

MASM22/FMSN30: Linear and Logistic Regression, 7.5 hp

FMSN40: ... with Data Gathering, 9 hp

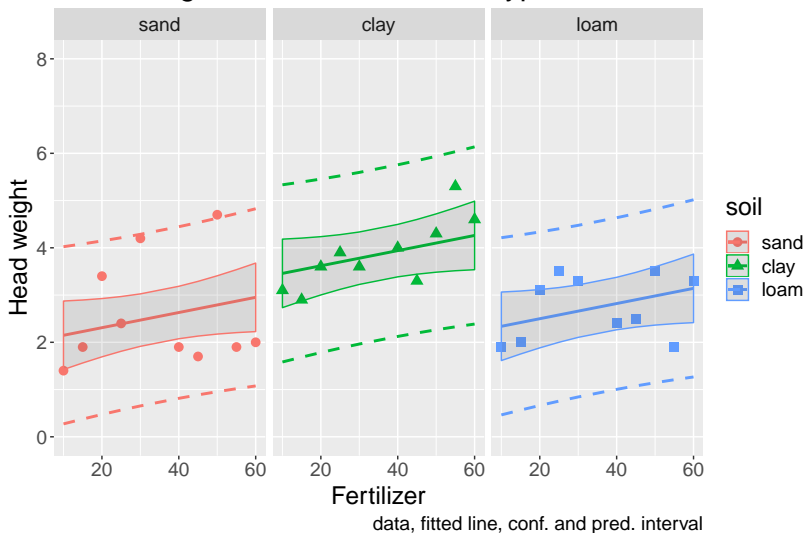
Lecture 4, spring 2023

Significance tests for model parameters

Mathematical Statistics / Centre for Mathematical Sciences
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29/3-23

Head weight as a function of soil type and fertilizer



Model: $Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i3} + \epsilon_i$

where $x_1 = 1$ if clay (0 if not clay), $x_2 = 1$ if loam (0 if not loam) and $x_3 = \text{fertilizer}$.

Questions

- ▶ Does the amount of fertilizer have a significant effect on the head weight, when we already have soil type in the model?
Test $H_0: \beta_3 = 0$ against $H_1: \beta_3 \neq 0$. **t-test**.
- ▶ Are there significant systematic differences between any of the soil types, when we have fertilizer in the model?
Test $H_0: \beta_1 = \beta_2 = 0$ against H_1 : “at least one of β_1 and β_2 is $\neq 0$ ”. **Partial F-test**.
- ▶ Is the model better than nothing?
Test $H_0: \beta_1 = \beta_2 = \beta_3 = 0$ against H_1 : “at least one of β_1 , β_2 and β_3 is $\neq 0$ ”. **Global F-test**.

Hypothesis testing

We want to **test** a null hypothesis H_0 against an alternative hypothesis H_1 . We will reject H_0 in favour of H_1 only if our results are very unlikely to appear by chance if H_0 were true. Thus, we want to either

- ▶ determine if the difference between what we observed and what we would expect to observe if H_0 were true, is too large to be due to just random variability, or, equivalently,
- ▶ determine if the chance of observing something this far away from what we expected if H_0 were true, by chance, is small.

Significance level

We define the **significance level** α of a test by

$$\alpha = P(\text{reject } H_0 \mid H_0 \text{ is true})$$

This is the probability of concluding that H_0 is wrong when it is in fact correct. False positive rate. Typically we choose $\alpha = 0.05$.

Test statistic

A **test statistic** is a suitable function of our observations that has a known distribution when H_0 is true, and can be expected to lie in the extreme end(s) of this distribution if H_0 is not true.

- ▶ Reject H_0 at significance level α if the test statistic lies above the α quantile (or outside the $1 - \alpha/2$ and $\alpha/2$ quantiles) in its distribution. One-sided or two-sided depends on the type of statistic.

P-value

The P-value of a test is defined as the probability of observing as large a result as the one produced by your data, **or an even more extreme result**, when H_0 is true. If the **P-value** is small our result is more extreme than would be expected by chance and H_0 is unlikely to be true.

- ▶ Reject H_0 at significance level α if **P-value** $< \alpha$.

We can use the test statistic and its distribution to calculate the P-value.

t-test for β_j

We want to test if one x -variable, x_j , has any relevance in the model, **given all the other variables**.

We want to test $H_0: \beta_j = 0$ against $H_1: \beta_j \neq 0$.

