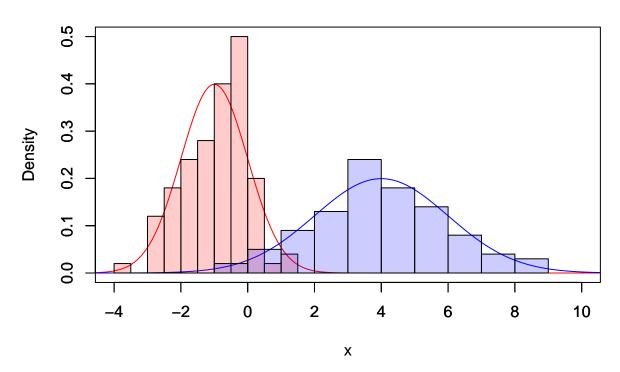
# Hypothesis Testing, Part 1

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### Two Normal Population Distributions, with Sample Data (N=100)



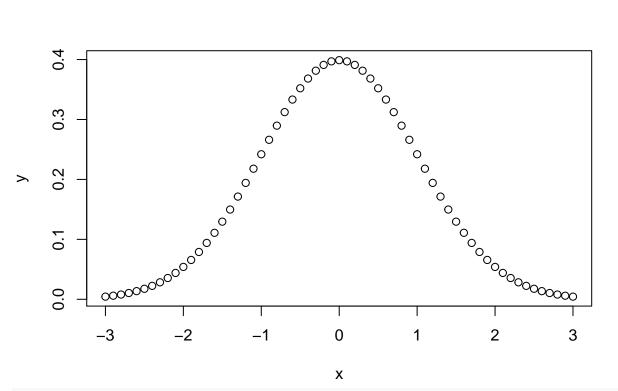
### Querying and Generating Probability Densities

R's built-in stats package comes with lots of functions for working with common probability distributions. The functions we will review here allow you to:

- Get the distribution's (d)ensity height, given x values: dnorm(), dpois(), dunif()
- Get (r)andom amples from the distribution: rnorm(), rpois(), runif()
- Get the distributions's quantiles:
  - The (q)uantile of the distribution, given a percentile: qnorm(), qpois(), qunif()
  - The **(p)ercentile** of the distribution, given a quantile: pnorm(), ppois(), punif()

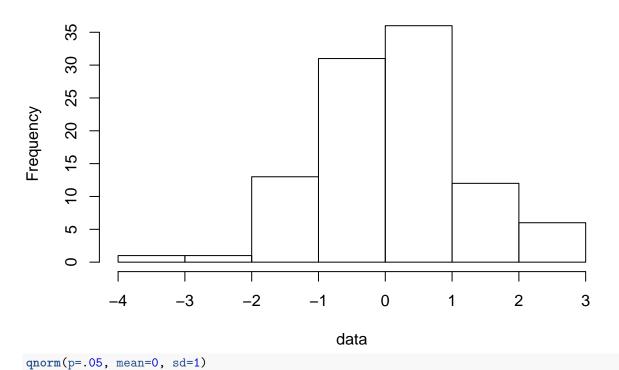
Examples of usage:

```
x <- seq(-3, 3, .1)
y <- dnorm(x, mean=0, sd=1)
plot(x, y)</pre>
```



data <- rnorm(100, mean=0, sd=1)
hist(data)</pre>

## Histogram of data



## [1] -1.644854
pnorm(q=-1.645, mean=0, sd=1)

## [1] 0.04998491

### **T-Tests**

The **t.test()** function does t-tests. Setting different options in the function arguments lets you use it as one-sided or two-sided test, a paired-sample or independent-sample test, assume equal variance, etc. It returns a table of the results:

```
x <- rnorm(100, mean=-0.5, sd=1)
y <- rnorm(100, mean=0.5, sd=1)
t.test(x, y, alternative = "two.sided")
##</pre>
```

```
##
## Welch Two Sample t-test
##
## data: x and y
## t = -5.9944, df = 197.76, p-value = 9.532e-09
## alternative hypothesis: true difference in means is not equal to 0
## 95 percent confidence interval:
## -1.0771268 -0.5438562
## sample estimates:
## mean of x mean of y
## -0.4332610 0.3772305
```

Nonparametric alternatives to the t-test are also available: - Kruskall-Wallis Test: kruskall.test() - Mann-Whitney U Test: wilcox.test() - Wilcoxon Rank Sum Test: wilcox.test(paired=TRUE)

If you save the result of the test as a variable, you can also access the details of the table and index the names of the result to access the data:

#### Correlations

Testing for correlations between paired samples can be done with the cor.test() function. By default it does a Pearson correlation, but can also do a Spearman or Kendall's Tau test by changing the "method" argument

#### **Exercises**

Exercises are available at the hypothesis\_testing\_exercises.R script.