### 4211 Homework 9

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1

(a)

```
x = data.frame(
    msce = c(8.27,8.20,8.25,8.14,9.00,8.10,7.20,8.32,7.70,8.50,9.48,8.65,8.16,8.83,7.76,8.63),
    eth = c(rep(0,9), rep(1,7))
)

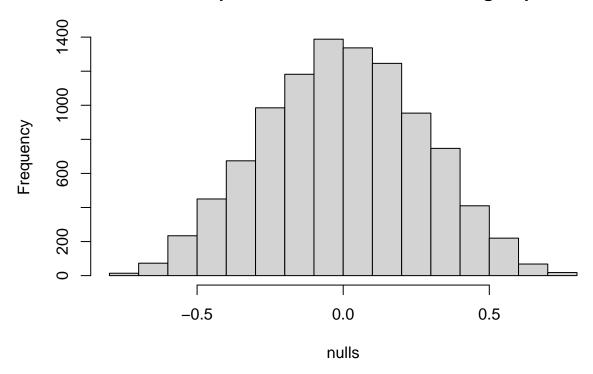
permtest = function(x,y){
    xstar = sample(x)
    return(mean(y[xstar==1])-mean(y[xstar==0]))
}

set.seed(488103)
observeddiff = mean(x$msce[x$eth==1]-x$msce[x$eth==0])

## Warning in x$msce[x$eth == 1] - x$msce[x$eth == 0]: longer object length is not
## a multiple of shorter object length

nulls = replicate(10000, permtest(x$eth, x$msce))
hist(nulls, main = "Distribution of permuted difference between group means")
```

# Distribution of permuted difference between group means



(b)

mean(abs(nulls)>abs(observeddiff))

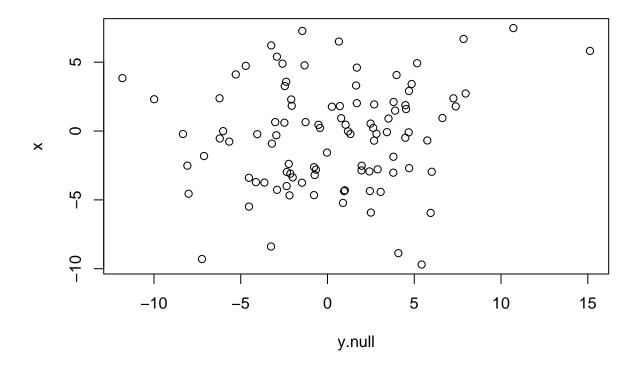
## [1] 0.0442

The difference is significant at the level alpha = 0.05.

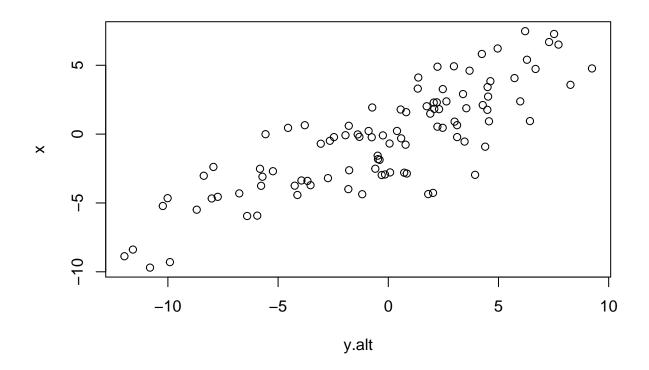
```
\mathbf{2}
```

(a)

```
set.seed(4211)
x=rnorm(100,0,4)
y.null=rnorm(100,0,5) # null hypothesis
y.alt=x+rnorm(100,0,3) # alternative hypothesis
## from a location model with normal errors
plot(y.null, x)
```



plot(y.alt, x)



#### cor.test(x, y.null)

Pearson's product-moment correlation

##

```
##
## data: x and y.null
## t = 1.5235, df = 98, p-value = 0.1309
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## -0.04567892  0.33841165
## sample estimates:
## cor
## 0.1521039

cor.test(x, y.alt)
```

