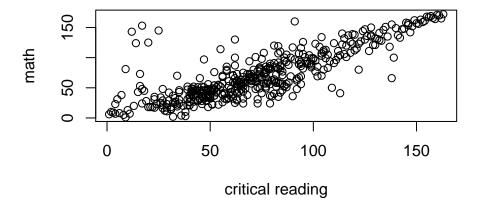
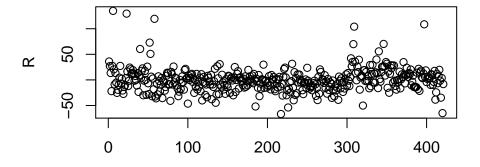
```
library(RCurl)
library(dplyr)
library(ggplot2)
library(lmtest)
x <- getURL("https://raw.github.com/cmm6/data605/main/2012_SAT_Results.csv",
            .opts=curlOptions(followlocation = TRUE))
sat_scores <- read.csv(text = x, header=TRUE)</pre>
colnames(sat_scores) <- c('dbn', 'school_name',</pre>
                           'num_test_takers',
                            'critical_reading',
                            'math','writing')
sat_scores$num_test_takers <- as.numeric(sat_scores$num_test_takers)</pre>
s_omitted <- sat_scores[- grep("s", sat_scores$math),]</pre>
x <- as.numeric(s_omitted$critical_reading)</pre>
y <- as.numeric(s_omitted$math)</pre>
math_as_cr <- lm(y ~ x)</pre>
summary(math_as_cr)
##
## Call:
## lm(formula = y \sim x)
##
## Residuals:
              1Q Median
##
       Min
                                3Q
                                        Max
## -66.620 -13.769 -1.266 10.843 134.850
##
## Coefficients:
              Estimate Std. Error t value Pr(>|t|)
##
## (Intercept) 2.30624 2.57867 0.894 0.372
                           0.03159 29.506 <2e-16 ***
## x
               0.93198
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 23.49 on 419 degrees of freedom
## Multiple R-squared: 0.6751, Adjusted R-squared: 0.6743
## F-statistic: 870.6 on 1 and 419 DF, p-value: < 2.2e-16
```

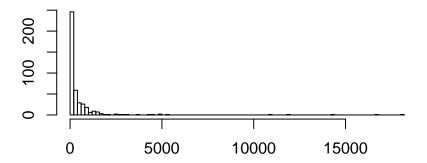


plot(math_as_cr\$residuals, xlab = "", ylab = "R")



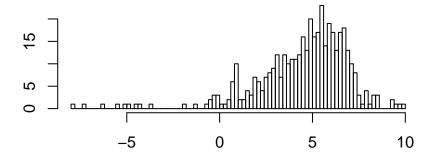
hist(math_as_cr\$residuals^2, breaks = 100, main = "log of R^2", xlab = "", ylab="")

log of R^2

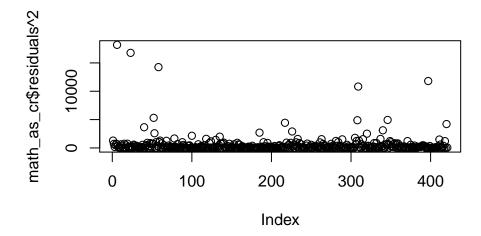


hist(log(math_as_cr\$residuals^2), breaks = 100, main = "log of R^2", xlab = "", ylab="")

log of R^2

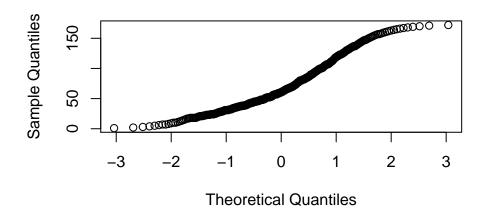


plot(math_as_cr\$residuals^2)



qqnorm(y)