Since $\hat{\beta}_j \sim N(\beta_j, \sigma^2(\mathbf{X}'\mathbf{X})_{jj}^{-1}) = N(0, \sigma^2(\mathbf{X}'\mathbf{X})_{jj}^{-1})$ when H_0 is true, a suitable test statistic is

$$t = \frac{\hat{\beta}_j - 0}{d(\hat{\beta}_j)} = \frac{\hat{\beta}_j}{s\sqrt{(\mathbf{X}'\mathbf{X})_{jj}^{-1}}} \sim t(n - (p + 1))$$

and we should reject H_0 at significance level α if

- ▶ $|t| > t_{\alpha/2, n-(p+1)}$ or
- ▶ $P\text{-value} = P(|t(n - (p + 1))| > |t|) < \alpha$ or
- ▶ $0 \notin I_{\beta_j}$.

All three variants are equivalent.

Cabbage: soil and fertilizer estimates

Variable	param	est	s.e.	t-value	P-value	95 % C.I.
intercept	β_0	1.99	0.42	4.74	$7 \cdot 10^{-5}$	(1.13, 2.85)
clay	β_1	1.31	0.38	3.48	0.002	(0.54, 2.08)
loam	β_2	0.19	0.38	0.50	0.62	(-0.58, 0.96)
fertilize	β_3	0.016	0.009	1.73	0.09	(-0.003, 0.035)
resid.std.dev	σ	0.84	df = 26			

- ▶ Since $|t| = \left| \frac{0.016}{0.009} \right| = |1.73| \not> t_{0.05/2, 26} = 2.06$ we should not reject $H_0: \beta_3 = 0$ at significance level $\alpha = 5\%$.
- ▶ Since $P\text{-value} = P(|t(26)| > 1.73) = 0.09 \not< 0.05$ we should not reject H_0 at significance level $\alpha = 5\%$
- ▶ Since the confidence interval for β_3 covers 0 we should not reject H_0 at significance level $\alpha = 5\%$.

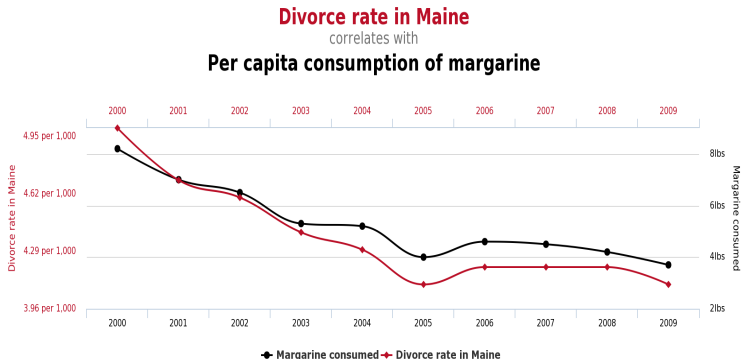
The amount of fertilizer does not have a significant effect on head weight.

R: two-sided quantile and P-value in a $t(f)$ -distribution

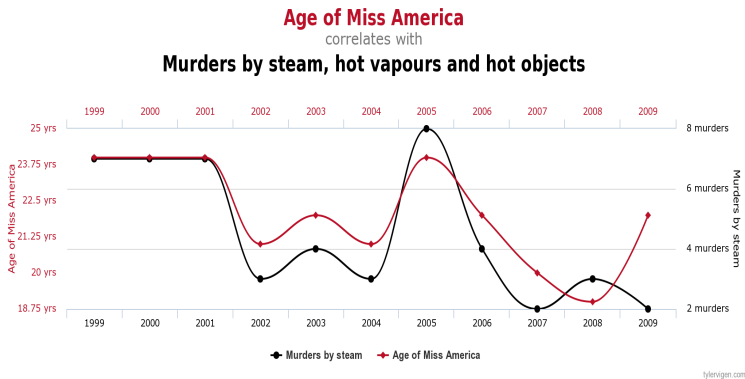
quantile = qt(alpha/2, f, lower.tail = FALSE) = qt(1 - alpha/2, f)

P-value = 2*pt(abs(t), f, lower.tail = FALSE)

Some relations that are absurd and irrelevant



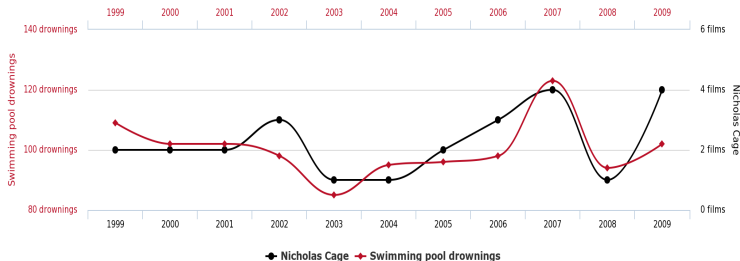
tylervigen.com



Number of people who drowned by falling into a pool

correlates with

Films Nicolas Cage appeared in



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Testing the significance of several variables at once

- ▶ When we have categorical variables with more than two categories, we replace one variable by two, or more, dummy-variables.
- ▶ How to test the significance of the original variable?
- ▶ Not *t*-test. That just tests one of the categories against the reference.
- ▶ Solution: Divide the variability in *Y* into different parts, depending on (groups of) different variables. ANOVA (ANalysis Of VAriance)
- ▶ This also works for testing several continuous variables at the same time.

