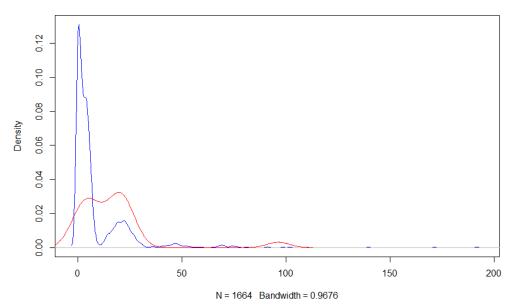
BACS HW (Week 5)

1. a) Divide customers into 2 groups and visualizing the ILEC & CLEC customers.

```
> time<-read.csv("verizon.csv",header = TRUE)
> ILEC<-time$Time[time$Group=="ILEC"]
> CLEC<-time$Time[time$Group=="CLEC"]
> plot(density(ILEC),col="blue",main="Response times")
> lines(density(CLEC),col="red")
```

Response times



b) Test the difference between the mean of ILEC and mean of CLEC by t.test function

- (i) In this case, H_0 : The mean of CLEC <=ILEC H_1 ::The mean of CLEC > ILEC
- (ii) Since p>0.01, cannot reject H_0 .

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c) Test the alternative and null values of t by bootstrapping compare and compare with t.test function.

```
> boot_mean_diff<-function(CLEC,ILEC){
  resample_CLEC<-sample(CLEC,length(CLEC),replace=TRUE)
  resample_ILEC<-sample(ILEC,length(ILEC),replace=TRUE)

  t_test_alt<- t.test(resample_CLEC,resample_ILEC, "greater",conf.level = 0.99)

  t_test_null<-t.test(resample_CLEC,CLEC)

  return(c(t_test_alt*statistic,t_test_null*statistic))
  }

  * t_test<-replicate(3000,boot_mean_diff(CLEC,ILEC))

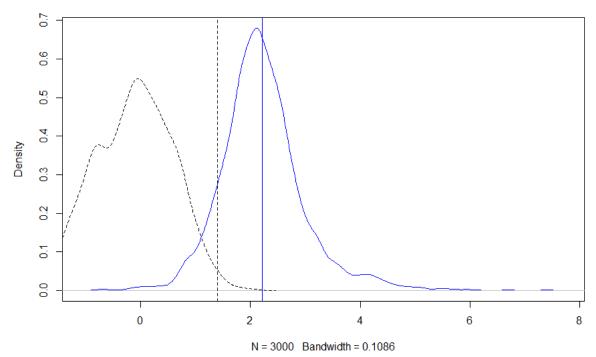
  * t_alt<-t_test[1,]

  * t_null<-t_test[2,]</pre>
```

(i) Plot the null hypothesis and alternative hypothesis, show the 1% rejection zone.

```
> plot(density(t_alt),col="blue",main="Bootstrapping the difference of mean")
> lines(density(t_null),lty="dashed")
> abline(v=quantile(t_null,c(0.99)),lty="dashed")
> abline(v=mean(t_alt),col="blue")
```

Bootstrapping the difference of mean



(ii) As the alternative t-value largely outside the 1% rejection zone, we could reject the null hypothesis.

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- 2. Test whether the variance of ILEC & CLEC response times are different.
 - a) The null & alternative hypothesis

H₀: There is no difference between the variance of ILEC & CLEC

H₁: The variance between ILEC & CLEC are not the same.

b)

```
#Identify which variance is higher
> var(CLEC)>var(ILEC)
[1] TRUE
```

(i) F-statistic of the ratio of variance=1.762717

(ii) The cut-off value of F=1.548476

```
> qf(p=0.95,df1=length(CLEC)-1,df2=length(ILEC)-1)
[1] 1.548476
```

- (iii) We can reject the null hypothesis since F-value is greater than the cut-off value.
- c) Bootstrapping

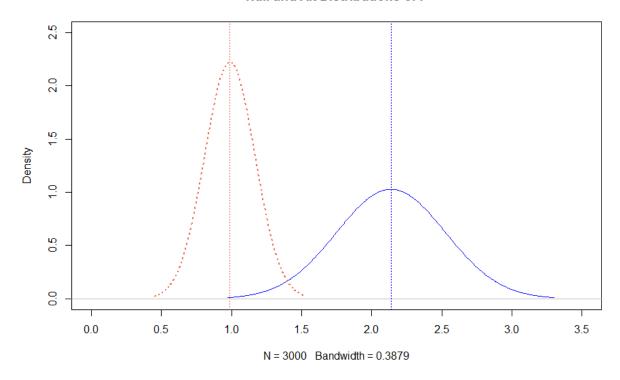
```
> boot_f_stat<-function(CLEC,ILEC) {
    resample_CLEC<-sample(CLEC,length(CLEC),replace=TRUE)
    resample_ILEC<-sample(ILEC,length(ILEC),replace=TRUE)
    f_alt<-var(resample_CLEC)/var(resample_ILEC)
    f_null<-var(resample_CLEC)/var(CLEC)
    c(f_alt,f_null)
}</pre>
```

(i) H_0 : The f-value <=1 H_1 :The f-value>1

(ii) The 95% cutoff value is 0.9890.

(iii) Visualize H₀, H₁ and the cut off value.

Null and Alt Distributions of F



(iv) We can reject the null hypothesis since F-value is greater than the cut-off value.