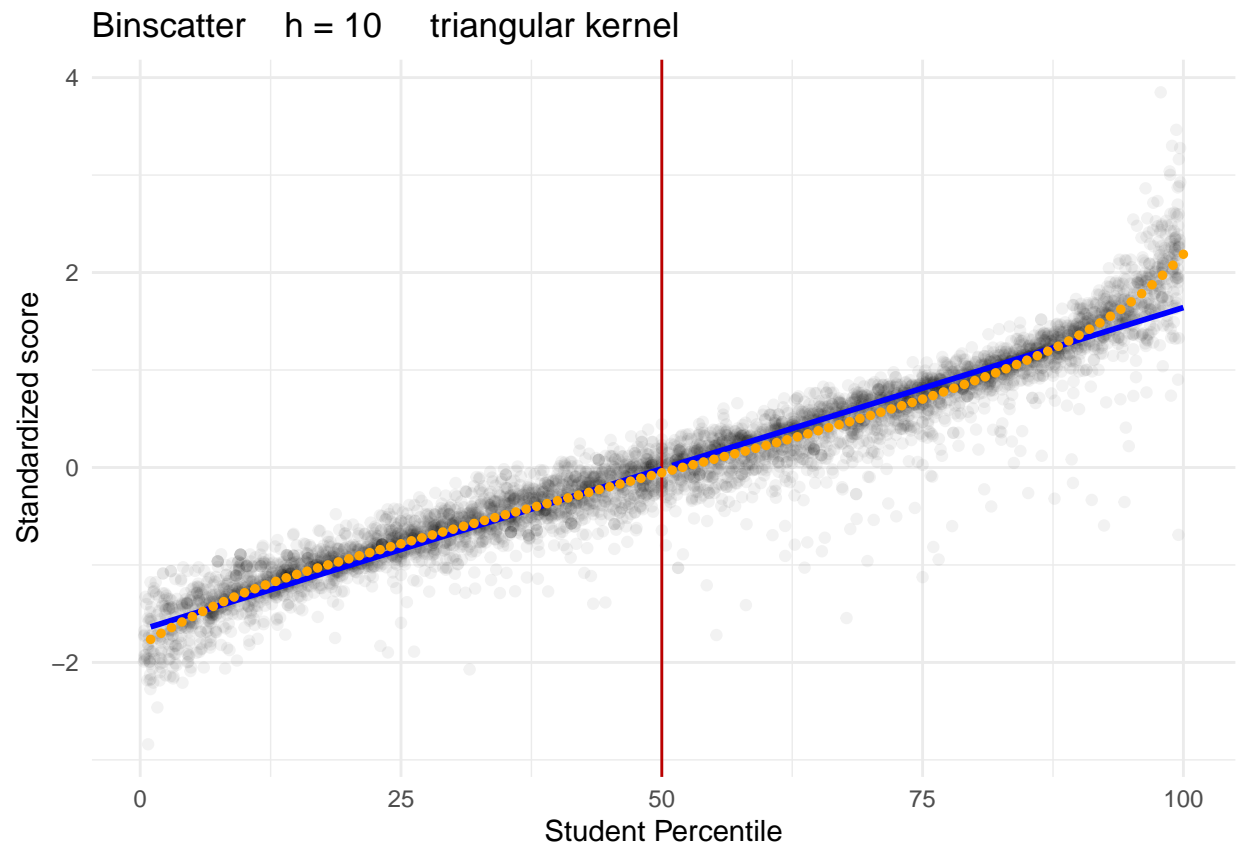
```
## Warning in binsreg(dat_use$y, dat_use$x, ci = c(3, 3)): To speed up computation,  
## bin/degree selection uses a subsample of roughly max(5,000, 0.01n) observations  
## if the sample size n>5,000. To use the full sample, set randcut=1.
```



```
# ci=c(3,3) is what the authors recommend to calculate standard errors
# can also play with many other options to get local linear, etc.

# for ggplot2 plot, grab data underlying figure:
df_breg <- data.frame(x=breg$data.plot$`Group Full Sample`$data.dots$x,
                      y=breg$data.plot$`Group Full Sample`$data.dots$fit#,
                      # ymin=breg$data.plot$`Group Full Sample`$data.ci$ci.l,
                      # ymax=breg$data.plot$`Group Full Sample`$data.ci$ci.r
                      )

ggplot(data=df_breg, aes(x=x,y=y)) +
  geom_point(data=dat_use, alpha=0.05) +
  # geom_errorbar(aes(ymin=min(y),ymax=max(y)), color='blue') +
  # geom_line(data=dat_curve, color="blue", linewidth=1.5) +
  # geom_point(color='orange',size=2) +
  geom_vline(aes(xintercept=50), colour="#BB0000") +
  geom_line(data = dat_curve1, color='blue', linewidth=1.) +
  geom_line(data = dat_curve2, color='blue', linewidth=1.) +
  geom_point(data=data.frame(x=xx,y=yy[,h]), color="orange", size=1) +
  xlab('Student Percentile') +
  ylab('Standardized score') +
  ggtitle(paste('Binscatter', ' ', "h", "=", h, " ", "triangular kernel")) +
  theme_mp()
```



```
saveplot(paste0(path_figure_save, "binscatter"))
```

