

# YSU ASDS, Statistics, Fall 2019

## Lecture 04

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09 Sep 2019

# Descriptive Statistics

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- ▶ Empirical CDF
- ▶ Histograms

# Last Lecture ReCap

- ▶ What is a **Frequency Table**?

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- ▶ What is the Definition of the **ECDF**?

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- ▶ What is a **Frequency Table**?
- ▶ What is the Definition of the **ECDF**?
- ▶ What is it for?

# Histograms

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To define the Histogram, first we divide the range of our Dataset into *class intervals* or *bins*:

- ▶ we take first the range: either  $I = [\min_i \{x_i\}, \max_i \{x_i\}]$  or  $I$  is an interval containing  $[\min_i \{x_i\}, \max_i \{x_i\}]$ ;

# Histograms

- ▶ we take a finite partition of  $I$ :  $I_1, I_2, \dots, I_k$ , i.e.  $I_j$ -s are disjoint, and their union is the interval  $I$ ;

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- ▶ we calculate the number  $n_j$  of datapoints  $x_i$  lying in  $I_j$ :

$$n_j = \text{the number of data points in } I_j \quad j = 0, 1, 2, \dots, k.$$

---

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# Histograms

**Definition:** The **frequency histogram** of our continuous (or a grouped) data  $x_1, \dots, x_n$  is the piecewise constant function

$$h_{freq}(x) = n_j, \quad \forall x \in I_j, \quad j = 1, 2, \dots, k.$$

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Frequency histogram shows the number of observations in our dataset in each bin, in each class interval. One also defines  $h_{freq}(x) = 0$  for all  $x \notin I$ .

## Example

*airquality* is a Dataset (standard Dataset in **R**) about the daily air quality measurements in New York, May to September 1973.

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Here is the header:

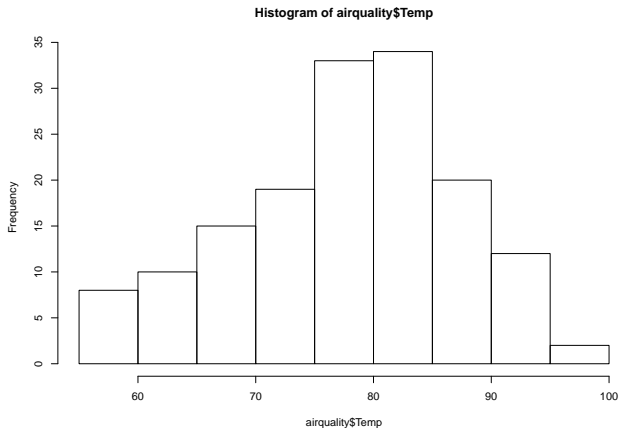
```
head(airquality)
```

##	Ozone	Solar.R	Wind	Temp	Month	Day
## 1	41	190	7.4	67	5	1
## 2	36	118	8.0	72	5	2
## 3	12	149	12.6	74	5	3
## 4	18	313	11.5	62	5	4
## 5	NA	NA	14.3	56	5	5
## 6	28	NA	14.9	66	5	6

## Example

Let's Plot the histogram of the *Temp* (Temperature) Variable:

```
hist(airquality$Temp)
```



## Notes on the Example

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- ▶ **R**'s *hist* command default bins have equal lengths;
- ▶ **R** is adding the default *OX* axis name and the Figure Title.

# Histograms

Next is the Relative Frequency Histogram definition:

**Definition** The **relative frequency histogram** of our continuous data  $x_1, \dots, x_n$  is the piecewise constant function

$$h_{\text{relfreq}}(x) = \frac{n_j}{n}, \quad \forall x \in I_j, \quad j = 1, 2, \dots, k.$$

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The Default **R** package has no Relative Frequency Histogram Plotting command (or I do not know ☺). But you can use, say, the *lattice* library's *histogram* command:

```
library(lattice)
histogram(airquality$Temp)
```

# The Density or Normalized Relative Frequency Histogram

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$$h_{dens}(x) = \frac{n_j}{n} \cdot \frac{1}{length(I_j)}, \quad \forall x \in I_j.$$

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$$h_{dens}(x) = \frac{n_j}{n} \cdot \frac{1}{length(I_j)}, \quad \forall x \in I_j.$$

Here  $length(I_j)$  is the length of the interval  $I_j$ . Also we define  $h(x) = 0$ , if  $x \notin I$ .



