

```

1)
f=file.choose()
DriveTest=read.table(f,header = TRUE)
Sober=DriveTest[c(1:24,41:80),1]
Drug=DriveTest[c(25:40,81:119),1]

> t.test(Sober,Drug,var.equal = T)

```

Two Sample t-test

```

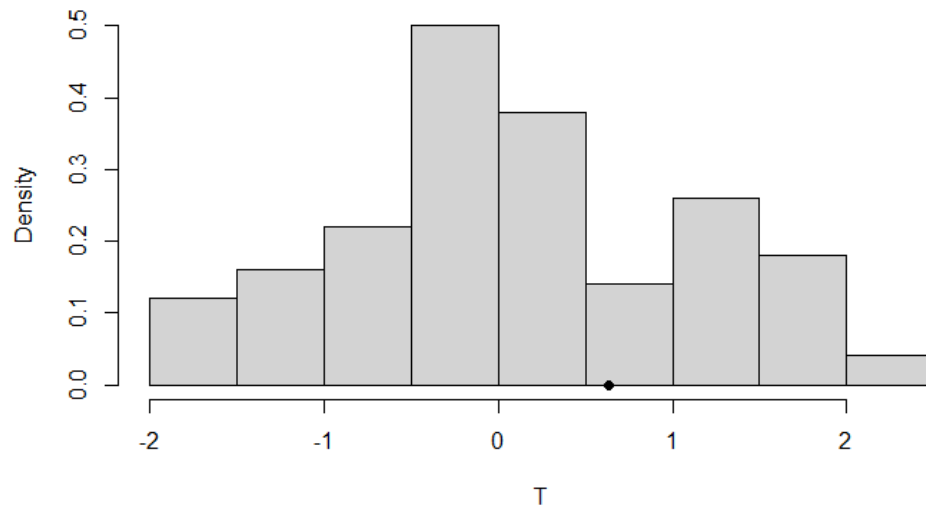
data: Sober and Drug
t = 0.6324, df = 117, p-value = 0.5284
alternative hypothesis: true difference in means is not equal to 0
95 percent confidence interval:
 -10.97359  21.26939
sample estimates:
mean of x mean of y
 43.3903  38.2424

```

```

> nm=DriveTest%>%filter(Drug=="1" | Drug=="0")%>%
+ arrange(Drug)%>%group_by(Drug)%>%summarise(n())%>%pull(2)
> z=DriveTest%>%filter(Drug=="1" | Drug=="0")%>%
+ arrange(Drug)%>%group_by(Drug)%>%pull(Distance)
> n=nm[1];m=nm[2];N=sum(nm)
> x=z[1:n];y=z[(n+1):N]
>
> t0 <- t.test(x, y)$statistic
> ts <- map(1:100,~sample.int(N,n))%>%map_dbl(~t.test(z[.x],z[-.x])$statistic)
> p <- mean(abs(ts) >= abs(t0))
> p
[1] 0.49
>
> hist(ts, main = "", freq = FALSE, xlab = "T",breaks = "scott")
> points(t0, 0, cex = 1, pch = 16)

```



```
> wilcox.test(Sober,Drug)
```

Wilcoxon rank sum test with continuity correction

data: Sober and Drug

W = 2184, p-value = 0.02399

alternative hypothesis: true location shift is not equal to 0

P-value is similar in the first two but not the third

2)

N = 100000

U = runif(N)

rand.samples = rep(NA,N)

```
for(i in 1:N){
```

```
  if(U[i]<.99){
```

```
    rand.samples[i] = rnorm(1,0,5)
```

```
  }else if(U[i]<.01){
```

```
    rand.samples[i] = rnorm(1,0,25)
```

```
  }
```

```
}
```

4)

```
library(car)
```

```
data("Ornstein")
```

Ornstein

```
p_glm=glm(interlocks~assets,family = poisson,data = Ornstein)
```

```
summary(p_glm)
```

Call:

```
glm(formula = interlocks ~ assets, family = poisson, data = Ornstein)
```

Deviance Residuals:

Min	1Q	Median	3Q	Max
-5.801	-3.397	-1.017	1.355	10.537

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)
(Intercept)	2.412e+00	1.916e-02	125.90	<2e-16 ***
assets	1.772e-05	4.113e-07	43.09	<2e-16 ***

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for poisson family taken to be 1)

Null deviance: 3737.0 on 247 degrees of freedom
Residual deviance: 2626.2 on 246 degrees of freedom
AIC: 3528.2

Number of Fisher Scoring iterations: 5

>