

Model validation

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- ▶ Normal distribution
- ▶ Linear models ...

Recap

Recall:

$$y_i = \alpha + \mathbf{x}_i \boldsymbol{\beta} + \epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

- ▶ the systematic component: the regression line
- ▶ the random component: leftover stuff

But more on that later

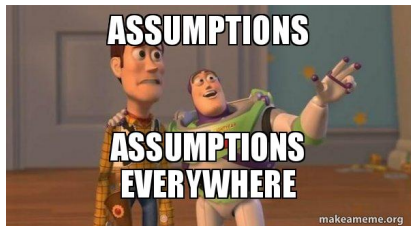
Linear regression assumptions

$$y_i = \alpha + \sum_{k=1}^p x_{ik} \beta_k + \epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

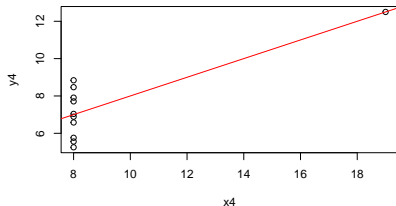
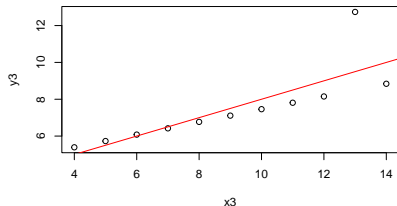
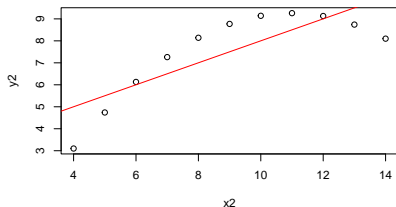
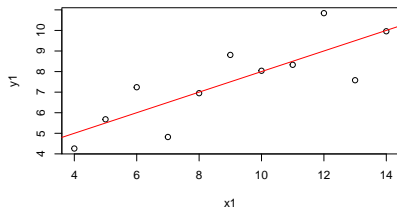
- ▶ Linearity
- ▶ Normality: normally distributed errors
- ▶ Homoscedasticity: same variance for all errors
- ▶ Independence of errors
- ▶ Lack of perfect multicollinearity ($\mathbf{x}_1 \neq \mathbf{x}_2$)
- ▶ No outliers
- ▶ No error in the explanatory variables

Violating assumptions

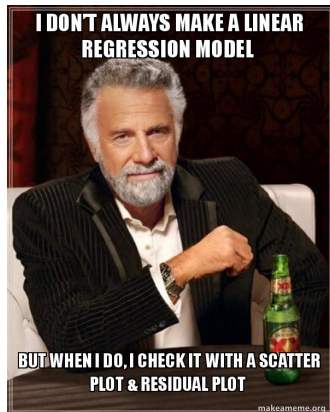
- ▶ Linearity: Eek
- ▶ Normality: cannot trust tests and confidence intervals
- ▶ Homoscedasticity: cannot trust tests and confidence intervals
- ▶ Independence of errors: Eek
- ▶ Lack of perfect multicollinearity
- ▶ No outliers: biased parameter estimates



Anscombe's quartet



All of these regressions have the same parameter estimates



Errors vs. Residuals

The true model:

$$y_i = \alpha + \sum_{k=1}^p x_{ik} \beta_k + \epsilon_i \sim \mathcal{N}(0, \sigma^2)$$

The fitted model:

$$y_i = \hat{\alpha} + \sum_{k=1}^p x_{ik} \hat{\beta}_k + \hat{\epsilon}_i$$

► $\epsilon_i = y_i - \alpha - \sum_{i=1}^n x_{ik}\beta$

► $\hat{\epsilon}_i = y_i - \hat{\alpha} - \sum_{k=1}^n x_{ik} \hat{\beta}$

Residuals are the **observed** deviation from the regression line.

In R

```
model <- lm(y~x)
residuals(model) # estimated random part
fitted(model) # estimated systematic part
```

Standardised residuals

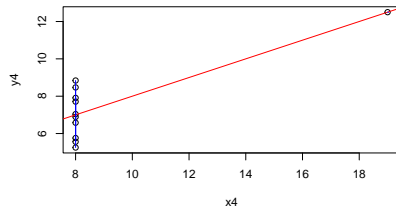
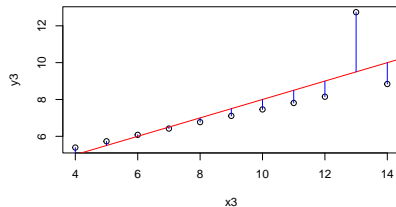
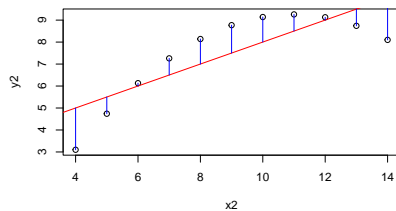
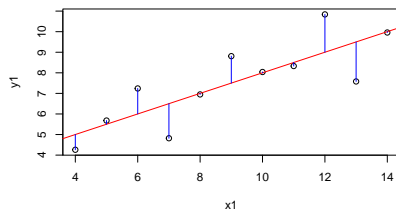
$$\frac{\hat{\epsilon}_i}{s\sqrt{1-h_i}}, \quad \text{where } h_i \text{ is "influence"} \quad (1)$$