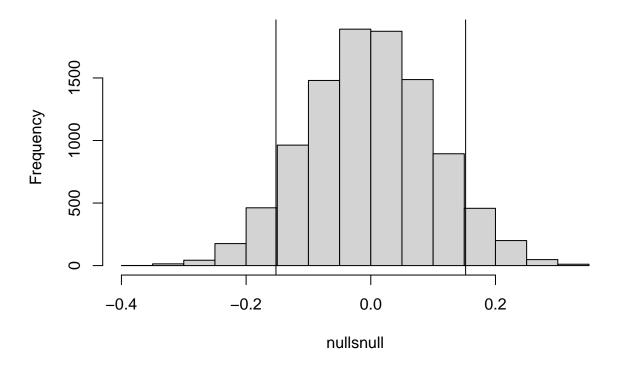
```
##
## Pearson's product-moment correlation
##
## data: x and y.alt
## t = 14.757, df = 98, p-value < 2.2e-16
## alternative hypothesis: true correlation is not equal to 0
## 95 percent confidence interval:
## 0.7576060 0.8828581</pre>
```

```
## sample estimates:
## cor
## 0.8304473
(b)
```

```
permtest = function(x,y){
    xstar = sample(x)
    return(cor(xstar, y))
}

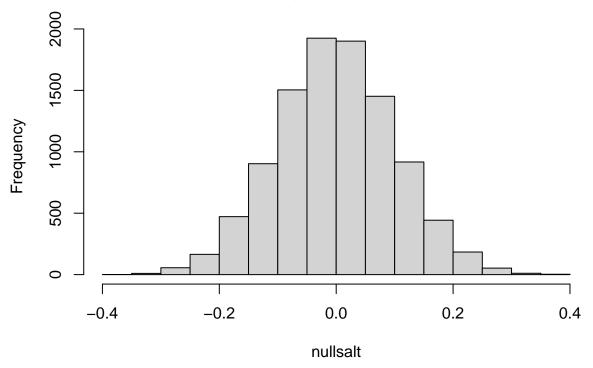
set.seed(488103)
    observedcornull = cor(x, y.null)
    observedcoralt = cor(x, y.alt)
    nullsnull = replicate(10000, permtest(x, y.null))
    nullsalt = replicate(10000, permtest(x, y.alt))
    hist(nullsnull)
    abline(v=observedcornull)
abline(v=-observedcornull)
```

## Histogram of nullsnull



hist(nullsalt)

# Histogram of nullsalt



mean(abs(nullsnull)>abs(observedcornull))

## [1] 0.1362

mean(abs(nullsalt)>abs(observedcoralt))

## [1] 0

3

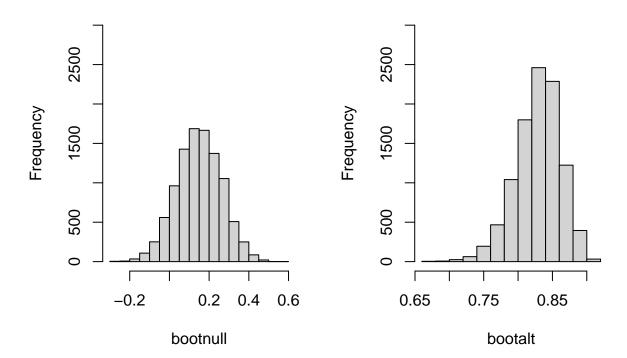
(a)

hist(bootalt, ylim=c(0,3000))

```
bootstrap = function(x,y){
  index = sample(1:length(x), replace = TRUE)
  xstar = x[index]
 ystar = y[index]
 return(cor(xstar, ystar))
}
set.seed(488103)
bootnull = replicate(10000, bootstrap(x, y.null))
bootalt = replicate(10000, bootstrap(x, y.alt))
var(bootnull)
## [1] 0.01283102
var(bootalt)
## [1] 0.001072049
(b)
par(mfrow=c(1,2))
hist(bootnull, ylim=c(0,3000))
```

# Histogram of bootnull

# Histogram of bootalt



The distribution of correlation under the null hypothesis appears to be normally distributed about 1.5, while the distribution of correlation under the alternative hypothesis is skewed left, with a peak around 0.83.