Variance decomposition

- ▶ Null model with only β_0 : $Y_i \sim N(\beta_0, \sigma^2)$ with $\hat{Y}_i = \hat{\beta}_0 = \bar{Y}$ and

$$\hat{\sigma}^2 = s_0^2 = \frac{\sum_{i=1}^n (Y_i - \bar{Y})^2}{n - 1} = \frac{\text{SS}(\text{Total}_{\text{corrected}})}{n - 1}$$

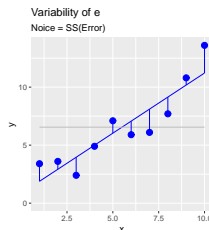
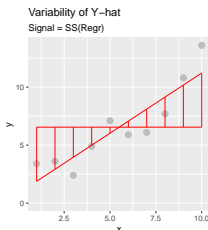
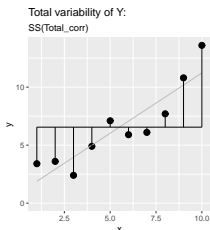
- ▶ How much of this variability in Y can be explained by the linear relationship with the x -variables?
- ▶ Full model: $Y_i \sim N(\mathbf{x}_i\boldsymbol{\beta}, \sigma^2)$ with $\hat{Y}_i = \mathbf{x}_i\hat{\boldsymbol{\beta}}$
- ▶ Idea: Divide Y_i into a noise part and a signal part:

$$Y_i = \underbrace{Y_i - \hat{Y}_i}_{e_i} + \underbrace{\hat{Y}_i}_{\text{signal}}$$

- ▶ Divide $\sum_{i=1}^n (Y_i - \bar{Y})^2$ into corresponding noise and signal parts.
If at least one $\beta_j \neq 0$, for $j = 1, \dots, p$, the signal should be stronger than the noise.

Variance decomposition (cont.)

$$\begin{aligned}
 \text{SS}(\text{Total}_{\text{corrected}}) &= \sum_{i=1}^n (Y_i - \bar{Y})^2 = \sum_{i=1}^n \underbrace{(e_i + \hat{Y}_i - \bar{Y})}_{Y_i}^2 \\
 &= \sum_{i=1}^n e_i^2 + \sum_{i=1}^n (\hat{Y}_i - \bar{Y})^2 - 2 \underbrace{\sum_{i=1}^n e_i (\hat{Y}_i - \bar{Y})}_{=0 \text{ see next slide}} \\
 &= \underbrace{\text{SS}(\text{Error})}_{\text{noise}} + \underbrace{\text{SS}(\text{Regr})}_{\text{signal}}
 \end{aligned}$$



(*) Proof of cross-term = 0

The Least squares estimation requires

$$\frac{\partial Q(\beta)}{\partial \beta_0} = \sum_{i=1}^n (Y_i - \mathbf{x}_i \beta) = 0 \Rightarrow \sum_{i=1}^n e_i = \sum_{i=1}^n (Y_i - \mathbf{x}_i \hat{\beta}) = 0.$$

Rewriting (see also Lecture 5)

$$\mathbf{X}\hat{\beta} = \underbrace{\mathbf{X}(\mathbf{X}'\mathbf{X})^{-1}\mathbf{X}'}_{=\mathbf{P}} \mathbf{Y} = \mathbf{P}\mathbf{Y}, \quad \mathbf{P}' = \mathbf{P}, \quad \mathbf{P}\mathbf{P} = \mathbf{P}$$

gives

$$\begin{aligned} \sum_{i=1}^n e_i \hat{Y}_i &= \mathbf{e}' \hat{\mathbf{Y}} = (\mathbf{Y} - \mathbf{X}\hat{\beta})' \mathbf{X}\hat{\beta} = (\mathbf{Y} - \mathbf{P}\mathbf{Y})' \mathbf{P}\mathbf{Y} = \\ &= (\mathbf{Y}' - \mathbf{Y}'\mathbf{P}') \mathbf{P}\mathbf{Y} = \mathbf{Y}' \mathbf{P}\mathbf{Y} - \mathbf{Y}' \underbrace{\mathbf{P}'\mathbf{P}}_{=\mathbf{P}} \mathbf{Y} = 0 \end{aligned}$$

and thus

$$\sum_{i=1}^n e_i (\hat{Y}_i - \bar{Y}) = \sum_{i=1}^n e_i \hat{Y}_i - \bar{Y} \sum_{i=1}^n e_i = 0 - \bar{Y} \cdot 0 = 0.$$

