

# Advanced Quantitative Methods: Hypothesis testing

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# Probability distributions

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We extensively discussed the **normal distribution**.

There are many different probability distributions, however, and many of them are related.

# Bernoulli trial

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“An experiment in which  $s$  trials are made of an event, with probability  $p$  of success in any given trial.”

(Weisstein, Eric W. “Bernoulli Trial.” <http://mathworld.wolfram.com/BernoulliTrial.html>)

# Binomial distribution

“The (...) probability distribution (...) of obtaining exactly  $n$  successes out of  $N$  Bernoulli trials.”

$$P(n|N) = \frac{N!}{n!(N-n)!} p^n (1-p)^{N-n}$$

(Weisstein, Eric W. “Binomial Distribution.” <http://mathworld.wolfram.com/BinomialDistribution.html>)

## Binomial distribution

$$P(n|N) = \frac{N!}{n!(N-n)!} p^n (1-p)^{N-n}$$

$$\lim_{N \rightarrow \infty} p(n) = \frac{1}{\sqrt{2\pi Npq}} e^{-\frac{(n-Np)^2}{2Npq}}$$

$$= \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(n-\bar{n})^2}{2\sigma^2}}$$

i.e. the **limiting distribution** of the binomial distribution is the **normal distribution**, with  $\sigma^2 \equiv Npq$ .

(Weisstein, Eric W. "Binomial distribution." <http://mathworld.wolfram.com/binomialdistribution.html>)

# Normal distribution

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Also called **Gaussian distribution**, but Gauss did not invent it.

(Davidson & MacKinnon 1999: 130-135)

## $\chi^2$ -distribution

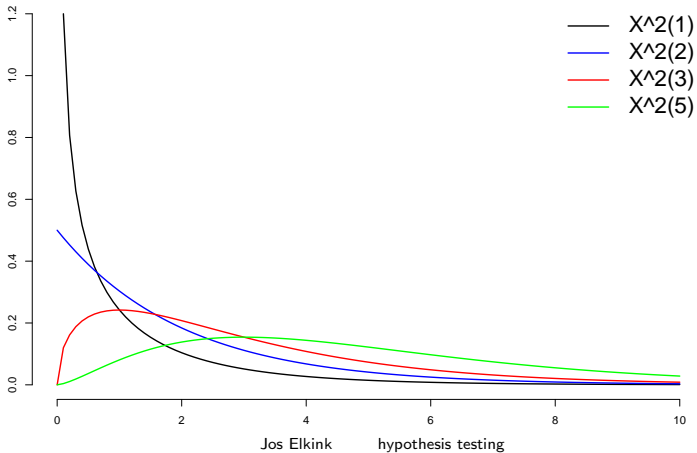
The sum of squares of  $r$  independent normal distributions, is distributed chi-squared with  $r$  degrees of freedom:

$$\chi^2(r) \equiv \sum_i^r x_i^2$$

(Weisstein, Eric W. "Chi-squared distribution." <http://mathworld.wolfram.com/chi-squaredistribution.html>)



# $\chi^2$ -distribution



# F-distribution

If

$$x \sim \chi^2(m)$$

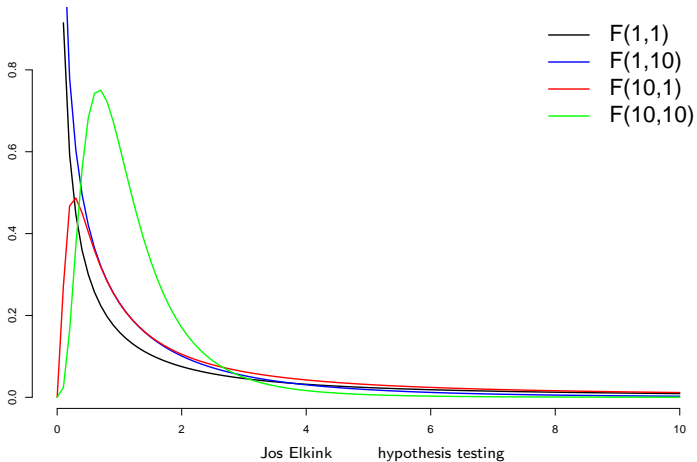
$$y \sim \chi^2(n)$$

with  $x, y$  independent, then

$$F(m, n) \equiv \frac{x/m}{y/n}$$

has an  $F$ -distribution with  $m$  and  $n$  degrees of freedom.