## Controlling other variables

Incorrect regression: partialing out covariates before binscatter regression

```
source("/Users/shaoyutong/Library/Mobile Documents/com~apple~CloudDocs/ECON883/HW/HW1/plot_funs.r")
```

```
nbins <- 20
colnames(dat_use)
```

```
## [1] "SD_std_mark" "MEAN_std_mark" "x" "xx"
## [5] "total_score" "y" "gender" "age"
## [9] "teacher"
```

```
# group 1
x1 <- group1$x
y1 <- group1$y
w_gp1 <- group1[,c('gender', 'age', 'teacher')]
names(w_gp1) <- c('w1', 'w2', 'w3')
# incorrect: residualize x and y w.r.t. w, then use binscatter on residuals
x_resid1 <- lm(x1 ~ w1 + w2 + w3, data=w_gp1)$residuals
```

```

y_resid1 <- lm(y1 ~ w1 + w2 + w3, data=w_gp1)$residuals
true1 <- predict(lm_1, newdata = data.frame(x=x_resid1))

df_gp1 <- data.frame(x=x1, y=y1, x_resid=x_resid1, y_resid=y_resid1,
                     true=true1)

# group 2
x2 <- group2$x
y2 <- group2$y
w_gp2 <- group2[,c('gender','age','teacher')]
names(w_gp2) <- c('w1', 'w2', 'w3')
# incorrect: residualize x and y w.r.t. w, then use binscatter on residuals
x_resid2 <- lm(x2 ~ w1 + w2 + w3, data=w_gp2)$residuals
y_resid2 <- lm(y2 ~ w1 + w2 + w3, data=w_gp2)$residuals
true2 <- predict(lm_2, newdata = data.frame(x=x_resid2))
df_gp2 <- data.frame(x=x2, y=y2, x_resid=x_resid2, y_resid=y_resid2,
                     true=true2)

df_all <- rbind(df_gp1, df_gp2)
colnames(df_all)