If H_0 is correct

If $H_0: \beta_1 = \dots = \beta_p = 0$ is correct then $E(Y_i) = \beta_0$ and $E(\hat{Y}_i) = E(\mathbf{x}_i\hat{\beta}) = \mathbf{x}_i\beta = \beta_0$.

- Using Y_1, \dots, Y_n with $\hat{\beta}_0 = \bar{Y}$ we know that

$$\hat{\sigma}^2 = \frac{\sum_{i=1}^n (Y_i - \bar{Y})^2}{n-1} = \frac{\text{SS}(\text{Total}_{\text{corrected}})}{n-1}$$

is an unbiased estimate of σ^2 (Basic statistics course).

- Using $\hat{Y}_1, \dots, \hat{Y}_n$ as a sample with $\hat{\beta}_0 = \bar{\hat{Y}} = \bar{Y}$ we get

$$\hat{\sigma}^2 = \frac{\sum_{i=1}^n (\hat{Y}_i - \bar{Y})^2}{p} = \frac{\text{SS}(\text{Regr})}{p} = \text{MS}(\text{Regr})$$

Note: the \hat{Y}_i are strongly dependent so we have to compensate by dividing by p in order to get an unbiased estimate (Advanced course in inference theory).

These two $\hat{\sigma}^2$ are only correct when H_0 is true. When H_0 is wrong they will be too large. We need another estimate that is always correct to compare with.

Full model

Using the full model with all the β -parameters we know that

$$\hat{\sigma}^2 = s^2 = \frac{\sum_{i=1}^n (Y_i - \hat{Y}_i)^2}{n - (p + 1)} = \frac{\text{SS(Error)}}{n - (p + 1)} = \text{MS(Error)}$$

is an unbiased estimate of σ^2 (Lecture 1). And this is true even if some (or all) of β_1, \dots, β_p happen to be 0!

Analysis of variances

The signal-to-noise comparison will use the two variance estimates $\hat{\sigma}^2 = \text{MS(Regr)} = \text{Signal}$ and $\hat{\sigma}^2 = \text{MS(Error)} = \text{Noise}$.

Chi-squared distribution, $\chi^2(f)$

If $Z_i \sim N(0, 1)$, $i = 1, \dots, n$ are independent then

$$\sum_{i=1}^n Z_i^2 \sim \chi^2(n)$$

If $Z_i \sim N(\mu, \sigma^2)$, $i = 1, \dots, n$ are independent then

$$\frac{1}{\sigma^2} \sum_{i=1}^n (Z_i - \bar{Z})^2 \sim \chi^2(n-1)$$

The parameter f is called the degrees of freedom.

F-distribution, $F(f_1, f_2)$

If $Z_1 \sim \chi^2(f_1)$ and $Z_2 \sim \chi^2(f_2)$ are independent then

$$\frac{Z_1/f_1}{Z_2/f_2} \sim F(f_1, f_2)$$

Distribution of Sums of Squares

We always have that

$$\frac{1}{\sigma^2} \text{SS}(\text{Error}) = \frac{1}{\sigma^2} \sum_{i=1}^n (Y_i - \hat{Y}_i)^2 \sim \chi^2(n - (p + 1))$$

If H_0 is true we also have that

$$\frac{1}{\sigma^2} \text{SS}(\text{Regr}) = \frac{1}{\sigma^2} \sum_{i=1}^n (\hat{Y}_i - \bar{Y})^2 \sim \chi^2(p)$$

Since they are independent it follows that, when H_0 is true,

$$\frac{\frac{1}{\sigma^2} \text{SS}(\text{Regr})/p}{\frac{1}{\sigma^2} \text{SS}(\text{Error})/(n - (p + 1))} = \frac{\text{MS}(\text{Regr})}{\text{MS}(\text{Error})} \sim F(p, n - (p + 1))$$

Global F-test

Test $H_0 : \beta_1 = \dots = \beta_p = 0$ vs $H_1 : \text{at least one } \beta_j \neq 0$,
 $j = 1, \dots, p$

