

Wilcoxon and Quantiles

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Wilcoxon Test with Ties

- ▶ Will show you warning message !

```
wilcox.test(c(-2, -2, 2, 2, 5, 5, 5), mu = 3)
```

```
## Warning in wilcox.test.default(c(-2, -2, 2, 2, 5, 5, 5),
```

```
## compute exact p-value with ties
```

```
##
```

```
## Wilcoxon signed rank test with continuity correction
```

```
##
```

```
## data: c(-2, -2, 2, 2, 5, 5, 5)
```

```
## V = 12, p-value = 0.7977
```

```
## alternative hypothesis: true location is not equal to 3
```

Suppress Warnings !

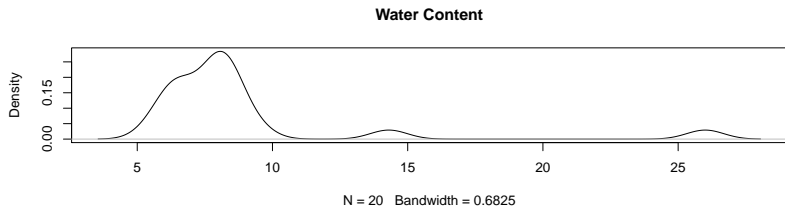
- ▶ We can suppress this warning message, but it's not recommended in practice

```
suppressWarnings(wilcox.test(c(-2,-2,2,2,5,5,5),mu = 3))
```

```
##  
##  Wilcoxon signed rank test with continuity correction  
##  
## data:  c(-2, -2, 2, 2, 5, 5, 5)  
## V = 12, p-value = 0.7977  
## alternative hypothesis: true location is not equal to 3
```

Another Example

```
x2 <- c(5.6, 6.1, 6.3, 6.4, 6.5, 6.6, 7.0, 7.5, 7.9, 8.0, 8.4)
## Plot the density
plot(density(x2), main = "Water Content")
```



Wilcoxon Signed Rank Test

```
suppressWarnings(wilcox.test(x2, mu=9, conf.int=TRUE))
```

```
##  
##  Wilcoxon signed rank test with continuity correction  
##  
## data:  x2  
## V = 41, p-value = 0.01774  
## alternative hypothesis: true location is not equal to 9  
## 95 percent confidence interval:  
##  7.150075 8.500071  
## sample estimates:  
## (pseudo)median  
##      7.810093
```

Sign Test

```
binom.test(sum(x2>9),length(x2),alternative = "two.sided")
```

```
##
```

```
## Exact binomial test
```

```
##
```

```
## data:  sum(x2 > 9) and length(x2)
```

```
## number of successes = 3, number of trials = 20, p-value
```

```
## alternative hypothesis: true probability of success is not equal to 0.5
```

```
## 95 percent confidence interval:
```

```
##  0.03207094 0.37892683
```

```
## sample estimates:
```

```
## probability of success
```

```
##                0.15
```

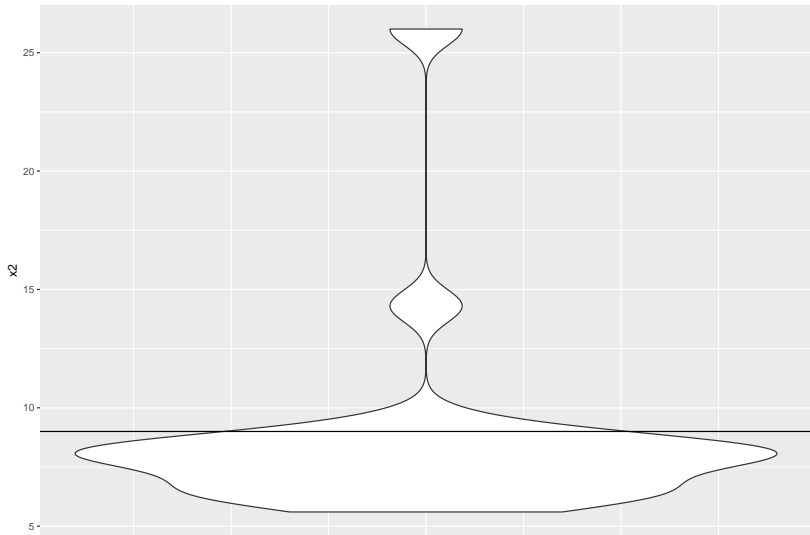
Parametric t-test

```
t.test(x2,alternative = "two.sided",mu=9)
```

```
##  
## One Sample t-test  
##  
## data: x2  
## t = -0.22154, df = 19, p-value = 0.827  
## alternative hypothesis: true mean is not equal to 9  
## 95 percent confidence interval:  
## 6.701535 10.858465  
## sample estimates:  
## mean of x  
## 8.78
```

