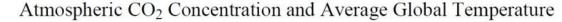
Lecture 8

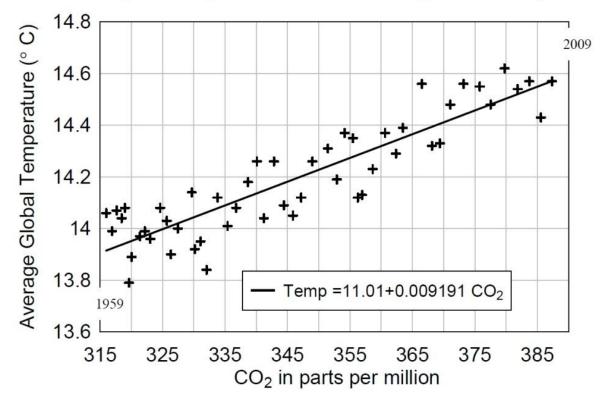
Correlation and linear regression

相关与线性回归

Linear regressions

- It is the simplest case of statistical modelling
- Linear regression is a very important and popular model
 - global warming
 - · trends in human longevity
 - etc.
- Despite all advances in statistical and computational methods, linear regression remains the option number 1 when trying to model or predict a variable as a function of another





变量之间的关联

Association between variables

变量可以在不同的级别相关联

- Variables may are associated at different levels
 - Covid-19 patients have been infected with coronavirus (always)
 - height is associated with weight (frequently)
 - fizzy drinks are associated with throat cancer (rarely)
 - ethnicity is not associated with IQ (ever)

相关分析 量化了变量之间的关联 1. 确定变量是否关联 2. 确定相关性是正还是负 3. 量化关联水平

- Correlation analysis quantifies associations between variables
 - 1. determine whether variables are associated
 - 2. establish whether correlation is positive or negative
 - 3. quantify levels of association

Pearson correlation

是两个变量之间线性相关性的度量=两个变量的变化相关联的程度

- Pearson (or linear) correlation is a measure of linear dependence between two variables
 - =the degree to which change in variable 1 is associated with change in variable 2
- How do we measure association?
 - take a sample and two variables:
 - *x* = male height
 - y = male weight
 - calculate average weight <u>X</u> and height <u>Y</u>
 - For each case *i* in the sample, calculate
 - difference between its height and average height

$$\bullet = (x_i - \underline{X})$$

difference between its height and average weight

$$\bullet = (y_i - \underline{Y})$$

高度与平均高度之差

身高与平均体重之差

Covariance

The product of the two quantities 两个量的乘积

$$(x_i - \underline{X})^*(y_i - \underline{Y})$$

gives an idea of how height and weight covary in one individual

样本中所有这些产品的平均值是两个性状的协方差

• The average of all those products in a sample is the *covariance* of the two traits

$$cov_{x,y} = \sum_{i=1}^{\infty} \frac{(x_i - X)(y_i - Y)}{n}$$

Exercise:

Manually calculate the covariance of hight and weight in the sample of three

cases Individual	height	weight	
1	173	61	
2	182	76	
3	165	58	

Pearson correlation

协方差收到变量的尺度和计量单位的影响

 But covariance is affected by scale and measurement units of variables

将协方差除以两个变量的标准差,得到皮尔逊相关系数r

 If we divide covariance by the standard deviations of the two variables, we obtain the Pearson correlation r

$$r = \frac{cov_{x,y}}{\sigma_x \sigma_y}$$

- i.e., correlation is the *standardised* covariance of *x* and *y*
 - for this reason, it varies between -1 and 1
 - r=1 means absolute association r=1表示正向绝对关联(实验中不可能出现的 , 如果出现就是 r=-1 means absolute (but inverse) association r=-1表示负向绝对相关 数据错了)
 - r=0 means no association r=0表示无关联

Exercise:

Now manually calculate the Pearson correlation between height and weight

Individual	height	weight
1	173	61
2	182	76
3	165	58

Significance test of correlation

- But correlation may or may not be significant 相关性可能显著可能不显著
 - as in the case of differences between means,
 - sample may be too small etc.
- We want to test whether the two variables are significantly correlated
 - null hypothesis: r=0 (no correlation) 零假设: r=0 无相关性

假设xy是正态分布的,定义一个t检验,检验值是r=0

• Parametric correlation test: we assume that x and y are normally distributed and define a t-test where test value is a correlation of r = 0

$$t = \frac{r-0}{sem} = r \frac{\sqrt{n-2}}{\sqrt{1-r^2}}$$

- Correlation test calculates probability that t is significantly different from 0
 - since this is a t-test, look for t < -1.96 or t > 1.96 for a significant difference

