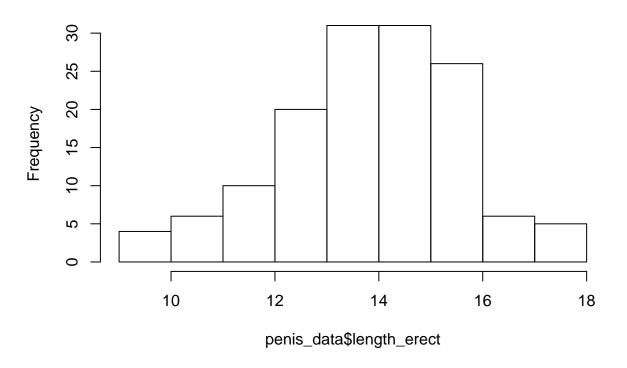
#### Penis Measurements Across the World

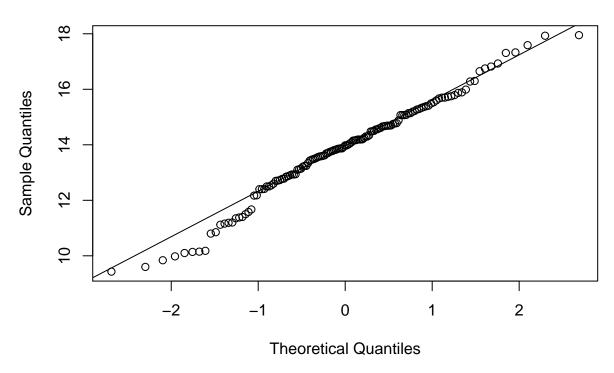
```
library(ggplot2)
library(tidyverse)
## -- Attaching packages
## v tibble 2.1.3
                      v dplyr
           1.0.2
## v tidyr
                      v stringr 1.4.0
           1.3.1
                      v forcats 0.4.0
## v readr
## v purrr
            0.3.3
## -- Conflicts -----
## x dplyr::filter() masks stats::filter()
## x dplyr::lag()
                    masks stats::lag()
# read in dataset
penis_data <- read.csv("/Users/Dohyun/Desktop/projects/Penis-Project/world_penis_dataset/penis.csv")</pre>
#check normality of erect length means
#using a histogram
hist(penis_data$length_erect)
```

## Histogram of penis\_data\$length\_erect



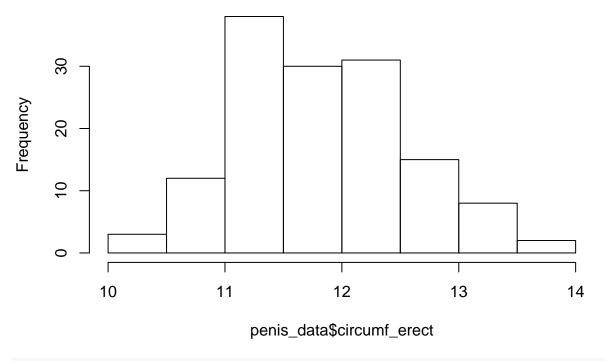
```
#NPP plot
qqnorm(penis_data$length_erect)
qqline(penis_data$length_erect)
```

# Normal Q-Q Plot



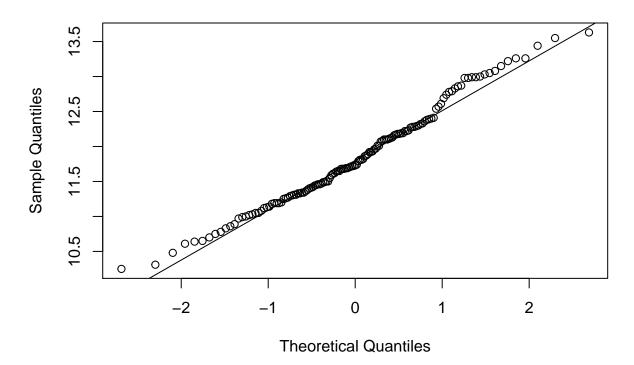
#check normality of erect girth means
#using a histogram
hist(penis\_data\$circumf\_erect)

## Histogram of penis\_data\$circumf\_erect



```
#NPP plot
qqnorm(penis_data$circumf_erect)
qqline(penis_data$circumf_erect)
```