Normal Q-Q Plot



```
normies <- s_omitted %>%
  mutate(r = math_as_cr$residuals) %>%
  filter(r^2 < 1200)

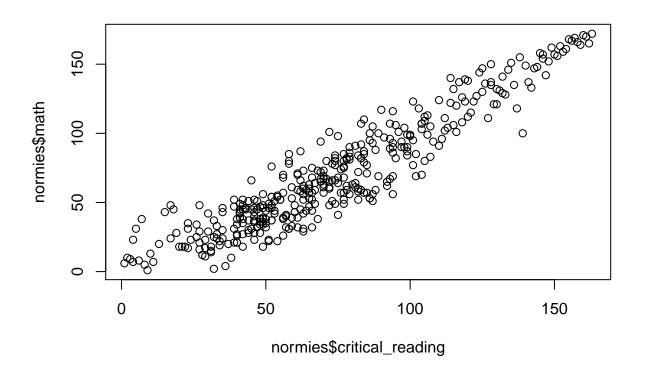
normies$math <- as.numeric(normies$math)
normies$critical_reading <- as.numeric(normies$critical_reading)
normies$writing <- as.numeric(normies$writing)

y <- normies$math
x <- normies$critical_reading

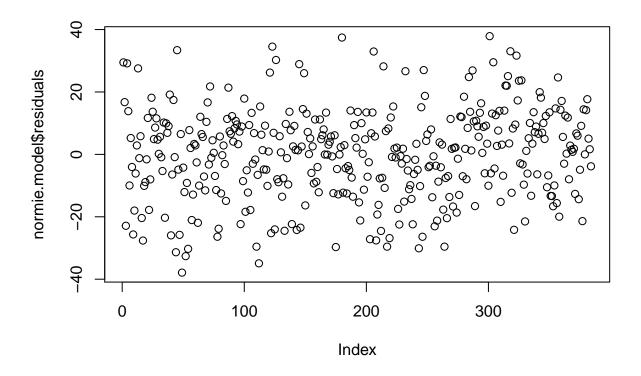
normie.model <- lm(y ~ x)
summary(normie.model)</pre>
```

```
##
## Call:
## lm(formula = y \sim x)
##
## Residuals:
##
       Min
                                3Q
                1Q Median
                                       Max
   -37.890 -9.741
                     1.320
                             9.256 37.880
##
## Coefficients:
##
              Estimate Std. Error t value Pr(>|t|)
## (Intercept) -7.18652
                           1.66723
                                     -4.31 2.08e-05 ***
                1.04372
                           0.02026
                                     51.52 < 2e-16 ***
## x
## ---
## Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
## Residual standard error: 14.47 on 382 degrees of freedom
## Multiple R-squared: 0.8742, Adjusted R-squared: 0.8738
## F-statistic: 2654 on 1 and 382 DF, p-value: < 2.2e-16
```

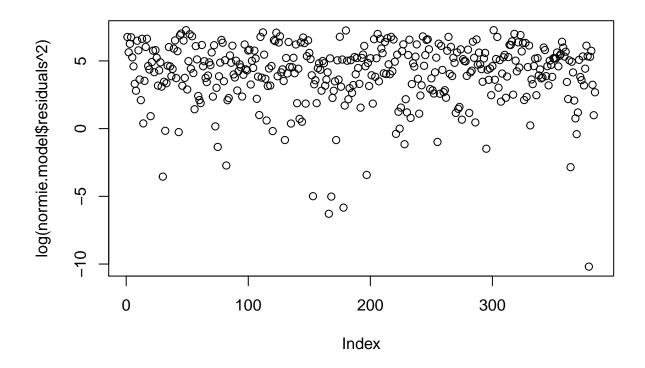
plot(normies\$critical_reading, normies\$math)



plot(normie.model\$residuals)