Studentized residuals

- ▶ omit observation i
- ▶ re-compute residual variance
- ▶ calculate standardised residual (with hat matrix)

`rstand` or `rstudent`

Anscombe's quartet



Residual plots

Residual plots should have no discernible patterns.

- ▶ No curvature
- ▶ No shapes whatsoever
- ▶ No outliers
- ▶ No change in variance

This can be difficult with few observations.

Our toolbox

- ▶ Residuals versus fitted: $\hat{\epsilon}_i$ vs. $\hat{\mu}_i$
- ▶ Observed versus theoretical quantiles:
 $Pr(X \leq y_i)$ vs. $Pr(X \leq x)$
- ▶ Cook's distance plot: $\hat{\epsilon}_i$ vs. $\hat{\mu}_i$
- ▶ Standardized residuals vs. fitted (Scale-location)
- ▶ Leverage and influence

Leverage: hat matrix

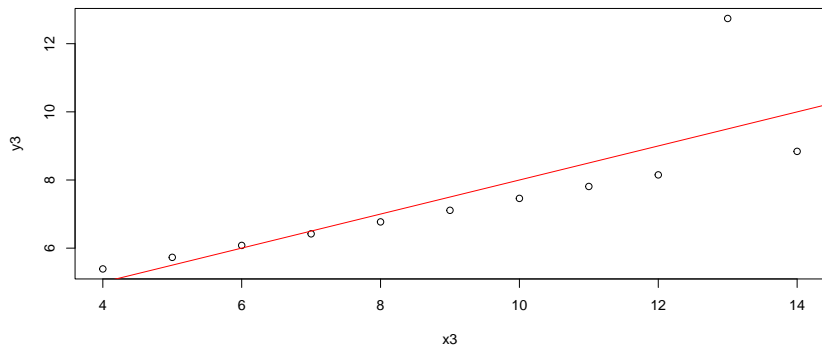
$$\hat{\mu}_i = \hat{\alpha} - \sum_{i=1}^n x_{ik} \hat{\beta}$$

but we can also write:

$$\hat{\mu} = \mathbf{H}\mathbf{y}$$

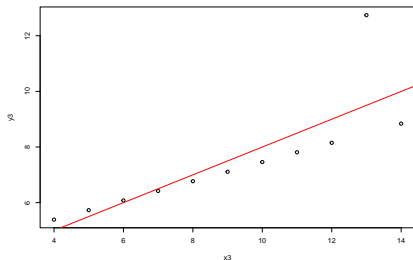
- ▶ **H** is the “hat” matrix
- ▶ **H** = $\mathbf{X}(\mathbf{X}^\top \mathbf{X})^{-1} \mathbf{X}^\top \mathbf{y}$
- ▶ Its diagonal gives “leverage” and is between 0 and 1
- ▶ Leverage: observation that is far from others on the explanatory variable

Influence



An observation that has undue impact on the regression line.

Outliers



- ▶ Outliers are observations far away from the regression line
- ▶ Can be erroneous
- ▶ Might be real data
- ▶ Remove with caution
- ▶ Better: sensitivity analysis

Cook's distance

Cook's distance: how much the fitted values change if we remove an observation

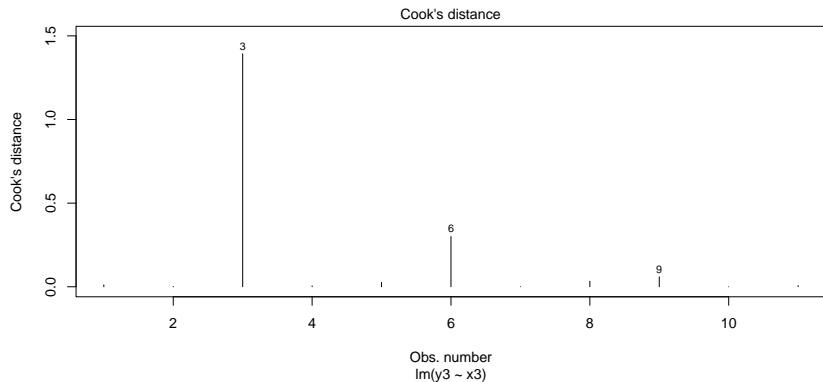
$$D_j = \frac{\sum_{i=1}^n (\hat{y}_i - \hat{y}_{i(-j)})^2}{\sigma^2}$$