### Normal Q-Q Plot



```
t.test(penis_data$length_erect)
##
##
    One Sample t-test
## data: penis_data$length_erect
## t = 91.633, df = 138, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 13.55726 14.15526
## sample estimates:
## mean of x
## 13.85626
t.test(penis_data$circumf_erect)
##
##
    One Sample t-test
##
## data: penis_data$circumf_erect
## t = 192.43, df = 138, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 11.71941 11.96275
## sample estimates:
## mean of x
## 11.84108
Confidence interval for mean erect length is 13.56-14.16 cm. Confidence interval for mean erect girth is
11.72-11.96 cm.
#check for overlaps between both methods
self_reported_data <- filter(penis_data, Method == "Self reported")</pre>
measured_data <- filter(penis_data, Method == "Measured")</pre>
t.test(self_reported_data$length_erect)
##
##
   One Sample t-test
## data: self_reported_data$length_erect
## t = 79.429, df = 50, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 14.33515 15.07897
## sample estimates:
## mean of x
## 14.70706
```

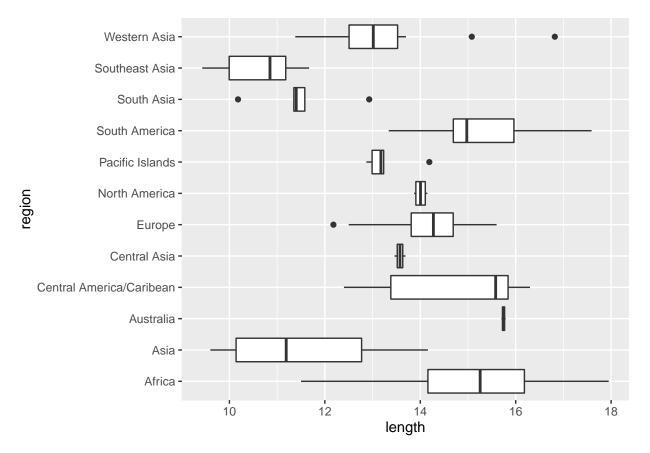
```
#CI for mean self-reported length: 14.33-15.08
t.test(measured_data$length_erect)
##
##
   One Sample t-test
##
## data: measured_data$length_erect
## t = 68.323, df = 87, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 12.97443 13.75193
## sample estimates:
## mean of x
## 13.36318
#CI for mean measured length: 12.97-13.75
t.test(self_reported_data$circumf_erect)
##
##
   One Sample t-test
##
## data: self_reported_data$circumf_erect
## t = 139.45, df = 50, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 11.90118 12.24902
## sample estimates:
## mean of x
     12.0751
##
#CI for mean self-reported girth: 11.55-11.86
t.test(measured_data$circumf_erect)
##
##
   One Sample t-test
## data: measured_data$circumf_erect
## t = 146.16, df = 87, p-value < 2.2e-16
## alternative hypothesis: true mean is not equal to 0
## 95 percent confidence interval:
## 11.54628 11.86463
## sample estimates:
## mean of x
## 11.70545
##CI for mean measure length: 11.90-12.25
```

Note that we we only care about the erect length and girth it provides a better standard of measurement. Flaccid measurements will always vary depending on body and outside temperature and different conditions like health.

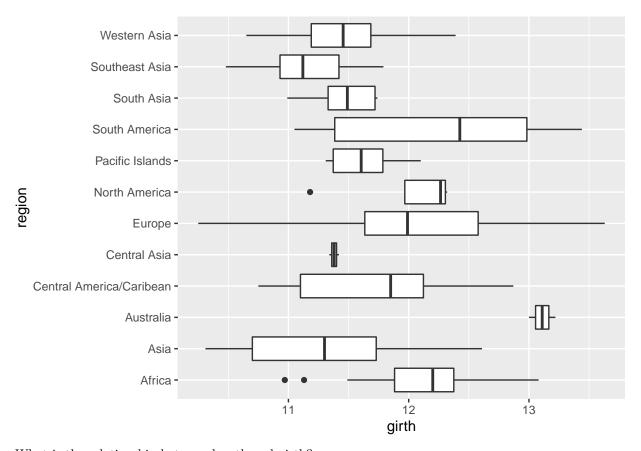
The confidence intervals for the self-reported and measured don't overlap, and on average the self-reported data shows a higher range of measurements. It is not certain whether the self-reported measurements are honest, but it is not out of the question whether they are.

```
size_length <- length(penis_data$length_erect)
size_girth <- length(penis_data$circumf_erect)
region <- penis_data[,"Region"]
length <- penis_data[,"length_erect"]
girth <- penis_data[,"circumf_erect"]

#boxplot of the regions
bp <- ggplot(penis_data, aes(x = region, y = length)) +
    geom_boxplot()
bp + coord_flip()</pre>
```



```
#boxplot of the regions vs girth
bp2 <- ggplot(penis_data, aes(x = region, y = girth)) +
   geom_boxplot()
bp2 + coord_flip()</pre>
```



What is the relationship between length and girth?

```
reg_data <- data.frame(length, girth)
fit1 <- lm(girth ~ length, reg_data)
plot(fit1)</pre>
```