# F-distribution



## $t$ -distribution

If

$$x \sim n(0, 1)$$

$$y \sim \chi^2(r)$$

with  $x$ ,  $y$  independent, then

$$t(r) \equiv \frac{x}{\sqrt{y/r}}$$

has a  $t$ -distribution with  $r$  degrees of freedom.

## $t$ -distribution

Imagine we have a sample of size  $n$  and we calculate the sample mean of  $x$ :

$$\bar{x} = \frac{1}{n} \sum_i^n x_i$$

with variance estimator

$$s^2 = \frac{1}{n-1} \sum_i^n (x_i - \bar{x})^2$$

then

$$\frac{(n-1)s^2}{\sigma^2}$$

is distributed  $\chi^2(n-1)$  (because  $s^2$  is a sum of squares).

## $t$ -distribution

$\bar{x}$  is a sum of normally distributed values, so is itself normally distributed;  $s^2$  has a chi-squared distribution, so

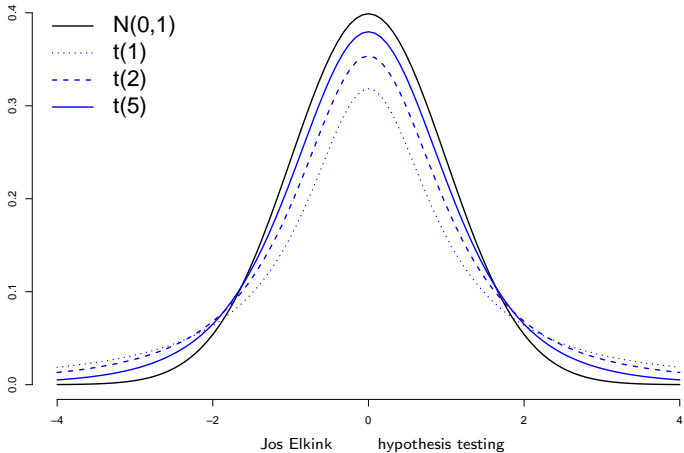
$$t(n) = \frac{\bar{x} - \mu}{\sqrt{s^2/n}}$$

has a  $t$ -distribution with  $n$  degrees of freedom.

As  $n$  increases, the  $t$ -distribution approaches a normal distribution. The  $t$ -distribution is the approximation for the normal distribution when  $n$  is small and  $\sigma^2$  unknown.

(The  $t(1)$  distribution is also called the **Cauchy distribution**.)

# $t$ -distribution

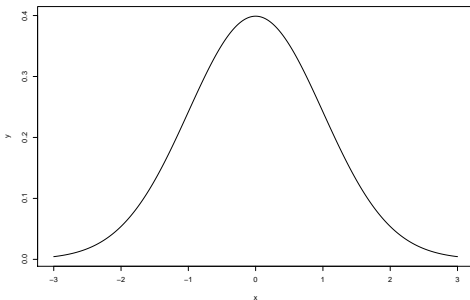


# Probability distributions in R

- You know  $x$  and want to know the **density** at that point: **d**
- You know  $x$  and want to know the **area** up to that point: **p**
- You know  $x$  and want to know the area beyond that point:  
**1-p**
- You know the area and want to know the  $x$  value: **q**
- You want **random** numbers drawn from that distribution: **r**

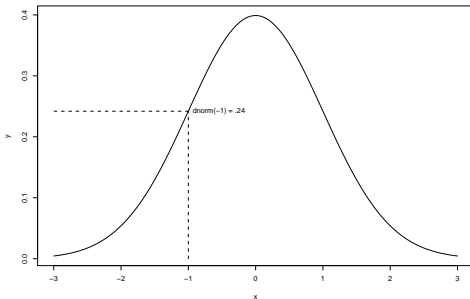


## Probability distributions in R: dnorm



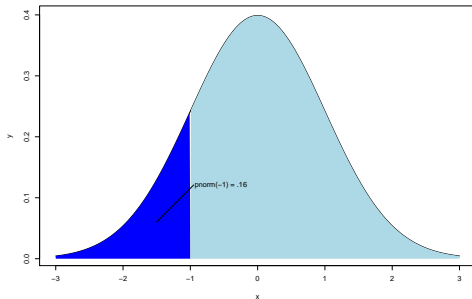
```
x <- seq(-3,3,.01)  
y <- dnorm(x)
```