- ▶ \hat{y}_i - prediction
- ▶ $\hat{y}_{i(-j)}$ - prediction for model with data point j removed
- ▶ σ^2 - residual variance

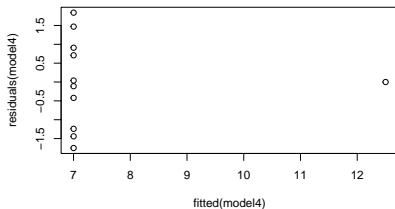
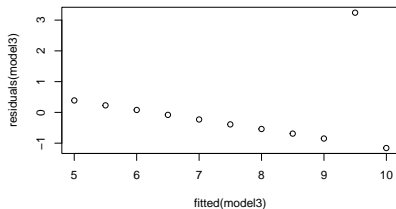
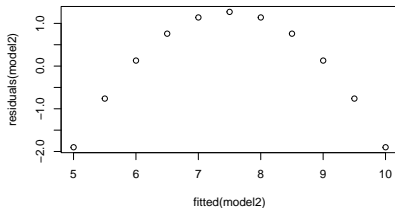
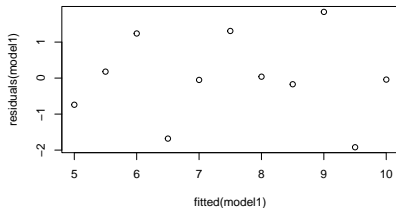
What is influential?

Large values of D_i mean a large influence

Usually $D_i > 1$, or $4/n$

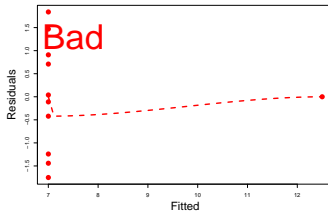
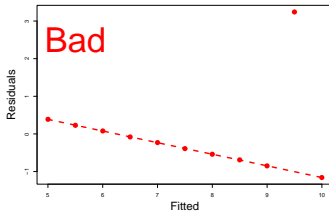
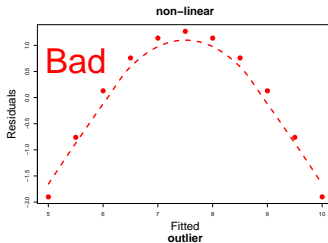
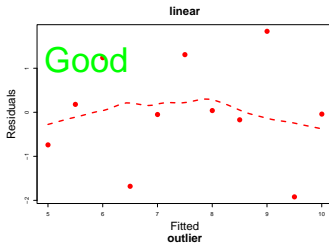


Anscombe's quartet: residuals

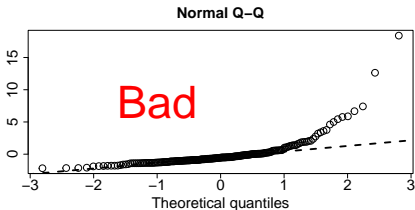
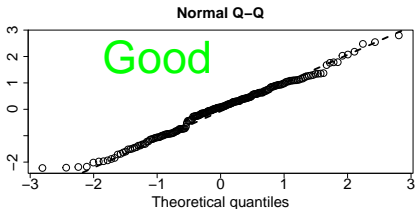


What do you see?

Residual diagnostics: residuals vs. fitted

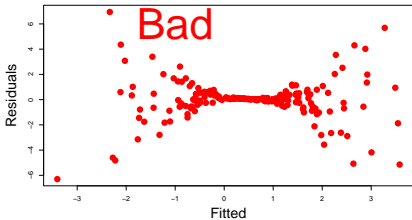
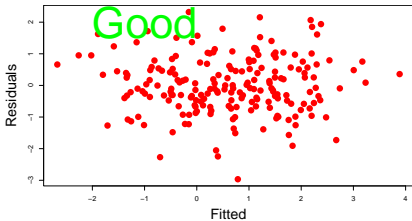


Residual diagnostics: normality with QQ-plot



1. compare residuals to theoretical quantiles
2. points should follow the line

Residual diagnostics: constant variance



1. compare residuals to fitted
2. points should be randomly distributed

Addressing assumption violations

- ▶ Curvature? Add quadratic term
- ▶ Outlier? Sensitivity analysis (if not error)
- ▶ Normality? GLM
- ▶ Constant variance? GLM

Alternative: data transformation

Use a Generalised linear model instead

Summary

Now we have a valid model, we can check **how** good it is (more on that tomorrow)