Significance test of correlation

- Example: is newborn head circumference and newborn weight (Swedish Birth Record) significantly correlated?
 - null hypothesis = no correlation (*r*=0)
 - File SBR

```
> cor.test(SBR$size, SBR$head)
Pearson's product-moment correlation
data: SBR$size and SBR$head
t = 319.6791, df = 186873, p-value < 2.2e-16
alternative hypothesis: true correlation is not equal to 0
95 percent confidence interval:
0.5916467 0.5975088
sample estimates:
cor
0.5945857
```

- Interpreting output: which are the three questions to ask?)
 - t= 319.7 p值接近零,拒绝零假设,两者是相关的
 - $P \sim 0$ => correlation is not zero; variables ARE correlated
 - correlation is positive: head size increases with weight 相关性是正的
 - association (r=0.59) is relatively strong 相关性r=0.59,相对较强

斯皮尔曼相关

Spearman's correlation ρ (rho)

代替皮尔逊相关的非参数检验办法

 = a nonparametric (rank) test alternative to Pearson's correlation

• To be used when

当样本量过小或样本变量不符合正态分布

- sample size is small
- distribution of variables is not normal

• Procedure:

- ranks the two variables 对两个变量进行排序
- replaces values with ranks 将值替换成等级,如果x和y的等级是相关的
- calculates Pearson correlation between the two rank distributions 那两个变量之间就是相关的(X是一级,Y也是一级)

Spearman's correlation ρ (rho)

- Running Spearman's correlation:
 - > cor.test(variable 1, variable 2, method="spearman")
- File *Brains2*: brain structures
 - Small sample of ape species (n=18)
 - what is the correlation between prefrontal white matter and prefrontal grey matter?

两个变量之间存在显著的、正的、强的关联

Conclusion: significant, positive, strong association between the two variables

Exercise:

Calculate the corrections between

- Lifespan and schooling
- Lifespan and income
- Income and schooling

using the full HDR2011 dataset

Which test do you use? Pearson or Spearman correlation?

hist(HDR2011\$lifespan)
hist(as.numeric(HDR2011\$schooling))
hist(as.numeric(HDR2011\$income))

Linear equation and linear regression

The linear equation

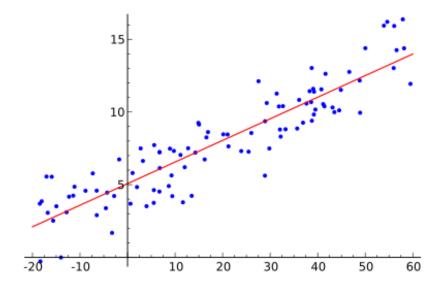
$$y = a + bx$$

relates variables y and x on the Cartesian plane

 Simple linear regression uses the linear equation to model (= predict) the dependent variable y from the independent variable x

$$y = a + bx + \varepsilon$$

- a = intercept 截距
 - where line crosses y axis
- *b* = slope or regression coefficient 斜率或回归系数
 - change in y per unit change in x
- ε = residual error 剩余误差,观察Y和预测Y之间的差异
 - difference between observed y and predicted y



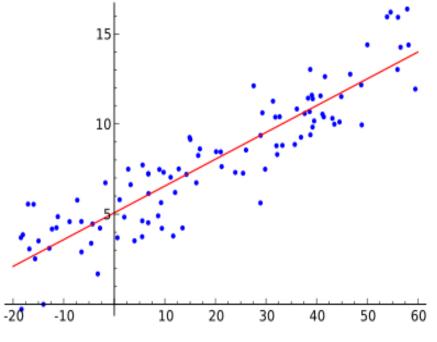
Estimation of linear regression

- Method of *least squares* estimates the 'best line' across sample of (x, y) points
- Best line is the one that minimises sum of squared differences (residuals) between observed y and predicted y:
 - SSres = Σ (observed y_i predicted y_i)² = $\Sigma(y_i - (a + bx_i))^2$ (since predicted y = a + bx)
 - best line always includes point (X, Y),
 where X = mean x, Y = mean y

最佳线过(x,y), xy是平均值

• Minimising SSres:

- Properties of the solution:
 - method rotates line around mean values (X, Y) to find combination of intercept and slope that reduces the sum of residuals
 - average residual = 0 平均残差=0