```

```
## [1] "x"      "y"      "x_resid" "y_resid" "true"
```

```

# df1 <- data.frame(y1=y1, x1=x1, x_resid=x_resid, y_resid=y_resid)
ggplot(data=df_all, aes(x=x_resid, y=y_resid)) +
  # geom_line(aes(x=dat_curve1$x, y=dat_curve1$y), color='blue', linewidth=1.) +
  # geom_line(aes(x=dat_curve2$x, y=dat_curve2$y), color='blue', linewidth=1.) +
  geom_line(aes(y=true), size=1, color='blue') +
  # geom_line(data = df_gp2, size=1, color='blue') +
  stat_summary_bin(fun.data = mean_se, bins=nbins, size= 0.4, color='orange') +
  stat_summary_bin(fun='mean', bins=nbins,
                  color='orange', size=1) +
  ggtitle('WRONG: Y-E(Y|Z) ~ bins(X - E(X|Z))') +
  theme_mp()

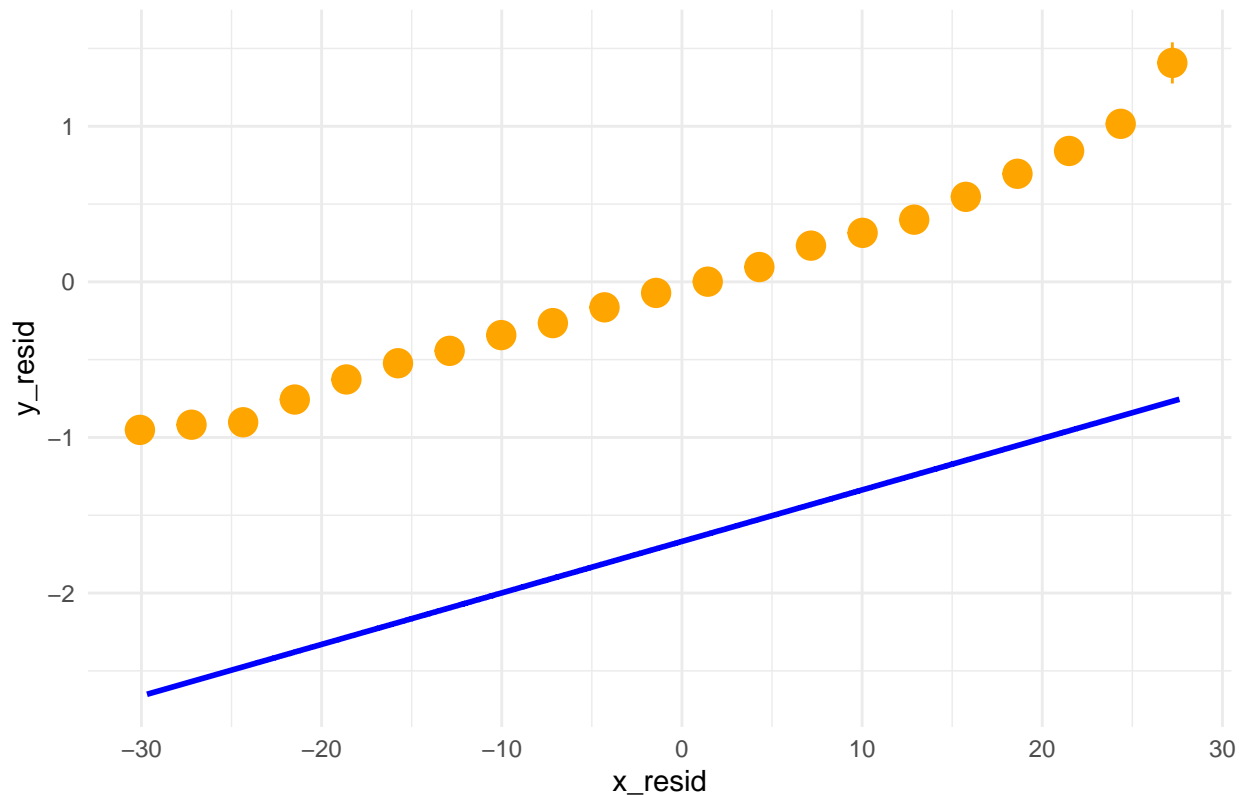
```

```
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
```

```
## Warning: Removed 1 rows containing missing values ('geom_segment()').
```

```
## Warning: Removed 21 rows containing missing values ('geom_segment()').
```