## Note

In the case (which is the mostly used one) when all intervals  $I_j$  have the same length:

$$\text{length}(I_j) = h,$$

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$$h_{dens}(x) = \frac{h_{relfreq}(x)}{h} = \frac{n_j}{n \cdot h}, \quad \forall x \in I_j.$$

## Idea of the Density Histogram

The idea of dividing to the length of the corresponding interval, in the definition of the Density Histogram, is that in this case, the Total Area of all rectangles of our Histogram is 1.

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Recall that all PDF functions integrate to 1.

# Idea of the Density Histogram

The idea of dividing to the length of the corresponding interval, in the definition of the Density Histogram, is that in this case, the Total Area of all rectangles of our Histogram is 1.

Recall that all PDF functions integrate to 1. And the Density Histogram is approximating (estimating) the unknown PDF behind our Data!

## Example

To draw the Density Histogram, we will use the *freq=FALSE* parameter in the *hist* command.

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We use here the *discoveries* Standard Dataset from **R**, which gives us the numbers of “great” inventions and scientific discoveries in each year from 1860 to 1959:

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We use here the *discoveries* Standard Dataset from **R**, which gives us the numbers of “great” inventions and scientific discoveries in each year from 1860 to 1959:

```
discoveries
```

```
## Time Series:
```

```
## Start = 1860
```

```
## End = 1959
```

```
## Frequency = 1
```

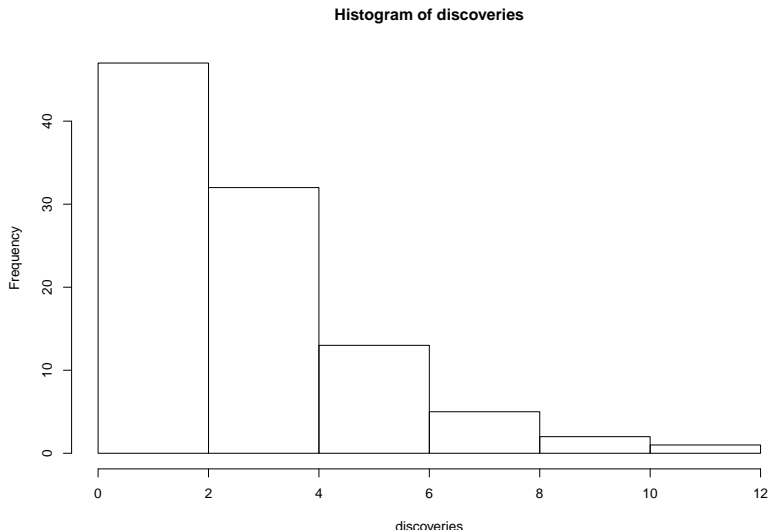
```
##   [1]  5  3  0  2  0  3  2  3  6  1  2  1  2  1  3  3  3
##  [24]  3  7 12  3 10  9  2  3  7  7  2  3  3  6  2  4  3
##  [47]  2  5  2  3  3  6  5  8  3  6  6  0  5  2  2  2  6
##  [70]  7  5  3  3  0  2  2  2  1  3  4  2  2  1  1  1  2
##  [93]  4  1  1  1  0  0  2  0
```



## Example

First, the Frequency Histogram:

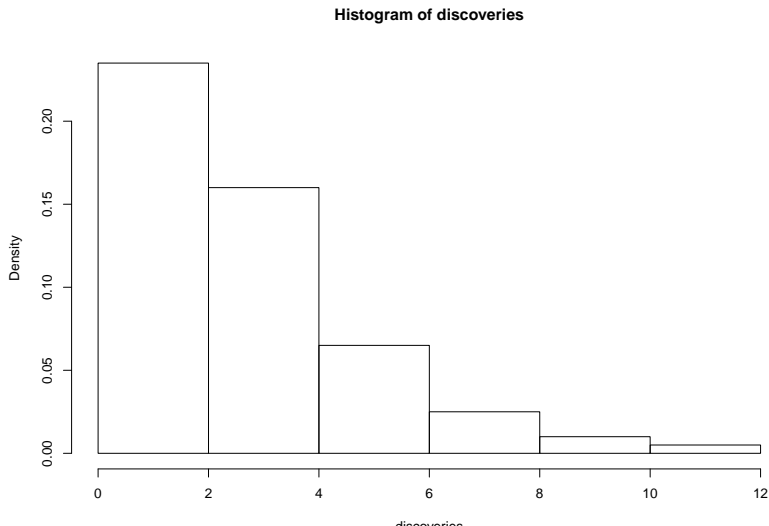
```
hist(discoveries)
```



## Example

Now, the Density Histogram:

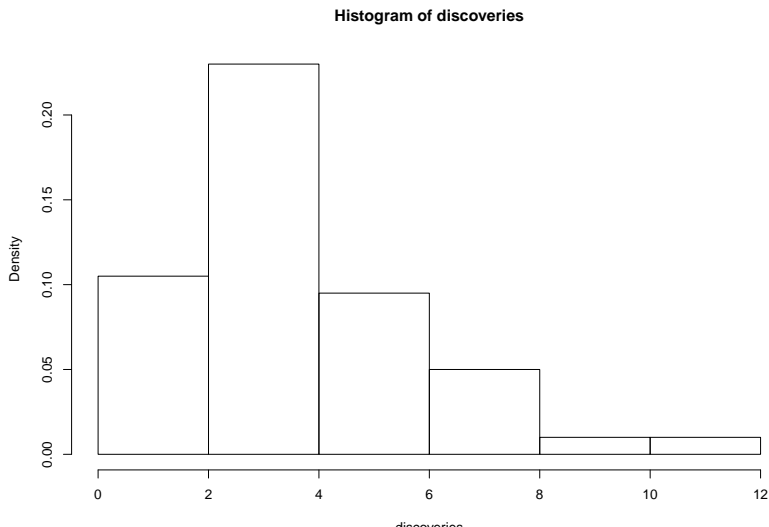
```
hist(discoveries, freq = FALSE)
```



## Example

Finally, the Density Histogram with the Bins left-endpoints included:

```
hist(discoveries, freq = FALSE, right = FALSE)
```



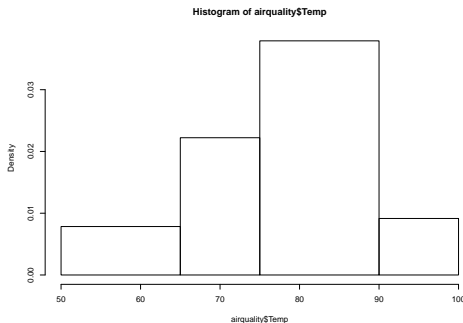
## Example

Now let us change the default bins for a Histogram.

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Now let us change the default bins for a Histogram. We can use the following - first define the vector of our class interval (Bins) endpoints: (note that you need to cover all Datapoints!)

```
bins.endpoints <- c(50, 65, 75, 90, 100)  
hist(airquality$Temp, breaks = bins.endpoints)
```



# Notes

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- ▶ By default, if we give custom bins with non-equal lengths, **R** is plotting the Density Histogram!
- ▶ You can give the *breaks* parameter either the vector of Bins' endpoints or the number of (equal-length) intervals