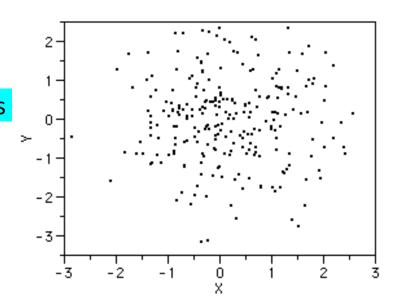


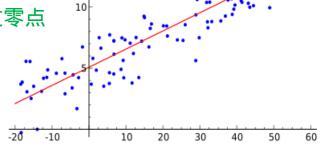
Significance of regression: slope test

- As in the case of means, proportions and correlations, significance of regression must be tested
- Key test is whether slope b is significantly different from 0
 - If b = 0, there is no linear relationship between variables! (i.e. there is no regression; slope not different from 0; 'best line' is horizontal)
 - A horizontal line (b=0) means that x has no effect on y
 - you cannot predict y from x 如果b=0,则变量之间不存在线性关系表示x对y没有影响,不能从x预测y
- We use a *t*-test for slope b

$$t = \frac{b-0}{sem(b)} = \frac{b}{sem(b)}$$

- We test whether b/sem(b) is within a 95% CI around the tested slope b=0
 - null hypothesis: b=0 (=no regression) 零假设: b=0(无回归)
- Intercept is also tested, but result is less important =0仍会得到回归,但不会过零点
 - If a=0, you still get a regression, but it does not cross origin (0,0)
 - Therefore, the regression test is the slope test!





Running linear regression in R

```
Is the amount of white matter in brains affected by the amount of grey matter?
 To run regression: function Im()
> Im(Brains$BrWhite ~ Brains$BrGray) <sub>V~X</sub>
or
> Im(BrWhite ~ BrGray, data = Brains)
or create an object
 > brainreg <- Im(Brains$BrWhite ~ Brains$BrGray)
Always run command summary either on Im command or object
> summary(Im(Brains$BrWhite ~ Brains$BrGray))
                                                             用smmary函数
or
> brainreg <- Im(Brains$BrWhite ~ Brains$BrGray)
> summary(brainreg)
```

Regression statistics: residuals

> summary(brainreg)

Call:

Im(formula = Brains\$BrWhite ~ Brains\$BrGray)

Residuals:

Min 1Q Median 3Q Max -25.367 -6.760 0.504 4.675 35.780

Coefficients:

Estimate Std. Error t value Pr(>|t|)
(Intercept) -1.44510 3.91407 -0.369 0.714
Brains\$BrGray 1.21928 0.03901 31.258 <2e-16 ***

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

Residual standard error: 13.72 on 33 degrees of freedom

Multiple R-squared: 0.9673, Adjusted R-squared: 0.9663

F-statistic: 977 on 1 and 33 DF, p-value: < 2.2e-16

Residuals: 根据定义,平均值=0,中位数应该是0

- mean=0 (by definition)
 - median should be ~0

如果第一个和第三个四分位数,或者最小和最大残差在数量上相差太大,则x和y之间的关系可能不是线性的可能存在异常值,有很多处理残差的技术(此处未提及)

- if 1st and 3rd quartile, or min and max residuals are too different in magnitude, relationship between x and y may not be linear
 - there may be outliers; many techniques to deal with residuals
 - (not addressed here)

Regression statistics: intercept

```
> summary(brainreg)
Call:
Im(formula = Brains$BrWhite ~ Brains$BrGray)
Residuals:
  Min
          1Q Median
                         3Q
                            Max
-25.367 -6.760 0.504 4.675 35.780
Coefficients:
               Estimate Std. Error t value Pr(>|t|)
(Intercept)
               -1.44510
                           3.91407 -0.369
                                                0.714
                                              <2e-16 ***
Brains$BrGray
               1.21928
                           0.03901 31.258
Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
Residual standard error: 13.72 on 33 degrees of freedom
Multiple R-squared: 0.9673, Adjusted R-squared: 0.9663
F-statistic: 977 on 1 and 33 DF, p-value: < 2.2e-16
```

Intercept test: 零假设: =0

• Null hypothesis: *a*=0

• t = -0.37

• *P*=0.714

Conclusion:

- a not different from 0
 - = curve goes through the origin
 - (as expected in this case)

与零无差异=曲线过原点

Regression statistics: coefficient

> summary(brainreg)

Call:

Im(formula = Brains\$BrWhite ~ Brains\$BrGray)

Residuals:

Min 1Q Median 3Q Max -25.367 -6.760 0.504 4.675 35.780

Coefficients:

Estimate Std. Error t value Pr(>|t|) (Intercept) -1.44510 3.91407 -0.369 0.714

Brains\$BrGray 1.21928 0.03901 31.258 <2e-16 ***

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Residual standard error: 13.72 on 33 degrees of freedom

Multiple R-squared: 0.9673, Adjusted R-squared: 0.9663

F-statistic: 977 on 1 and 33 DF, p-value: < 2.2e-16

Slope test:

- null hypothesis: *b*=0
- *t*-statistic=31.3
- P~0

斜率b明显不同于0

Conclusion:

- slope b is significantly different from 0
- b > 0: there is a positive effect of grey matter volume on white matter

Interpretation

 an extra gram of grey matter in primate brains predicts an extra 1.219 g of white matter

<u>IMPORTANT</u>

- Slope test is the regression test!
 - regression of white matter on grey matter IS significant
 - =we have a significant regression

我们有一个显著的回归

Confidence intervals

- Function confint calculates 95% confidence intervals of a and b estimates
 - Significant *b* => 95% CI excludes *b*=0

95%的置信区间不包括b=0

- b is significantly different from 0
 - regression is significant

可视化回归

Visualising regression

首先绘制y和x的关系图

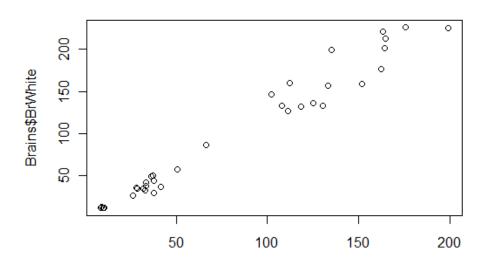
First plot y against x

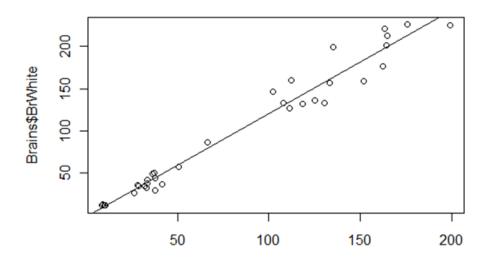
>plot(Brains\$BrWhite ~ Brains\$BrGray)

- Now plot line from the linear model
 - save your model as an object (in this case, brainreg)
 - plot regression line with command abline (=line defined by parameters a=intercept and b=slope)

用abl i ne绘制回归线

>brainreg <- lm(Brains\$BrWhite ~ Brains\$BrGray)
>abline(brainreg)





Goodness of fit

两条回归线可能是显著的,但他们与观测数据的匹配程度可能不同=模型的拟合度

- Two regression lines may be significant, but they may differ in how closely they match observed data
 - = the 'goodness of fit' of the model

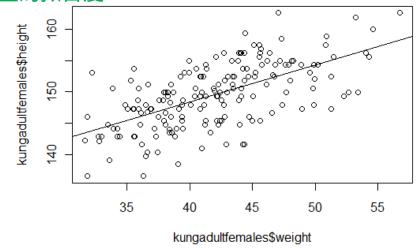
这反映了变量之间的线性关系=在回归线附近扩散

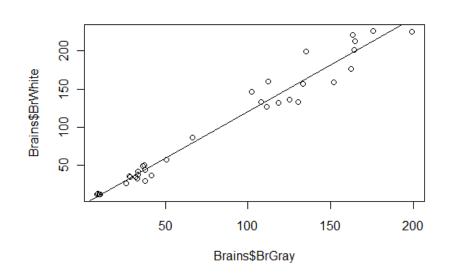
- This reflects how linear the relationship between the variables is
 - = dispersal around the regression line

拟合优度的主要测量是基于方差分析=R方

 Main measure of 'goodness of fit' is based on a generalisation of analysis of variance (ANOVA)

= (Multiple) R²



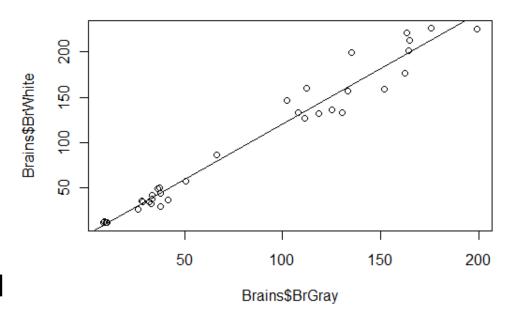


Generalised ANOVA

ANOVA可以用于计算模型的决定系数

- ANOVA can be used to calculate a coefficient of determination (COD) of a model
- COD is the fraction of variance of Y explained by model (=by the independent variable X)
- COD is estimated after partition of total variance into:
 - S = sum of squares explained by model 模型解释的平方和 = squared differences between predicted y and Y (general Y mean)
 - R = residual sum of squares 残差平方和=观测y与预测y之间的平方差 = squared differences between observed y and predicted y

```
COD = \frac{S = sum \ of \ squares \ explained \ by \ model}{S + R = total \ sum \ of \ squares}
```