## Probability distributions in R: dnorm



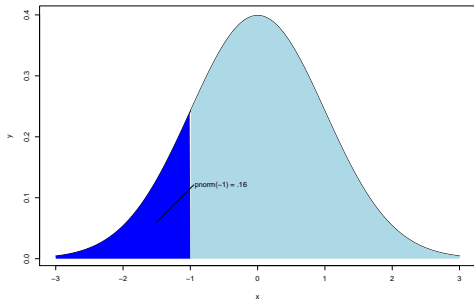
```
> dnorm(-1)
[1] 0.2419707
```

## Probability distributions in R: pnorm



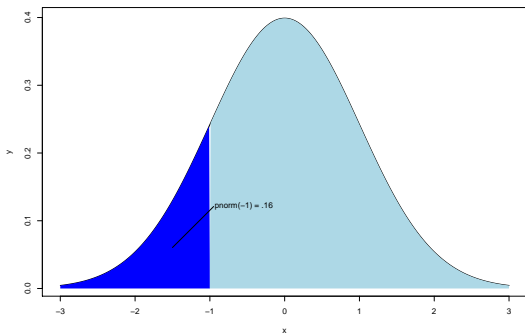
```
> pnorm(-1)  
[1] 0.1586553
```

## Probability distributions in R: pnorm



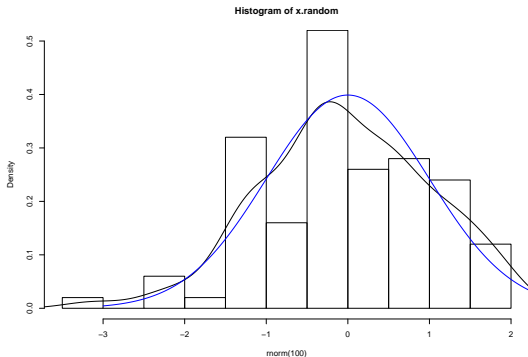
```
> 1 - pnorm(-1)
[1] 0.8413447
```

## Probability distributions in R: pnorm



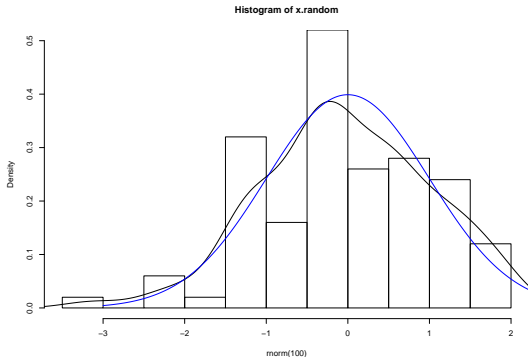
```
> qnorm(.1586553)  
[1] -0.9999998
```

# Probability distributions in R: rnorm



```
x.random <- rnorm(100)
```

# Probability distributions in R: rnorm



```
hist(x.random, freq=false)
```

# Probability distributions in R

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These functions work on many distributions:

<code>rnorm()</code>	random from normal distribution
<code>pchisq()</code>	get area under $\chi^2$ -distribution
<code>1-pf()</code>	get area under $F$ -distribution
<code>rbinom()</code>	draw randomly from binomial distribution
<code>1-dt()</code>	get $p$ -value from $t$ -distribution (one-tailed)



## Probability distributions in R

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Twenty throws (“Bernoulli trials”) with a coin:

```
> x <- rbinom(20,1,.5)
> factor(x, labels=c("head","tails"))
[1] head  head  head  tails head  tails tails head
[9] tails head  head  head  tails tails tails head
[17] tails tails tails head
```

## Type I and II errors

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**Type I error:** rejecting a null hypothesis that is true (e.g.  $\alpha = .05$ )

**Type II error:** not rejecting a null hypothesis that is false

(Davidson & MacKinnon 1999: 126)

# Power

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**Power:** probability of rejecting a hypothesis that is false

$$1 - P(\text{Type II error})$$

The power of a test increases when:

- the true value is further from the null hypothesis value;
- the variance is lower;
- the sample size is larger.

