## **Computer Practical 1**

### Q1 and Q2

```
n <- 13 p <-0.31 # we will carry out 100 trials in which each trial we observe 10 Binomial(n,p) sampleSize <- 10 trials <- 100 # varHat stores the true variance of the MLE of p varHat <- (1/n) * p * (1-p) print(paste("Variance of pHat(Y) is " ,varHat))
```

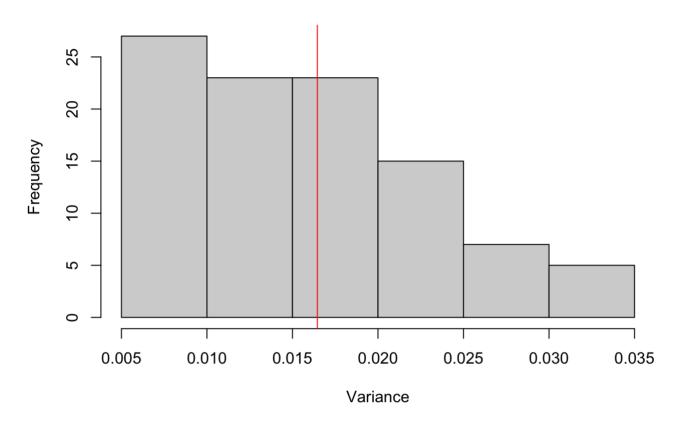
```
## [1] "Variance of pHat(Y) is 0.0164538461538462"
```

```
# phat is the MLE of p function
phat <-function(Y) {
    Y/n
}

# we generate 1000 binomials and organise them into 10 rows of 100
# this means each column is a trial
xValues <- rbinom(sampleSize*trials, n, p)
xSamples <- matrix(xValues, nrow=sampleSize) # 10x100
# we pass each row into phat to get the MLE function transformation
pHatSamples<-apply(xSamples, 1, phat) # it becomes 100x10
# we then get the sample variance of each row
pHatVar <- apply(pHatSamples, 1, var) # stays 100x1
hist(pHatVar, xlab = "Variance")
abline(v = varHat, col='red')</pre>
```

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#### Histogram of pHatVar



## Q3 (Q4 and Q5 done in working out section)

year.data <- read.csv("year\_data.csv")
knitr::kable(year.data)</pre>

year	births	deaths	clinic
1841	3036	237	1
1842	3287	518	1
1843	3060	274	1
1844	3157	260	1
1845	3492	241	1
1846	4010	459	1
1841	2442	86	2
1842	2659	202	2
1843	2739	164	2
1844	2956	68	2
1845	3241	66	2
1846	3754	105	2

```
n1 <- sum(year.data[year.data$clinic==1,]$births)
y1 <- sum(year.data[year.data$clinic==1,]$deaths)
n2 <- sum(year.data[year.data$clinic==2,]$births)
y2 <- sum(year.data[year.data$clinic==2,]$deaths)
print(paste("The number of births in clinic 1 is ", n1))</pre>
```

## [1] "The number of births in clinic 1 is 20042"

```
print(paste("The number of deaths in clinic 1 is ", y1))
```

```
## [1] "The number of deaths in clinic 1 is 1989"
```

```
print(paste("The number of births in clinic 2 is", n2))
```

```
## [1] "The number of births in clinic 2 is 17791"
```

```
print(paste("The number of deaths in clinic 2 is", y2))
```

```
## [1] "The number of deaths in clinic 2 is 691"
```

```
print(paste("The ML estimates of p1hat(y1) is y1/n1 which is 1989/20042 which is",y1/n1))
```

## [1] "The ML estimates of plhat(y1) is y1/n1 which is 1989/20042 which is 0.0992415 926554236"

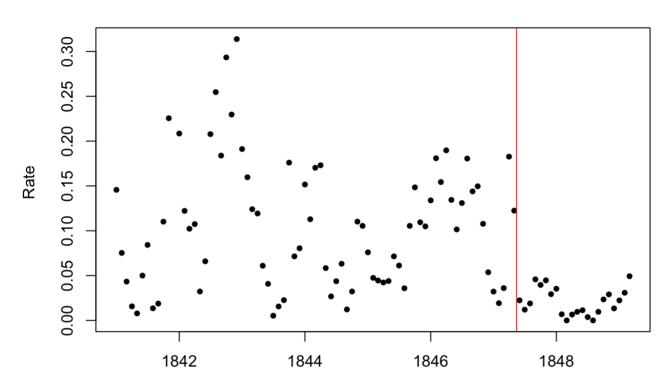
```
print(paste("The ML estimates of p2hat(y2) is y2/n2 which is 691/17791 which is " , y 2/n2))
```