## Estimation of the PDF through the Density Histogram

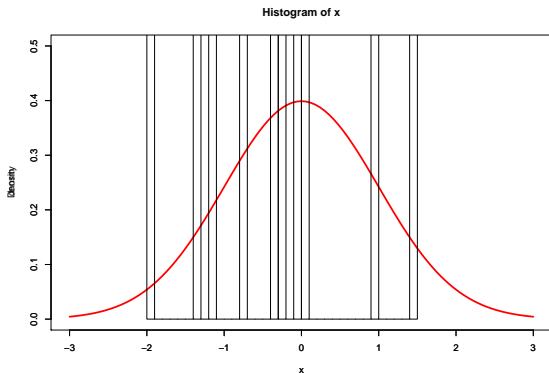
As it was stated above, the Density Histogram is an approximation (estimate) of the PDF of the Data unknown Distribution. To check this, let us take a synthetic Dataset from the Distribution we know:



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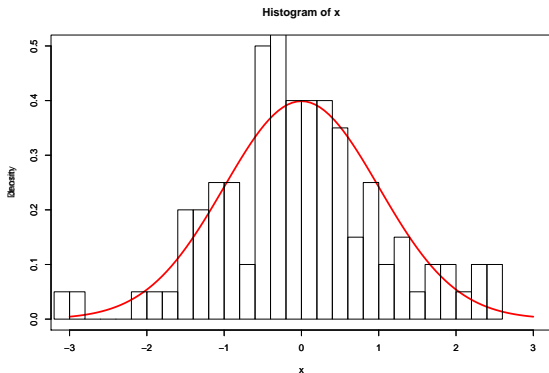
```
plot(dnorm, lwd = 3, col= "red", xlim=c(-3,3), ylim=c(0,0.5))  
x <- rnorm(10)  
par(new = TRUE)  
hist(x, breaks = 40, freq = FALSE, xlim=c(-3,3), ylim=c(0,0.5))
```



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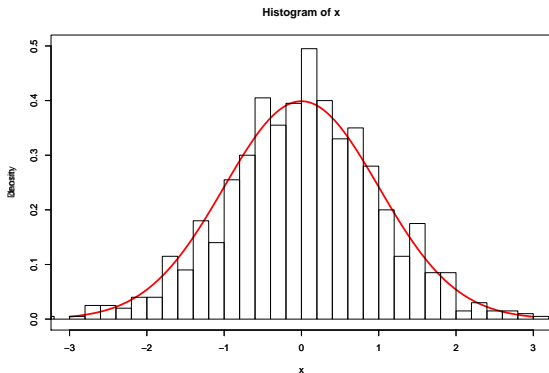
```
plot(dnorm, lwd = 3, col= "red", xlim=c(-3,3), ylim=c(0,0.5))  
x <- rnorm(100)  
par(new = TRUE)  
hist(x, breaks = 40, freq = FALSE, xlim=c(-3,3), ylim=c(0,0.5))
```



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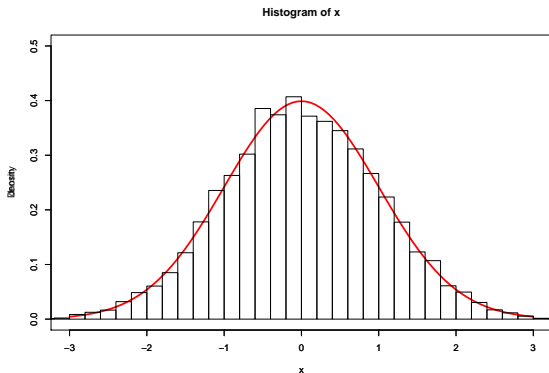
```
plot(dnorm, lwd = 3, col= "red", xlim=c(-3,3), ylim=c(0,0.5))  
x <- rnorm(1000)  
par(new = TRUE)  
hist(x, breaks = 40, freq = FALSE, xlim=c(-3,3), ylim=c(0,0.5))
```



# Estimation of the PDF through the Density Histogram

As it was stated above, the Density Histogram is an approximation (estimate) of the PDF of the Data unknown Distribution. To check this, let us take a synthetic Dataset from the Distribution we know:

```
plot(dnorm, lwd = 3, col= "red", xlim=c(-3,3), ylim=c(0,0.5))  
x <- rnorm(10000)  
par(new = TRUE)  
hist(x, breaks = 40, freq = FALSE, xlim=c(-3,3), ylim=c(0,0.5))
```



## Choosing Bin sizes correctly

It is important to choose the Bin sizes (lengths of the Bin, class, intervals) wisely. Otherwise you will skip some info or you will not get any valuable info.