WRONG:  $Y - E(Y|Z) \sim \text{bins}(X - E(X|Z))$



Correct regression

```
# colnames(dat_use)
# group 1
# group 1
lm_ctrl1 <- lm(y1 ~ x1 + gender + age + teacher, data=group1)

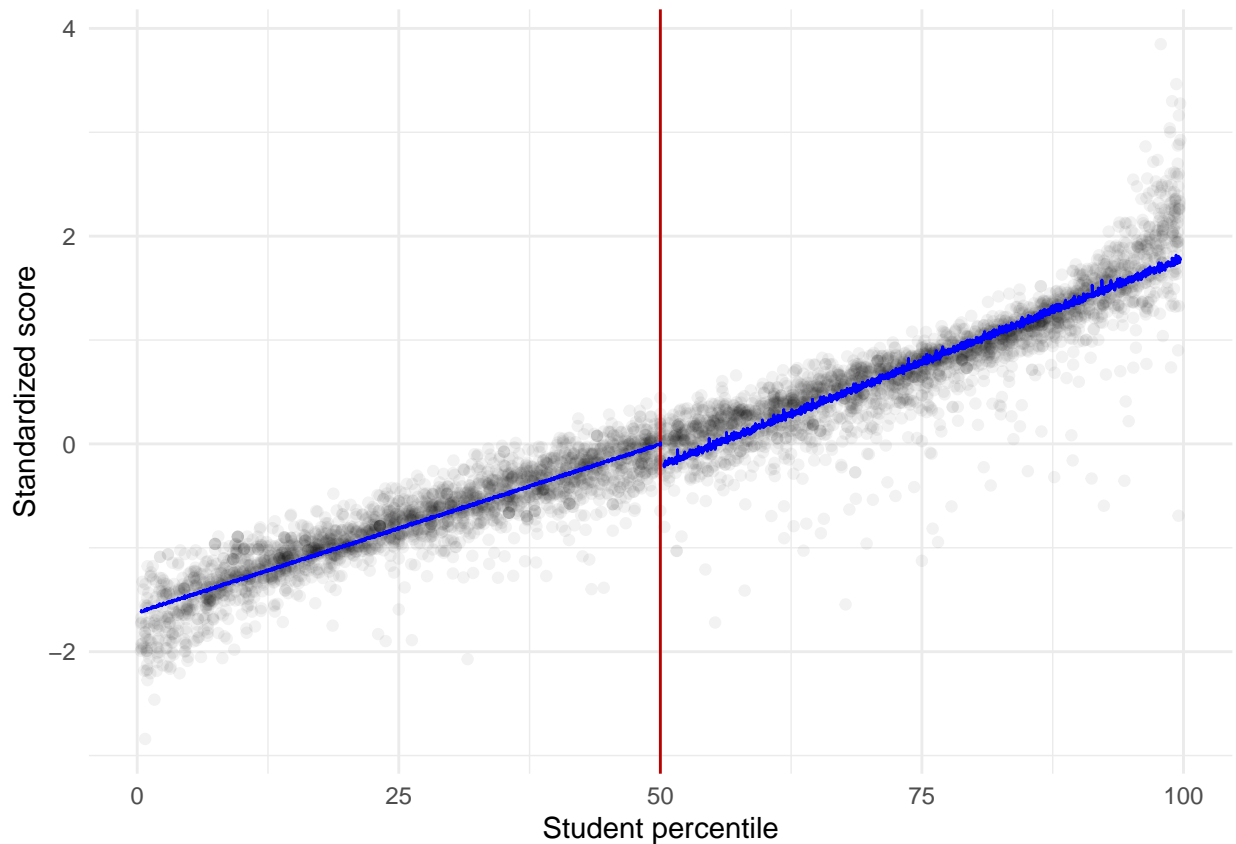
# colnames(group1)
# group 2
lm_ctrl2 <- lm(y2 ~ x2 + gender + age + teacher, data=group2)

y_ctrl_curve1 <- predict(lm_ctrl1, newdata = data.frame(x1=group1$x,
                                                         gender=group1$gender,
                                                         age=group1$age,
                                                         teacher=group1$teacher))

y_ctrl_curve2 <- predict(lm_ctrl2, newdata = data.frame(x2=group2$x,
                                                         gender=group2$gender,
                                                         age=group2$age,
                                                         teacher=group2$teacher))

ctrl_curve1 <- data.frame(x=group1$x, y=y_ctrl_curve1)
ctrl_curve2 <- data.frame(x=group2$x, y=y_ctrl_curve2)
```

```
ggplot(mapping=aes(x=x,y=y)) +
  geom_point(data=dat_use, alpha= 0.05) +
  geom_vline(aes(xintercept=50), colour="#BB0000") +
  geom_line(data = ctrl_curve1, color='blue') +
  geom_line(data = ctrl_curve2, color='blue') +
  xlab('Student percentile') +
  ylab('Standardized score') +
  theme_mp()
```



```
xbin1 <- cut(x1, breaks=quantile(x1,probs=seq(0,1,length.out=nbins+1)),
            labels=FALSE)
xbin1[is.na(xbin1)] <- 1