Make a nice looking plot

```
library(ggplot2)
qplot(y=x2, x= 1, geom = "violin")+geom_abline(slope=0,inte
```



Quantiles

The `ecdf()` function in R

Description

Compute an empirical cumulative distribution function, with several methods for plotting, printing and computing with such an “ecdf” object.

Usage

Arguments

`x`, object

numeric vector of the observations for `ecdf`; for the methods, an object inheriting from class “ecdf”.

...

arguments to be passed to subsequent methods, e.g., `plot.stepfun` for the plot method.

`ylab`

label for the y-axis.

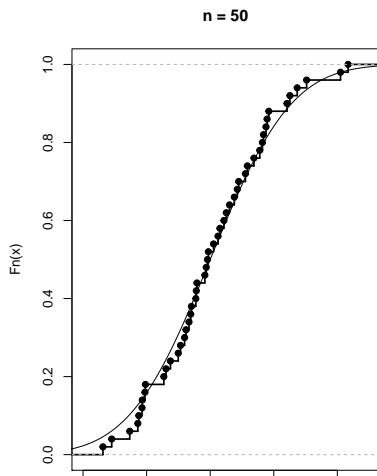
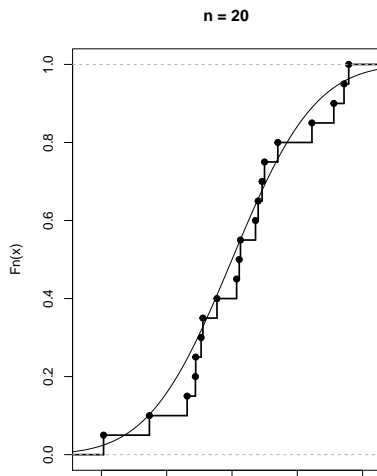
`verticals`

Convergence

`https://jdatta.shinyapps.io/eCDFdemo/`

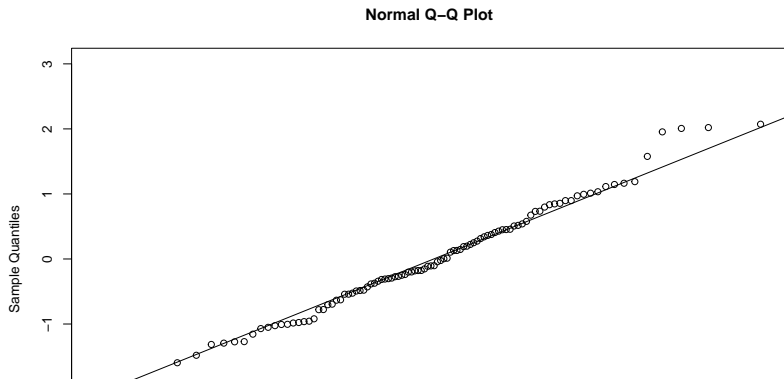
Empirical CDF

```
set.seed(123)
emp1 <- ecdf(rnorm(20)) # E-CDF of 20 samples from  $N(0,1)$ 
emp2 <- ecdf(rnorm(50)) # E-CDF of 20 samples from  $N(0,1)$ 
```