- ▶ If H_0 is true then none of the x -variables are needed and the signal is just noise and $\frac{MS(\text{Regr})}{MS(\text{Error})} \approx 1$
- ▶ If H_0 is wrong a consistent part of the response variation is due to the signal and $\frac{MS(\text{Regr})}{MS(\text{Error})} \gg 1$
- ▶ If H_0 is true then:

$$F = \frac{MS(\text{Regr})}{MS(\text{Error})} \sim F(p, n - (p + 1)) \quad (\text{one-sided test})$$

and we can reject H_0 , in favour of H_1 at significance level α if

$$F > F_{\alpha, p, n-(p+1)}$$

Example: cabbage weight, soil type and fertilizer

Are any of the variables in the model necessary?

$H_0: \beta_1 = \beta_2 = \beta_3 = 0.$

In R: run the `summary()` for your model

On the last line you have

F-statistic: 5.721 on 3 and 26 DF, p-value: 0.003814.

This means that

- ▶ $F = \frac{MS(\text{Regr})}{MS(\text{Error})} = 5.721$
- ▶ The test statistic F follows an $F(3, 26)$ -distribution if H_0 is true.
- ▶ $P\text{-value} = P(F(3, 26) > 5.721) = 0.003814$ if H_0 is true.

Since $P\text{-value} < 0.05 = \alpha$ we should reject H_0 . Our data suggest that the model does contain at least one relevant covariate (it doesn't say which one!)

Partial F -test: testing a subset of parameters

Model: $Y_i = \beta_0 + \beta_1 x_{i1} + \dots + \beta_p x_{ip} + \epsilon_i$ where $\epsilon_i \sim N(0, \sigma^2)$.

Question: are k specific β -parameters $= 0$

(e.g. the k last: $\beta_{p-k+1} = \dots = \beta_p = 0$)?

Procedure:

- ▶ Estimate the full model with all $p + 1$ parameters:
 $SS(\text{Error}_{\text{full}})$ with $df = n - (p + 1)$.
- ▶ Estimate the reduced model with $p + 1 - k$ parameters:
 $SS(\text{Error}_{\text{reduced}})$ with $df = n - (p + 1 - k)$.
- ▶ Calculate the increase in $SS(\text{Error})$:
 $Q = SS(\text{Error}_{\text{reduced}}) - SS(\text{Error}_{\text{full}})$ with
 $df = n - (p + 1 - k) - (n - (p + 1)) = k$.
- ▶ Is this increase too large? Reject H_0 at $\alpha = 5\%$ if

$$F = \frac{Q/k}{s_{\text{full}}^2} > F_{\alpha, k, n-(p+1)}$$

Example: do we need the soil type(s)?

Compare the full model: $Y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i2} + \beta_3 x_{i3} + \epsilon_i$
against the reduced model $Y_i = \beta_0 + \beta_3 x_{i3} + \epsilon_i$.

Test $H_0: \beta_1 = \beta_2 = 0$ against $H_1: "$ $\beta_1 \neq 0$ and/or $\beta_2 \neq 0$ "

In R: `anova(model.reduced, model.full)` gives

$$F = \frac{Q/k}{s_{\text{full}}^2} = \frac{(28.427 - 18.405)/2}{18.405/26} = 7.08,$$

$$\text{P-value} = 0.0035$$

Since $F = 7.08 > F_{0.05, 2, 26} = 3.37$ or $\text{P-value} = 0.0035 < 0.05$ we should reject H_0 . Yes, we need the soil types in the model.

Important!

- ▶ Partial F-test for comparison between a large and a reduced model can only be performed if the models are **nested**. All variables in the reduced model must be present in the full model.
- ▶ Data used must be the same for both models.

Warning: Running `anova(model)` on just one model builds the ANOVA table sequentially using Type I Sums of Squares, assigning as much of the variability as possible to the x -variable that comes first in the `lm()`-function, then it assigns as much of the remaining variability as possible to the second x -variable, etc. **This is not what we want!**

We should use Type II in models without interaction and Type III in models with interaction. This will automatically be the case when using `anova(model.reduced, model.full)`.

Note:

- ▶ A t-test is for the significance of a single β -parameter, given all others in the model.
- ▶ An F-test tests several β -parameters at the same time but does not say **which** β_j 's are significantly $\neq 0$.
- ▶ An F-test is always one-sided because a larger model can never explain less than a reduced model.

R:

quantile $F_{\alpha, f_1, f_2} =$ `= qf(alpha, f1, f2, lower.tail = FALSE) = qf(1 - alpha, f1, f2),``P-value = pf(F, f1, f2, lower.tail = FALSE)`