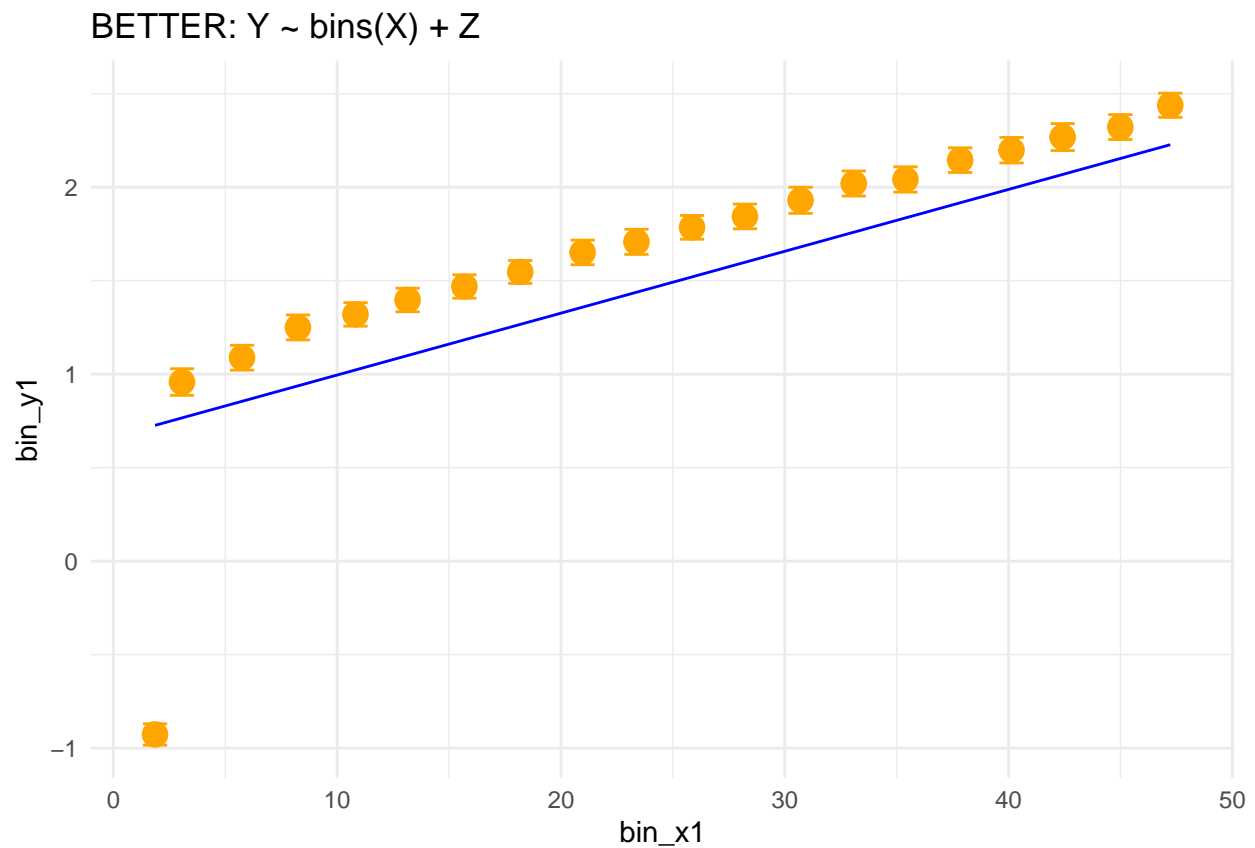
# xbin1

df_bin1 <- data.frame(y=y1, x1=x1,
                     w11=group1$gender,
                     w22=group1$age - mean(group1$age),
                     w33=group1$teacher,
                     xbin1=as.factor(xbin1))
reg_bins1 <- lm(y~1+xbin1+w11+w22+w33, data=df_bin1,x=TRUE)
reg_bins_xval1 <- lm(x1~1+xbin1, data=df_bin1)
df2_1 <- data.frame(bin_x1 = reg_bins_xval1$coefficients[1:nbins],
                   bin_y1 = reg_bins1$coefficients[1:nbins] - mean(y1),
                   bin_y_se1 = sqrt(diag(vcovHC(reg_bins1)))[1:nbins])
```

```

ggplot(data=df2_1, aes(x=bin_x1, y=bin_y1)) +
  geom_line(
    data =
      data.frame(
        x = seq(min(df2_1$bin_x1), max(df2_1$bin_x1), length.out=100),
        y = predict(lm_1, newdata = data.frame(
          x = seq(min(df2_1$bin_x1), max(df2_1$bin_x1), length.out=100)
        )) - 3*mean(y1)
      ),
    aes(x=x, y=y), color='blue') +
  # geom_line(data = ctrl_curve2, color='blue') +
  geom_errorbar(aes(ymin=bin_y1-1.96*bin_y_se1,ymax=bin_y1+1.96*bin_y_se1),
    color='orange') +
  geom_point(color='orange',size=4) +
  ggtitle('BETTER: Y ~ bins(X) + Z') +
  theme_mp()

```



```

saveplot(paste0(path_figure_save,"sim_residualize_better"))

```

## Reference

1. Duflo, E., Dupas, P., & Kremer, M. (2011). Peer effects, teacher incentives, and the impact of tracking: Evidence from a randomized evaluation in Kenya. *American economic review*, 101(5), 1739-74.