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It is important to choose the Bin sizes (lengths of the Bin, class, intervals) wisely. Otherwise you will skip some info or you will not get any valuable info.

Let us use another **R** standard dataset to show the effect of the choice of the bin size: *precip*. This Dataset shows the average amount of precipitation (rainfall) in inches for each of 70 United States (and Puerto Rico) cities.

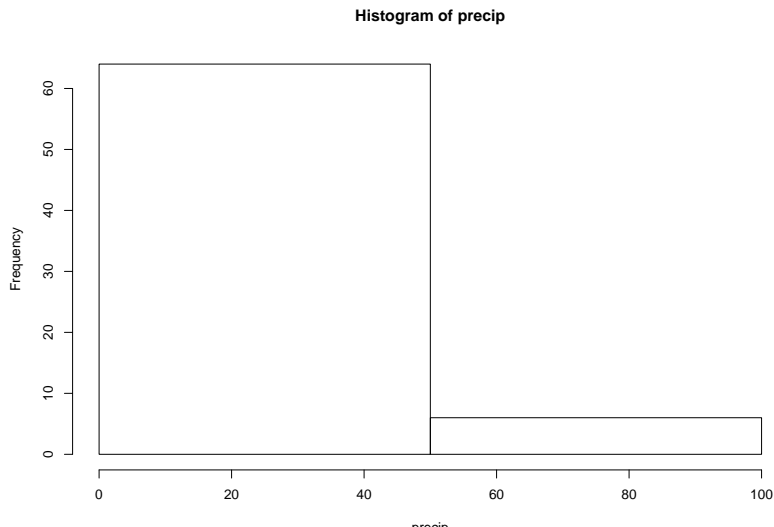
```
head(precip)
```

##	Mobile	Juneau	Phoenix	Little Rock	Los Angeles
##	67.0	54.7	7.0	48.5	1

## Version 1, Small bins

Here, we just use 2 bins:

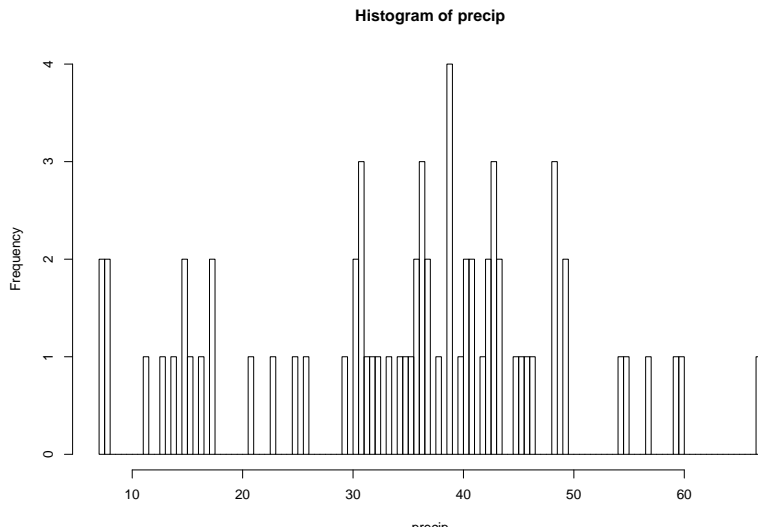
```
hist(precip, breaks = 2)
```



## Version 2, large bins

Here, we use 200 bins:

```
hist(precip, breaks = 200)
```

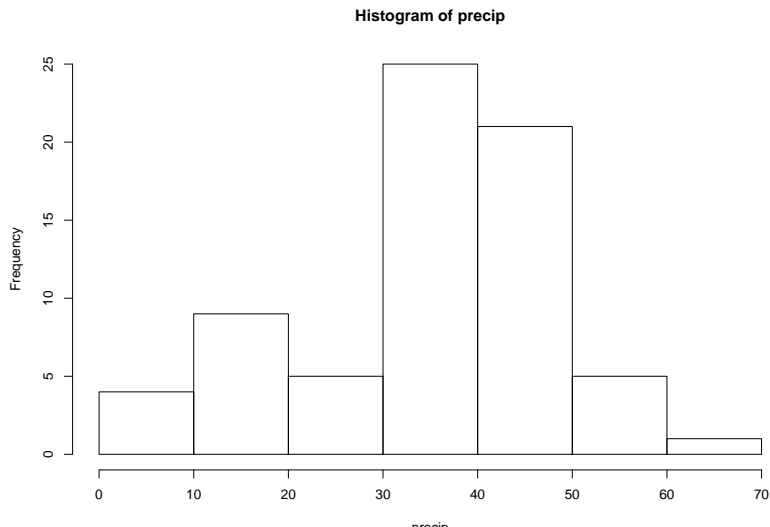




## Version 2, large bins

Now, the default:

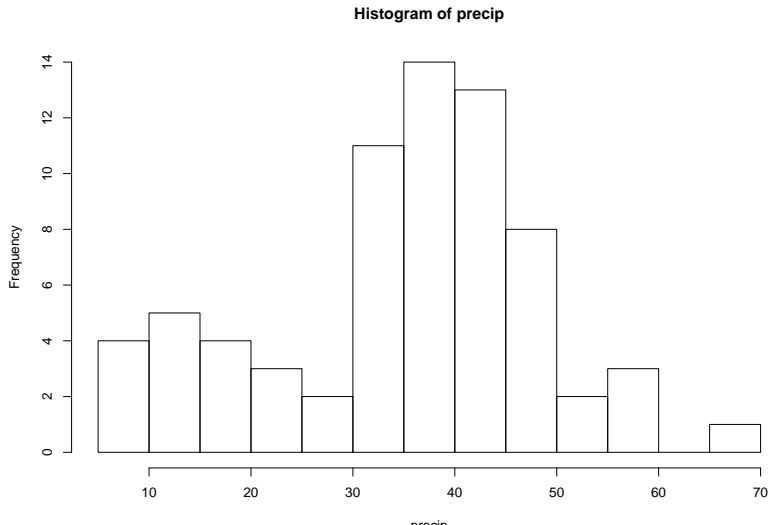
```
hist(precip)
```



## Version 3

Now, let us change to 20 bin intervals:

```
hist(precip, breaks = 20)
```



# Choosing the Bin Length

In fact, choosing the correct Bin width is not an easy job. See, for example, [the Histogram Wiki page](#).

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- ▶ *Barplot* is for a categorical or Discrete Data, *Histogram* is for both Discrete and Continuous

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- ▶ *Barplot* is for a categorical or Discrete Data, *Histogram* is for both Discrete and Continuous
- ▶ We can exactly reconstruct the Dataset from the *Barplot*, but not the *Histogram*

## Addition to the Histogram

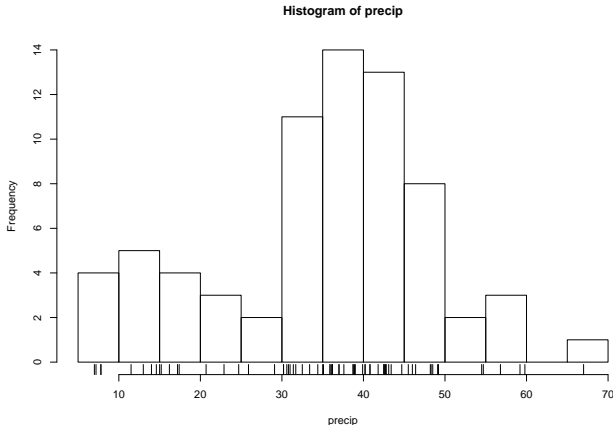
Nice addition to your Histogram Plot is to add, in some way, the Datapoints:



## Addition to the Histogram

Nice addition to your Histogram Plot is to add, in some way, the Datapoints:

```
hist(precip, breaks = 20)  
rug(precip)
```



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- ▶ is spread out or concentrated at some point
- ▶ has some gaps
- ▶ has values far apart from others, has outliers (anomalies)
- ▶ is unimodal, bimodal or multimodal

# KDE

Another estimate for the unknown Distribution PDF is the **Kernel Density Estimator**, KDE.

