

What on earth is going on with my linear models??!

March 8, 2018

1 Linear model, reminder

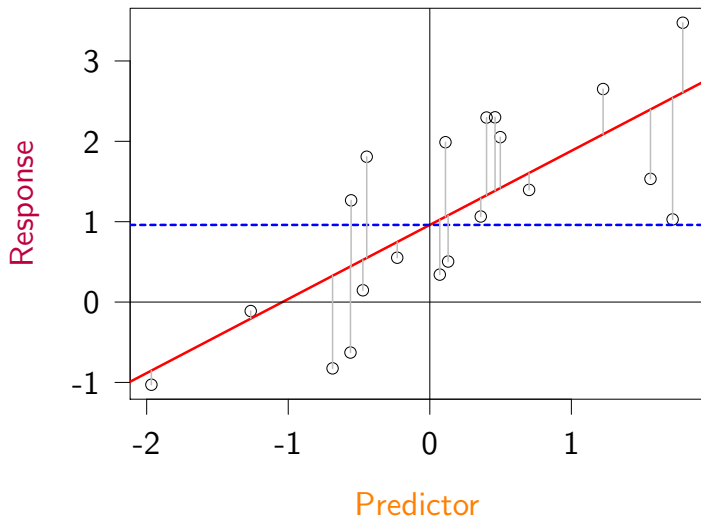
2 Checks and prediction

3 Cures

4 Bonus fun

A simple linear model

$$\text{Response} = \text{Intercept} + \text{Slope} \times \text{Predictor} + \text{Error}$$



A simple linear model

$$\text{Response} = \text{Intercept} + \text{Slope} \times \text{Predictor} + \text{Error}$$

In R:

```
lm(response ~ 1 + predictor1 + predictor2, data=data)
# equivalent to
lm(response ~ predictor1 + predictor2, data=data)
```

- Intercept can be explicit or implicit
- Can remove intercept with $\dots \sim 0 + \dots$
- Error is implicit
- Feed the option `data=` to keep code short, reliable and flexible
- Order of predictors do not matter

1 Linear model, reminder

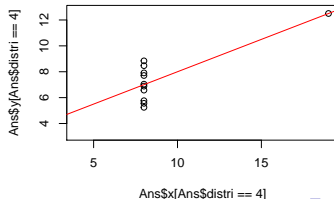
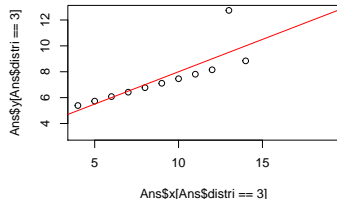
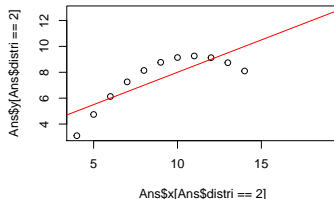
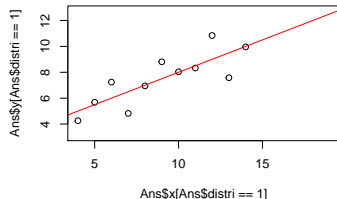
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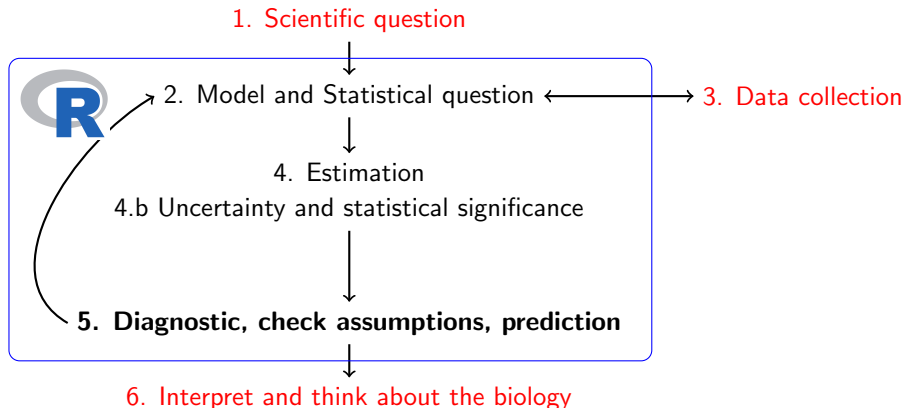
4 Bonus fun

Why we need checks: summary(lm) isn't enough

```
Ans <- read.csv(file = "Anscombe.csv")
```



General approach



Why we need checks: missing a relationship

```
forprediction <- read.csv(file = "forprediction.csv")
```

Does "predictor" predict "obs"?

Why we need checks: missing a relationship

```
forprediction <- read.csv(file = "forprediction.csv")
```

Does "predictor" predict "obs"?

```
summary(lm(obs ~ 1 + predictor, data=forprediction) )
```

Why we need checks: missing a relationship

Does "predictor" predict "obs"? Apparently not:

```
summary(lm(obs ~ 1 + predictor, data=forprediction) )
```

Call:

```
lm(formula = obs ~ 1 + predictor, data = forprediction)
```

Residuals:

Min	1Q	Median	3Q	Max
-2.1962	-0.5326	0.1378	0.5785	1.8664

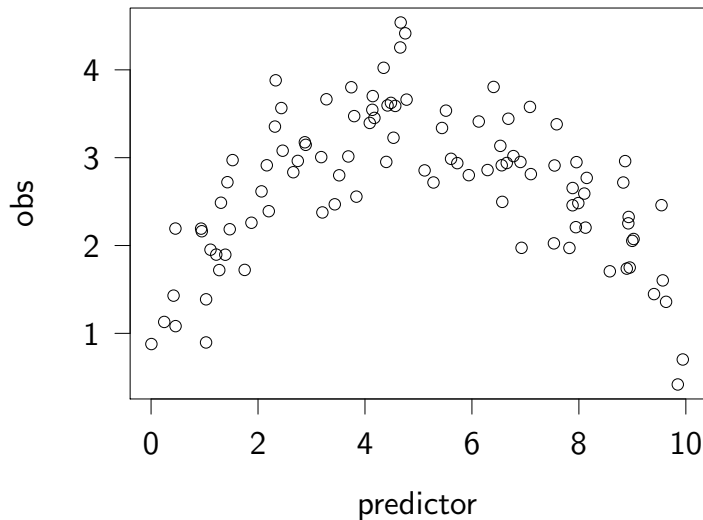
Coefficients:

	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	2.72530	0.16953	16.076	<2e-16 ***
predictor	-0.01129	0.02956	-0.382	0.703

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 0.8382 on 98 degrees of freedom

Why we need checks: missing a relationship



How to check?

```
m0 <- lm(obs ~ 1 + predictor, data=forprediction)
summary(m0)
```

Call:

```
lm(formula = obs ~ 1 + predictor, data = forprediction)
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Residuals:

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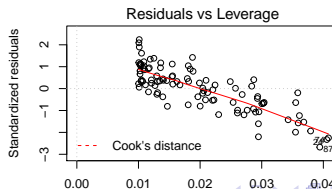
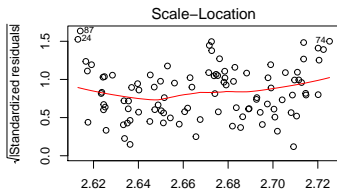
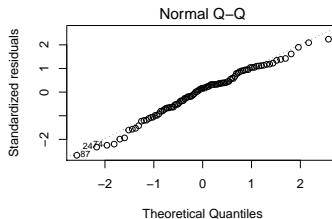
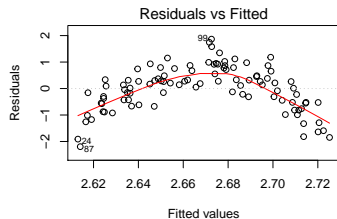
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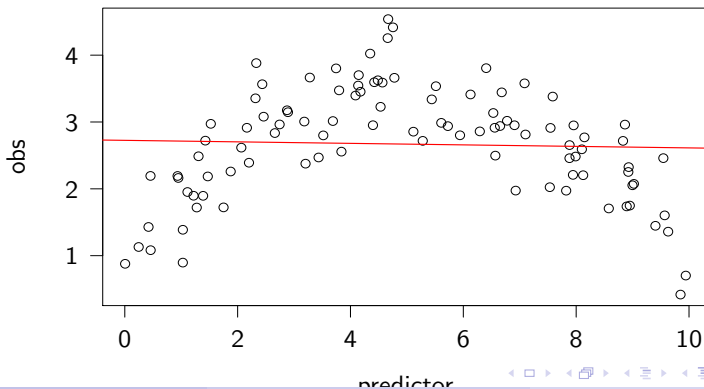
```
par(mfrow=c(2,2))  
plot(m0)
```



How to check?

```
m0 <- lm(obs ~ 1 + predictor, data=forprediction)
```

```
setPar()  
plot(x=forprediction$predictor, y=forprediction$obs, xlab="predictor",  
     abline(m0, col="red", lwd=3) #simple prediction, without SE
```



Check checklist

- **Visualize your data**
- Residual in summary(): are they symmetrical?
- plot(lm):
 - ① trend residual/fitted?
 - ② Normal residuals?
 - ③ trend in residual variance?
 - ④ outliers?
- Predictions: range and biological meaning

Practice lm() with parasites

What explains variation in parasitic load?

You collected ecto-parasites on some furry large mammals at three locations. Parasites break easily when we collect them and are impossible to count, so we decide to measure parasitic load as their mass. **Why do some mammals have larger parasitic load?**

Practice `lm()` with parasites

What explains variation in parasitic load?

You collected ecto-parasites on some furry large mammals at three locations. Parasites break easily when we collect them and are impossible to count, so we decide to measure parasitic load as their mass. **Why do some mammals have larger parasitic load?**

- Load the `Para.csv` data (don't forget: `str()`, `summary()`, `plot()`...)
- Model `Parasite_Mass` using `lm()`
- Find what variables predict `Parasite_Mass`
- How good are your models? Assumptions? Prediction?
- What biological interpretation can you imagine?

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Multiple regression

Transformations

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Not necessarily wrong, but typical interpretation assumes:

- Linear combination of parameters (including transformation, polynoms, interactions. . .)
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- Homoscedasticity (constant error variance)
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Risk: Poor predictions
- Homoscedasticity (constant error variance)
Risk: Over-optimistic uncertainty, unreliable predictions
- Independence of error
Risk: Bias and over-optimistic uncertainty

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Extra exercises

General R coding

- 1 What is the fastest way to get row averages in a data-frame?
- 2 Create a function called `colVars`, like `colMeans` but for variance
- 3 Create nice plots to visualize iris data (ideally journal-quality)

Linear models

- 1 Load `Cdata.csv`, fit models of y predicted by x_1 and x_2 , or x_2 and x_3 . Something is weird, what is going on? What to do?
- 2 For model that can be fitted with `t.test`, `aov`, and `lm`, is one of the function faster?
- 3 Write your own code to obtain a prediction from a `lm` (that is, a simpler version of the `predict` function), with confidence interval. (extra toughness: do it using the matrix formulation of the analytical solution to a linear model)

What do you want to learn about?

Topics