Goodness-of-fit

> summary(brainreg)

Call:

Im(formula = Brains\$BrWhite ~ Brains\$BrGray)

Residuals:

```
Min 1Q Median 3Q Max -25.367 -6.760 0.504 4.675 35.780
```

Coefficients:

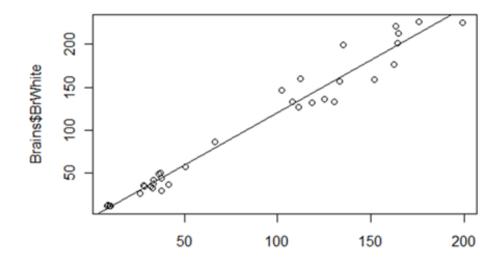
	Estimate	Std. Error	t value	Pr(> t)
(Intercept)	-1.44510	3.91407	-0.369	0.714
Brains\$BrGray	1.21928	0.03901	31.258	<2e-16 ***

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

Residual standard error: 13.72 on 33 degrees of freedom

Multiple R-squared: 0.9673, Adjusted R-squared: 0.9663

F-statistic: 977 on 1 and 33 DF, p-value: < 2.2e-16



- Back to our summary table:
- In linear regression analysis, COD is called Multiple R²

COD被称为multiple R方

- Multiple R squared = R^2 = 0.9673
- Very high value! Almost all variance in y can be predicted by x through the regression

很高的值,y中几乎所有的方差都可以通过回归用x预测

R^2 and r^2

在线性回归中,决定系数是两个变量之间的皮尔逊相关系数的平方

• In linear regression, the coefficient of determination is the square of the Pearson correlation coefficient between the two variables

$$R^2 = r^2$$

Calculating Pearson correlation r between x and y:

> cor(Brains\$BrWhite, Brains\$BrGray)
[1] 0.9835284

皮尔逊系数r可能是正的也可能是负的 一平方就和R一样了

And its square:

- > (cor(Brains\$BrWhite, Brains\$BrGray))^2
 [1] 0.9673282
- Squared Pearson coefficient = r^2 = COD = same value as R^2

r is the standardised regression slope

- if x and y are expressed in standard deviation units (z-scores), regression slope is the Pearson coefficient r
 - if correlation is perfect (r=1), z-scores of x and y are the same for all cases
 - if there is no correlation, result is *r*=0 (horizontal line) 如果相关性是完美的(r=1),则x和y的z分数对于所有情况都是相同的如果不存在相关性,则结果是r=0(水平线)

Summary

To create a linear regression model:

绘制变量y和x并且目测检查数据:看起来是否有线性关系?

- Plot variables y and x and visually inspect data
 - Does it seem that there is a linear relationship?

回归斜率的显著性检验,斜率显著表示线性模型有效

- Test significance of regression slope;
 - significant slope means linear model is valid
- If slope is significant, write down model y = a + bx; interpret meaning of intercept and slope
- Report confidence intervals of slope b and goodness-of-fit R²

Exercises

Predicting !Kung adult male weight from height (file 'Kungadultmales')

y是weight, x是height

• Let's say you want to predict body weights of !Kung men from their heights. What is the dependent variable y?

yes

- Plot variables y against x
 - does the relationship look linear? 0. 429**0. 5

```
plot(weight ~ height, data=Kungadultmales)
reg1 <- Im(weight ~ height, data=Kungadultmales)
summary(reg1)
0.429**0.5
cor.test(Kungadultmales$weight, Kungadultmales$height)
abline(reg1)</pre>
```

Run a linear regression of weight on height

- is the regression significant? 看b=0否
- how much of variance in data is explained by the model?
- what is the correlation between weight and height? multiple r-square开根号
- what is the final model? Write it as a line equation y = a + bx
- Add regression line to points
- Based on your model, what is the predicted weight of a !Kung man whose height is 165 cm?

数据中有多少方差<mark>是</mark> 中横刑解释的?