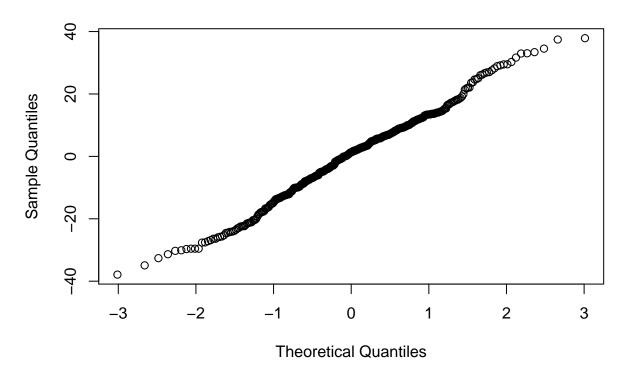


plot(log(normie.model\$residuals^2))



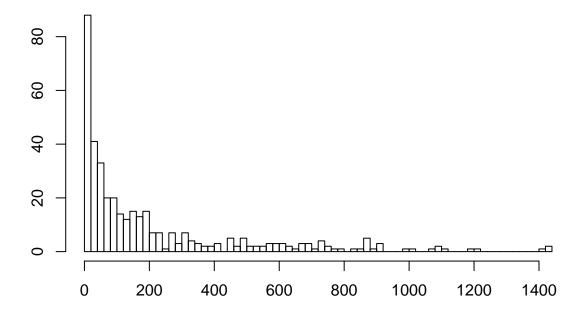
qqnorm(normie.model\$residuals)

Normal Q-Q Plot



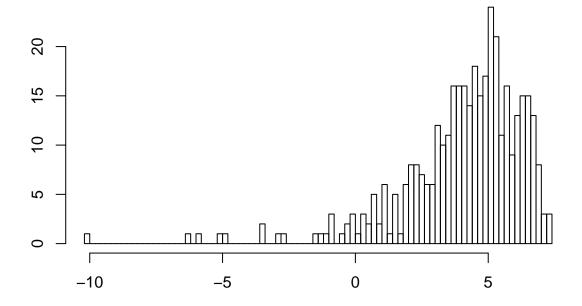
hist(normie.model\$residuals^2, breaks = 100, main = "log of R^2", xlab = "", ylab="")

log of R^2



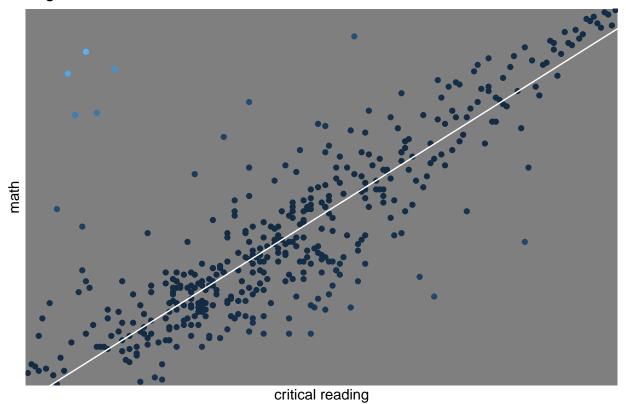
hist(log(normie.model\$residuals^2), breaks = 100, main = "log of R^2", xlab = "", ylab="")

log of R^2



```
math_as_cr$math <- as.numeric(math_as_cr$math)</pre>
math_as_cr$critical_reading <- as.numeric(math_as_cr$critical_reading)</pre>
math_as_cr$writing <- as.numeric(math_as_cr$writing)</pre>
ggplot(s_omitted, aes(x = critical_reading,
                    y = math,
                    color = math_as_cr$residuals^2)) +
  geom_point() +
  geom_abline(intercept = normie.model$coefficients[1],
              slope = normie.model$coefficients[2],
              color = "white") +
  theme_dark() +
  labs(title = "original data set with 'Normie Line'",
       x = "critical reading",
       y = "math") +
  theme(legend.position = "none") +
  scale_x_discrete(breaks=NULL) +
  scale_y_discrete(breaks=NULL)
```

original data set with 'Normie Line'



bptest(normie.model)

```
##
## studentized Breusch-Pagan test
##
## data: normie.model
## BP = 0.68361, df = 1, p-value = 0.4083
```

It's still not very good!