(Davidson & MacKinnon 1999: 126)

## $p$ -value

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The  $p$ -value is the probability of a Type I error when rejecting the null hypothesis.

You can say a test is “statistically significant” if  $p < \alpha$ , but the  $p$ -value contains more information by itself.

(Davidson & MacKinnon 1999: 128)

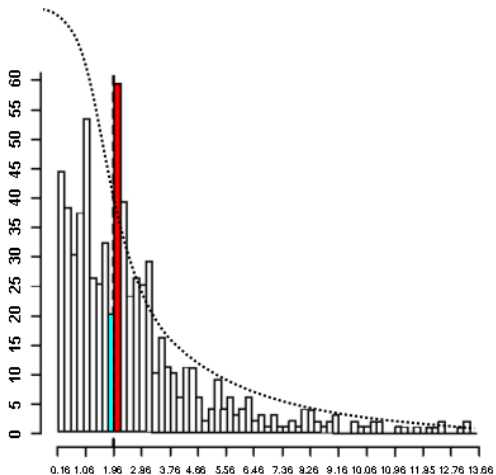
$$\alpha = .05$$

Note that his value is absolutely arbitrary and just habit since the publication of Fisher (1923).

The  $p$ -value is, one could argue, just a complicated way of measuring the sample size.

“Another interesting example (...) is the propensity for published studies to contain a disproportionately large number of Type I errors; studies with statistically significant results tend to get published, whereas those with insignificant results do not.” (Kennedy 2008: 61)

$$\alpha = .05$$



# Confidence intervals

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Instead of an arbitrary threshold it is often more illuminating to present **confidence intervals** or graphical presentations of levels of uncertainty.

## t-test

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$$h_0 : \beta = 0$$

$$h_1 : \beta \neq 0$$

We can calculate the  $t$ -value by subtracting the value under the null and dividing by the standard error:

$$t = \frac{\hat{\beta}}{\sigma_{\hat{\beta}}}$$

Because  $\hat{\beta}$  has a normal distribution and  $\sigma_{\hat{\beta}}^2$  a  $\chi^2$ -distribution,  $t$  has the  $t$ -distribution with  $n - k$  degrees of freedom.



## t-test

```
bhat <- solve(t(x) %*% x) %*% t(x) %*% y
e <- y - x %*% bhat
vhat <- (1/(n-k) * t(e) %*% e) %x% solve(t(x) %*% x)
se <- sqrt(diag(vhat))
p <- 2 * (1 - pt(abs(bhat / se), n-k))
cbind(bhat, se, p)
```

abs() absolute value

2 \* because it is a two-tailed test

## t-test

$$h_0 : \beta = 0$$

$$h_1 : \beta \neq 0$$

$$t = \frac{\hat{\beta}}{\sigma_{\hat{\beta}}}$$

$$h_0 : \beta = a$$

$$h_1 : \beta \neq a$$

$$t = \frac{\hat{\beta} - a}{\sigma_{\hat{\beta}}}$$

$$h_0 : \beta_3 = \beta_2$$

$$h_1 : \beta_3 \neq \beta_2$$

$$t = \frac{\hat{\beta}_3 - \hat{\beta}_2}{\sigma_{\hat{\beta}_3}}$$

## Sums of squares

SST Total sum of squares  $\sum (y_i - \bar{y})^2$

SSE Explained sum of squares  $\sum (\hat{y}_i - \bar{y})^2$

SSR Residual sum of squares  $\sum e_i^2 = \sum (\hat{y}_i - y_i)^2 = \mathbf{e}'\mathbf{e}$

The key to remember is that **SST = SSE + SSR**

Sometimes instead of “explained” and “residual”, “regression” and “error” are used, respectively, so that the abbreviations are swapped (!).

