

Data-analyysi R-ohjelmistolla

Tommi Härkänen

Terveyden ja hyvinvoinnin laitos (THL), Helsinki
E-mail: tommi.harkanen@helsinki.fi

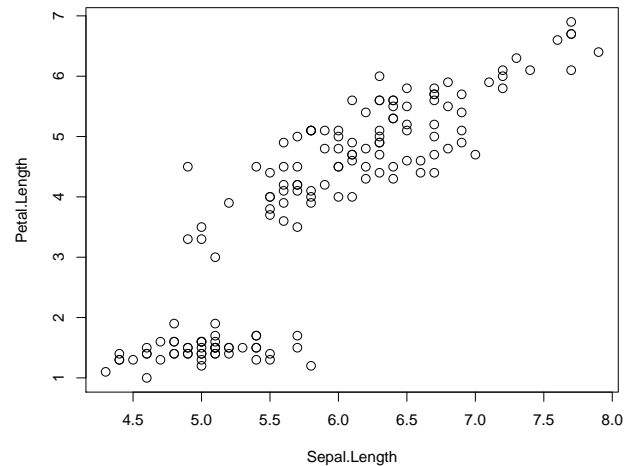
Helsingin yliopisto, 2.4.2014

Contents

Linear models

Association of continuous variables

Example: The iris data



Regression modeling

What is the average value of the outcome variable?

A researcher wants to know, what is the association of two (or more) continuous variables.

Simple questions:

- ▶ If the researcher measures e.g. sepal length, then what is the **average petal length**?
- ▶ How much does the petal length **change** on average, if the measured sepal length increases by 1 cm (unit of measurement)?

More complicated questions:

- ▶ Are the associations listed above different for **different species**?
- ▶ How well does the model **predict** petal length given sepal length (and possibly other variables)?

Regression modeling

Linear model for one explanatory variable (a.k.a **covariate** or independent variable) x_i for individual $i = 1, 2, \dots, n$ is often defined as

$$Y_i = \overbrace{\beta_0 + \beta_1 x_i}^{\text{expectation}} + \epsilon_i. \quad (1)$$

The **outcome** variable is Y_i

Regression coefficients are β_0 and β_1 :

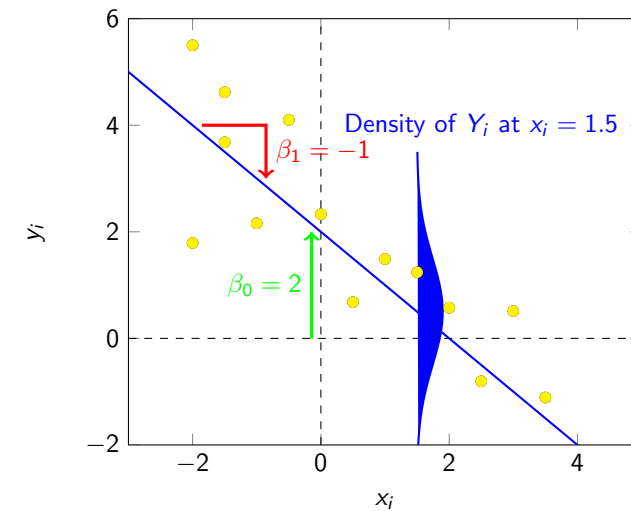
The **intercept** term β_0 controls the average level of the outcome values at $x_i = 0$. The expected value of the outcome is $\mathbb{E}[Y_i | x_i = 0] = \beta_0$.

The **slope** term β_1 controls the **association** of the outcome and the covariate. Note that if x_i increases by 1 unit, then the outcome value increases by β_1 on average.

Error term is ϵ_i , which is often assumed to be a normally distributed random variable with mean 0 and variance σ^2 .

Regression modeling

$Y_i = \beta_0 + \beta_1 x_i + \epsilon_i = 2 + (-1) \times x_i + \epsilon_i$ and $\epsilon_i \sim N(0, \sigma^2)$ where $\sigma^2 = 1$.



Linear regression modeling in R

The basic command is `lm`:

```
lm(formula, data, subset, ...)
```

Some of the most important options are

formula The model description as a formula: `outcome ~ terms` where **terms** are the covariates separated by '+' and their interactions defined using '*' or ':'.

data Optional data frame, list or environment name.

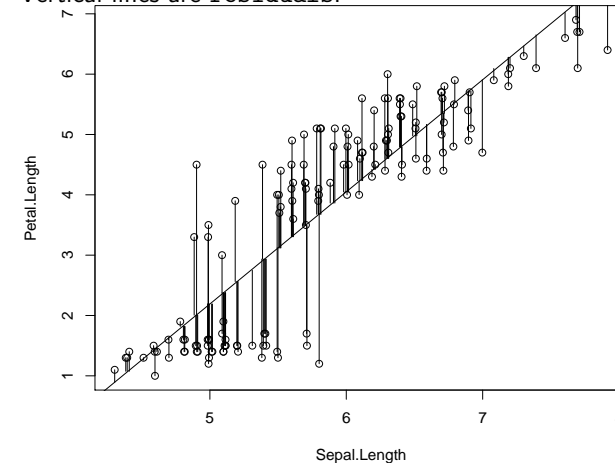
subset Optional vector specifying a subset of observations.

Example:

```
lm(Petal.Length ~ Sepal.Length, data = iris, subset = Species ==
  "setosa")
##
## Call:
## lm(formula = Petal.Length ~ Sepal.Length, data = iris, subset = Species ==
##   "setosa")
##
## Coefficients:
## (Intercept) Sepal.Length
##          0.803          0.132
```

Observed vs. predicted values

Vertical lines are residuals.



Observed vs. predicted values

Separate analyses for each species