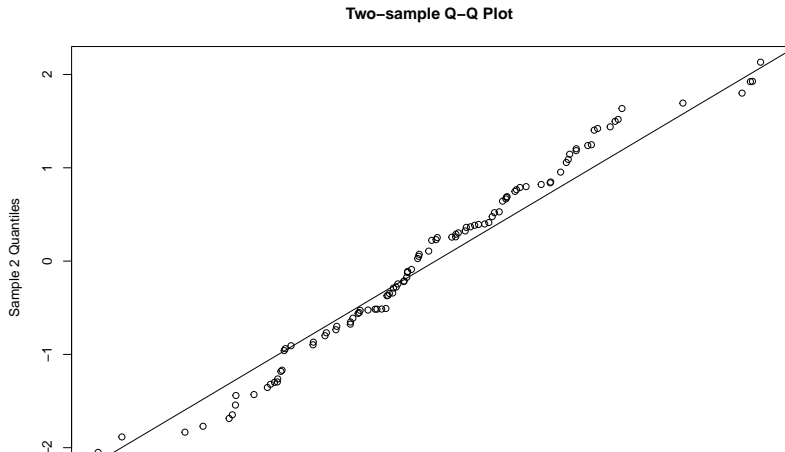
Q-Q Plots (One-sample)

```
set.seed(12) # Reproducibility
y <- rnorm(100)
qqnorm(y, ylim=c(-3,3), main = "Normal Q-Q Plot",
       xlab = "Theoretical Quantiles", ylab = "Sample Quantiles",
       plot.it = TRUE)
qqline(y, distribution = qnorm)
```



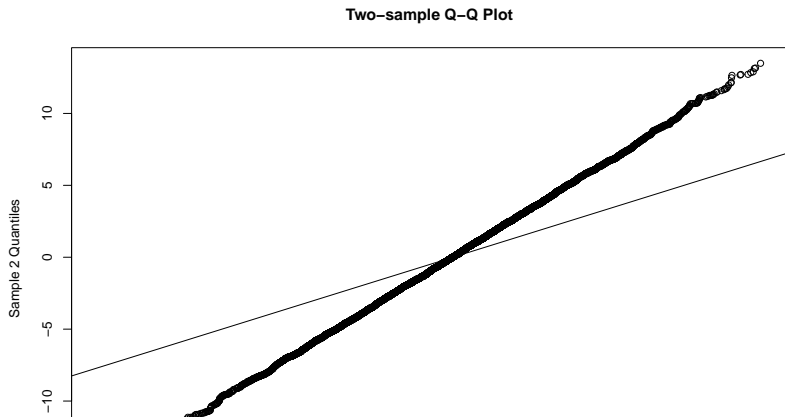
Two samples same

```
z <- rnorm(100)
qqplot(y,z,main = "Two-sample Q-Q Plot",
       xlab = "Sample 1 Quantiles", ylab = "Sample 2 Quantiles",
       abline(0, 1))
```



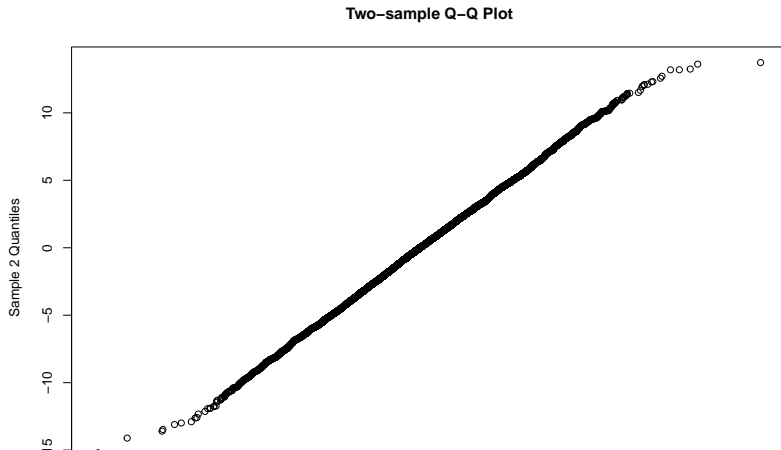
Two samples different

```
x = rnorm(10000,0,2)
y = rnorm(10000,0,4)
qqplot(x,y,main = "Two-sample Q-Q Plot",
        xlab = "Sample 1 Quantiles", ylab = "Sample 2 Quantiles",
        abline(0, 1))
```