# Anova

Variation	SS	df	MS
Explained	$\sum(\hat{y}_i - \bar{y})^2$	$k - 1$	$SSE/(k - 1)$
Residual	$\sum(\hat{y}_i - y_i)^2$	$n - k$	$SSR/(n - k)$
Total	$\sum(y_i - \bar{y})^2$	$n - 1$	

$$\frac{SSE/(k - 1)}{SSR/(n - k)} \sim F_{k-1, n-k}$$

## F-test

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```
ssr <- t(e) %*% e
yhat <- X %*% bhat
sse <- t(yhat - mean(y)) %*% (yhat - mean(y))
F <- (sse/(k-1))/(ssr/(n-k))
1 - pf(F, k-1, n-k)
```

## Restrictions

$$H_0 : \mathbf{y} = \mathbf{X}_{n \times k}^{(1)} \boldsymbol{\beta}_1 + \boldsymbol{\varepsilon}$$

$$H_1 : \mathbf{y} = \mathbf{X}_{n \times k}^{(1)} \boldsymbol{\beta}_1 + \mathbf{X}_{n \times r}^{(2)} \boldsymbol{\beta}_2 + \boldsymbol{\varepsilon}$$

USSR Unrestricted sum of squared residuals

RSSR Restricted sum of squared residuals

$$F_{\beta_2} = \frac{(RSSR - USSR)/r}{USSR/(n - k - r)} \sim F(r, n - k - r)$$

## Chow test

You might have a situation where you want to know whether the coefficients differ for two groups in the sample.

You could test this with the following regression:

$$\begin{bmatrix} \mathbf{y}_1 \\ \mathbf{y}_2 \end{bmatrix} = \begin{bmatrix} \mathbf{X}_1 \\ \mathbf{X}_2 \end{bmatrix} \beta + \begin{bmatrix} \mathbf{0} \\ \mathbf{X}_2 \end{bmatrix} \gamma + \varepsilon,$$

whereby the difference between the coefficients for the two groups are captured by  $\gamma$ .

# Chow test

It turns out you can just run two separate regressions (“unrestricted”), for the two groups, as well as (“restricted”) regression  $\mathbf{y}$  on  $\mathbf{X}$ , so that we get:

$$F_{\gamma} = \frac{(RSSR - SSR_1 - SSR_2)/k}{(SSR_1 + SSR_2)/(n - 2k)} \sim F(k, n - 2k)$$

(Davidson & MacKinnon 1999: 146-147)



# Likelihood

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“The likelihood that any parameter (or set of parameters) should have any assigned value (or set of values) is proportional to the probability that if this were so, the totality of observations should be that observed.”

(Fisher 1922, 310)

# Maximum Likelihood

The likelihood is proportional to the probability of observing the data you have (give or take some arbitrarily small deviation), given some parameter estimate:

$$L(\beta|\mathbf{y}, \mathbf{X}) = \alpha P(\mathbf{y}, \mathbf{X}|\beta)$$

The **likelihood function** is thus proportional to the **probability density function** of the given sample, as a function of the parameter values.

The estimator that maximizes this function also maximizes this probability density function and is the **Maximum Likelihood Estimator** (MLE or ML).

(Fisher 1922)

# Three tests

Three tests are based on this likelihood:

- **Likelihood Ratio (LR)** test:

$$H_0 : 2 \log \frac{L_U}{L_R} = 2(\log L_U - \log L_R) = 0$$

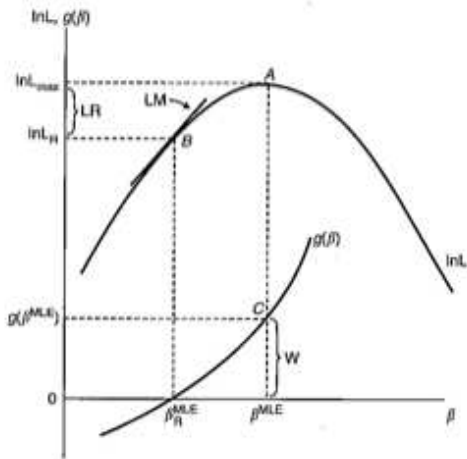
- **Wald (W)** test:

$$H_0 : \frac{(\hat{\beta}^{MLE} - \beta^*)^2}{\text{var}(\hat{\beta}^{MLE})} = 0$$

- **Lagrange Multiplier (LM)** test:

$$H_0 : \frac{\partial \ln L}{\partial \beta} = 0$$

# Three tests



## Three tests

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Asymptotically, LR, W, and LM are all  $\chi^2$ -distributed.

In small samples,  $W \geq LR \geq LM$ .

We return to these tests and how to do them in R later in the course.

(Kennedy 2008: 64)