## [1] "The ML estimates of p2hat(y2) is y2/n2 which is 691/17791 which is 0.0388398 628520038"

# Q6 First Part (second part done in working out section)

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#### Mortality rate by month



Date; red line indicated start of intervention period

```
before.interventions <- month.data[month.data$date < intervention.date,]
after.intervention <- month.data[month.data$date > intervention.date,]

n1 <- sum(before.interventions$births)
y1 <- sum(before.interventions$deaths)
n2 <- sum(after.intervention$births)
y2 <- sum(after.intervention$deaths)
print(paste("The number of births before intervention is ", n1))</pre>
```

```
## [1] "The number of births before intervention is 19571"
```

```
print(paste("The number of deaths before intervention is ", y1))
```

## [1] "The number of deaths before intervention is 2060"

```
print(paste("The number of births after intervention is", n2))
```

## [1] "The number of births after intervention is 6595"

```
print(paste("The number of deaths after intervention is", y2))
```

## [1] "The number of deaths after intervention is 142"

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print(paste("The ML estimates for the mortality rates before intervention is y1/n1 wh ich is " , y1/n1))

## [1] "The ML estimates for the mortality rates before intervention is y1/n1 which i s 0.105257779367431"

print(paste("The mortality rates after intervention is y2/n2 which is ", y2/n2))

## [1] "The mortality rates after intervention is y2/n2 which is 0.0215314632297195"

Q4) 
$$IE[w] = IE[\rho, (Y_1) - \rho_2(Y_2)]$$
  
 $= [E[\rho, (Y_1)] - [E[\rho_2(Y_2)]]$   
 $= \rho - \rho = 0$   
 $Var(w) = Var(\rho, (Y_1) - \rho_2(Y_2))$   
 $= Var(\rho, [Y_1)) + Var(\rho_2(Y_2))$   
 $= \frac{1}{n_1} \rho_1(1-\rho_1) + \frac{1}{n_2} \rho_2(1-\rho_2)$ 

$$=\left(\frac{1}{n_1}+\frac{1}{n_2}\right)P(1-P)$$

QS)

We are assuming the true underlying mortallity rates are the same so let
$$p_1 = p_2 = p = 1989 + 691 = 2680$$

$$p_1 = p_2 = p = 1989 + 691 = 2680$$
  
 $20042 + 17791 = 37833$ 

= 0.67683763  
Then IP 
$$(|W - \mu_w| > \rho_i(y_i) - \rho_2(y_2))$$

$$= 1P(|w|) > 0.09924159 - 0.0388)$$

$$= 1P(|w|) > 0.06040173)$$

 $= \frac{(\frac{1}{n_1} + \frac{1}{n_2}) P(1-P)}{(0.06040)^2}$   $= \frac{(\frac{1}{n_1} + \frac{1}{n_2}) P(1-P)}{(0.06040)^2}$ 

Thos we have shown that there is a small probability we will see such a large difference between the observed mortallity rate under the assomption that the true underlying mortality rates are actually the same.

= 0-00/9/42

Q6) let P be true underlying mortallity rate then as before we assume

$$P = \frac{2060 + 142}{19571 + 6595} = \frac{2202}{26166}$$

= 0.0841550)

Similarly to before,  

$$(E(w) = 0)$$
  
 $(ar(w) \neq 1 + 1)$   
 $(ar(w) = 0)$   
 $(ar(w) \neq 1 + 1)$   
 $(ar(w) = 0)$   
 $(ar(w$ 

assoming underlying mortallity rore same for both pardom variables

Then 
$$P(1W - \mu_W | 7, \hat{p}_1(y_1) - \hat{p}_2(y_2))$$

$$= P(1W | 7, 0.052777 - 0.021531)$$

$$= P(1W | 7, 0.08374639)$$

$$\leq \left(\frac{1}{11} + \frac{1}{12}\right) P(1-P) \text{ Changeur Inequality}$$

$$= \left(\frac{1}{19571} + \frac{1}{6595}\right) (0.08415501)$$

$$= \left(\frac{1}{19571} + \frac{1}{6595}\right) (0.08415501)$$

Thus we have shown there is a Small probability we will see such a large difference between the observed morrally rate under the assemption the inderlying martally rates are the same.

Q7) 
$$L(0) \times \prod_{i=1}^{2} f_{Y_{i}}(y_{i}, n_{i}, x_{i}; 0)$$