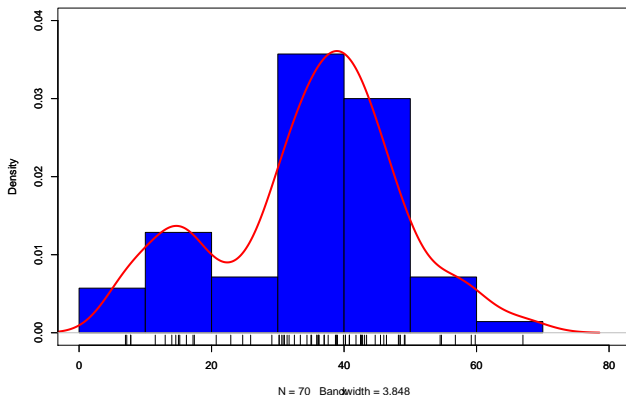


# KDE

Another estimate for the unknown Distribution PDF is the **Kernel Density Estimator**, KDE. It is, in some sense, the smoothed version of the Histogram: Histogram is a piecewise-constant function, with jumps, so it is not a smooth function.

## KDE Example

```
x <- precip; d <- density(x)
hist(x, freq = FALSE, xlim = c(0, 80), ylim = c(0,0.04),
     col = "blue", main = "")
rug(x); par(new = TRUE)
plot(d, lwd = 3, col = "red", xlim = c(0,80), ylim = c(0,0.04),
     main = "")
```



## KDE

To define the KDE, we first choose a smooth Kernel function  $K(t)$ , here, a function with

$$K(t) \geq 0, t \in \mathbb{R}, \quad \text{and} \quad \int_{-\infty}^{+\infty} K(t) dt = 1.$$

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For example, we can take the Gaussian Kernel

$$K(t) = \frac{1}{\sqrt{2\pi}} \cdot e^{-t^2/2}, \quad t \in \mathbb{R},$$

or any other PDF.

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For example, we can take the Gaussian Kernel

$$K(t) = \frac{1}{\sqrt{2\pi}} \cdot e^{-t^2/2}, \quad t \in \mathbb{R},$$

or any other PDF.

Next, one defines the Kernel Density Estimator with Kernel  $K$  as

$$KDE_K(x) = KDE(x) = \frac{1}{nh} \cdot \sum_{i=1}^n K\left(\frac{x - x_i}{h}\right).$$

# KDE

It is easy to see that  $KDE(x)$  will give a PDF, i.e., will be nonnegative and will integrate to 1:

$$\int_{-\infty}^{+\infty} KDE(x) dx =$$

# KDE

It is easy to see that  $KDE(x)$  will give a PDF, i.e., will be nonnegative and will integrate to 1:

$$\begin{aligned}\int_{-\infty}^{+\infty} KDE(x) dx &= \frac{1}{nh} \cdot \sum_{i=1}^n \int_{-\infty}^{+\infty} K\left(\frac{x - x_i}{h}\right) dx = \\&= \frac{1}{n} \cdot \sum_{i=1}^n \int_{-\infty}^{+\infty} K\left(\frac{x - x_i}{h}\right) d\frac{x - x_i}{h} \stackrel{u = \frac{x - x_i}{h}}{=} \\&= \frac{1}{n} \cdot \sum_{i=1}^n \int_{-\infty}^{+\infty} K(u) du = \frac{1}{n} \cdot \sum_{i=1}^n 1 = 1.\end{aligned}$$

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**Note:** Like in the case of the Density histogram, where that histogram was depending on the bins choice, the KDE depends on the choice of  $h > 0$ .  $h$  is called the **bandwidth**, and its estimation is another story.