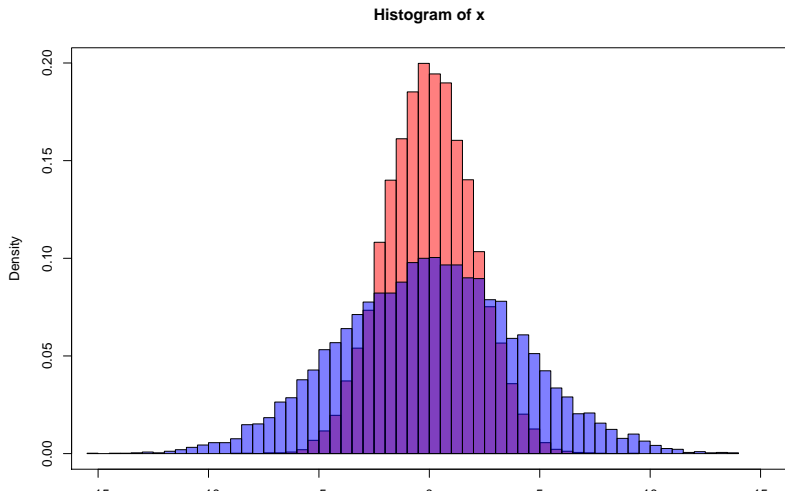
A Common Mistake

```
x = rnorm(10000,0,2)
y = rnorm(10000,0,4)
qqplot(x,y,main = "Two-sample Q-Q Plot",
        xlab = "Sample 1 Quantiles", ylab = "Sample 2 Quantiles")
```



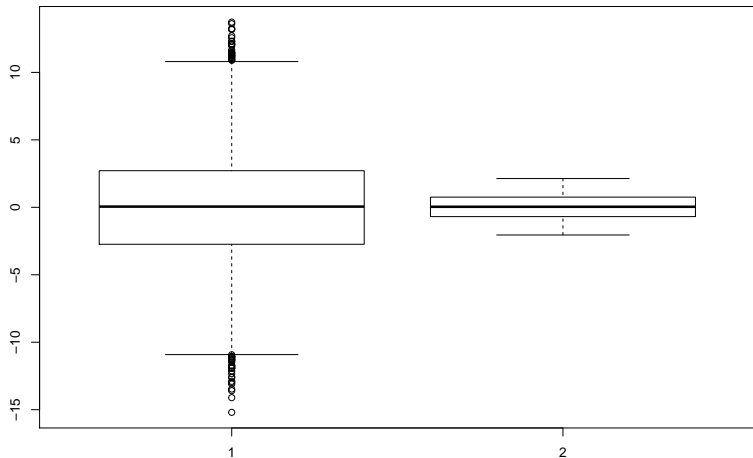
Histograms

```
hist(x,breaks=50,freq=F,col=rgb(1,0,0,0.5),xlim=c(-15,15))  
hist(y,breaks=50,freq=F,col=rgb(0,0,1,0.5),add=T)  
box()
```



Or, Boxplots

```
boxplot(y,z)
```



Even better, test a hypothesis

```
ks.test(rnorm(20),pnorm)
```

```
##  
##  One-sample Kolmogorov-Smirnov test  
##  
## data:  rnorm(20)  
## D = 0.24294, p-value = 0.1593  
## alternative hypothesis: two-sided
```

In our case

```
ks.test(x,y)
```

```
##  
##  Two-sample Kolmogorov-Smirnov test  
##  
## data:  x and y  
## D = 0.1691, p-value < 2.2e-16  
## alternative hypothesis: two-sided
```

Next Time

- ▶ We will dig deeper into these tests !
- ▶ Kolmogorov-Smirnov, Lilliefors, Chi-square GoF etc.
- ▶ These are important as they tell you whether you should use NPStats or Normal Stats.