# Lying with Statistics Project

## Contents

1	Data manipulation	2
2	A fair figure depicting data	4
3	Conducting a statistical test such as t-squared test on girls vs boys	5
4	Compare differences across genders by creating a t-test for genders function	5
5	Display statistical analysis results of gender variances across score differences	7
6	Interpretation of results	8
7	Figure and analysis that provides a distorted version of what we actually would find in the data.	8
Q	Figure that significantly simplies the data	c



Diploma

PROOF:)

## 1 Data manipulation

1.1 Cleans up the RMD output and files by ensuring they don't fall off the page

1.2 Install and load all required packages (Don't install in rmd)

```
# install.packages('dplyr') install.packages('tidyverse')
# install.packages('knitr') install.packages('readr')
# install.packages('finalfit')

library("tidyverse")
library("dplyr")
library("knitr")
library("readr")
```

1.3 Read in files based on type, headers, and with specific NA values accounted for

1.4 Add transposed variable names to w4\_child based off of the provided text file

```
names(w4_child) <- t(names_w4_child)</pre>
```

1.5 Make all headers into lower case for easier merging

```
names(w1_child) <- tolower(names(w1_child))
names(w2_child) <- tolower(names(w2_child))
names(w3_child) <- tolower(names(w3_child))
names(w4_child) <- tolower(names(w4_child))
names(educinc) <- tolower(names(educinc))</pre>
```

1.6 Merge files by famid and select specified variables

```
w1234 <- (list(w1_child, w2_child, w3_child, w4_child, educinc) %>%
    reduce(full_join, by = "famid")) %>%
    dplyr::select(famid, c01cohort, c01gender, c01school, c01sibli,
        contains("atts"), contains("pcmp"), contains("attt"),
        contains("dscr"), contains("atod"), fameduc, income,
        c01sibli, contains("edex"))
```

1.7 Reverse code for pcmp 1 and 2

1.8 Compute averages based on sets of columns for variable sets and place average into a new variable

```
w1234$c01attt <- rowMeans(w1234[c(grep("c01attt", names(w1234)))],
    na.rm = TRUE)
w1234$c04attt <- rowMeans(w1234[c(grep("c04attt", names(w1234)))],
    na.rm = TRUE)
w1234$c01pcmp <- rowMeans(w1234[c(grep("c01pcmp", names(w1234)))],
    na.rm = TRUE)
w1234$c04pcmp <- rowMeans(w1234[c(grep("c04pcmp", names(w1234)))],
    na.rm = TRUE)
w1234$c01dscr <- rowMeans(w1234[c("c01dscr07", "c01dscr08", "c01dscr09",
    "c01dscr10")], na.rm = TRUE)
w1234$c04dscr <- rowMeans(w1234[c("c04dscr07", "c04dscr08", "c04dscr09",</pre>
```

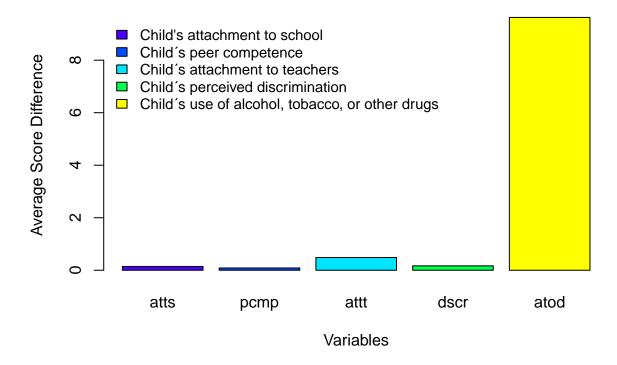
### 1.9 Create difference scores between Waves 1 and 4

### 1.10 Function to generate a random vector of colors in order from

```
color_vector <- function(num) {
   return(topo.colors(num))
}</pre>
```

## 2 A fair figure depicting data

# Average Score Differences in Variables from Wave 1 to Wave 4



- 3 Conducting a statistical test such as t-squared test on girls vs boys
- 3.1 Recode girls and boys

```
w1234\$c01gender[w1234\$c01gender == 1] = "girl"
w1234\$c01gender[w1234\$c01gender == 2] = "boy"
```

4 Compare differences across genders by creating a t-test for genders function

```
t_test_gender <- function(score_var) {
    return(t.test(score_var ~ c01gender, paired = FALSE, data = w1234))</pre>
```

```
}
tatod <- t_test_gender(w1234$difference_atod)</pre>
tattt <- t_test_gender(w1234$difference_attt)</pre>
tatts <- t_test_gender(w1234$difference_atts)</pre>
tdscr <- t_test_gender(w1234$difference_dscr)</pre>
tpcmp <- t_test_gender(w1234$difference_pcmp)</pre>
tatod
##
## Welch Two Sample t-test
##
## data: score_var by c01gender
## t = -0.122, df = 669, p-value = 0.9
## alternative hypothesis: true difference in means between group boy and group girl is not equal to 0
## 95 percent confidence interval:
## -0.87090 0.76865
## sample estimates:
## mean in group boy mean in group girl
              -9.6409
                                 -9.5898
##
tattt
##
## Welch Two Sample t-test
##
## data: score_var by c01gender
## t = 1.32, df = 581, p-value = 0.19
## alternative hypothesis: true difference in means between group boy and group girl is not equal to 0
## 95 percent confidence interval:
## -0.043818 0.221895
## sample estimates:
## mean in group boy mean in group girl
##
              0.52987
                                 0.44083
tatts
##
## Welch Two Sample t-test
## data: score_var by c01gender
## t = 0.055, df = 666, p-value = 0.96
## alternative hypothesis: true difference in means between group boy and group girl is not equal to 0
## 95 percent confidence interval:
## -0.087517 0.092561
## sample estimates:
## mean in group boy mean in group girl
              0.14484
                                 0.14232
```

```
tdscr
##
   Welch Two Sample t-test
##
##
## data: score_var by c01gender
## t = 0.0432, df = 572, p-value = 0.97
## alternative hypothesis: true difference in means between group boy and group girl is not equal to 0
## 95 percent confidence interval:
## -0.077353 0.080834
## sample estimates:
## mean in group boy mean in group girl
##
              0.16956
                                0.16782
tpcmp
##
## Welch Two Sample t-test
##
## data: score_var by c01gender
## t = 1.19, df = 579, p-value = 0.23
## alternative hypothesis: true difference in means between group boy and group girl is not equal to 0
## 95 percent confidence interval:
## -0.031859 0.130578
## sample estimates:
## mean in group boy mean in group girl
##
            0.111693
                               0.062333
```

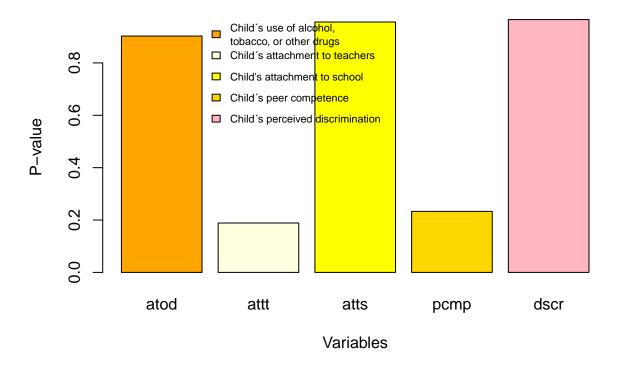
# 5 Display statistical analysis results of gender variances across score differences

### 5.1 Function to extract p values from t-test result text

```
extract_pval <- function(ttest) {
    return(ttest$p.value)
}</pre>
```

### 5.2 Graph p values

### Gender Differences in Score Differences from Wave 1 to Wave 4



## 6 Interpretation of results

- 6.1 S
- 7 Figure and analysis that provides a distorted version of what we actually would find in the data.
- 7.1 Code to determine which p\_values are stastically significant i.e., <0.05

```
p_values[p_values < 0.05] = "yes"
p_values[p_values > 0.05] = "ABSOLUTELY NO CHANGE"
percent_not_significant <- c(length(p_values[p_values > 0.05])/5 *
100, length(p_values[p_values < 0.05])/5 * 100)</pre>
```

## 8 Figure that significantly simplies the data

```
pie(percent_not_significant, col = c("red", "blue"), labels = percent_not_significant,
    main = "% of Score Difference Variables that Statistically Differed by Gender")

legend(-2.1, 1.05, c("STATISTICALLY NO DIFFERENCE", "yes"), cex = 0.8,
    fill = c("red", "blue"))
```

## % of Score Difference Variables that Statistically Differed by Gende

