# Regression

Part 1: Simple linear regression

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# Aims and Objectives

- Understand linear regression with one predictor
- Understand how we assess the fit of a regression model
  - Total Sum of Squares
  - Model Sum of Squares
  - Residual Sum of Squares
  - $-R^2$
  - F

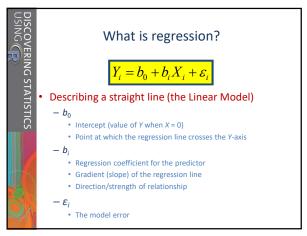
Know how to perform linear regression and interpret the model

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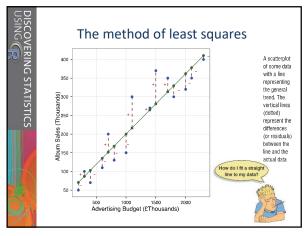
# What is regression?

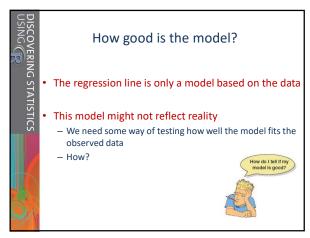
- A way of predicting the value of one variable from another
  - It is a hypothetical model of the relationship between two variables
  - The model used is a linear one
  - Therefore, we describe the relationship using the equation of a straight line

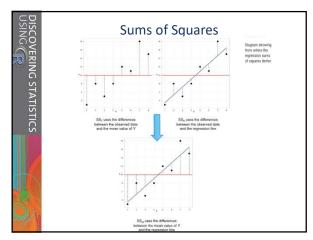




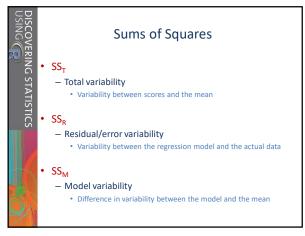
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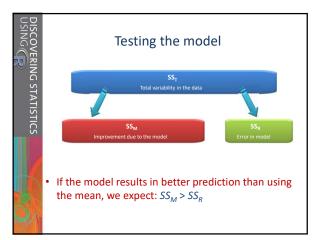
Testing the model

• How well does our model 'fit' the data?

-  $R^2$ • The proportion of the total variability in the data accounted for by the regression model

• Equivalent to the Pearson correlation coefficient squared  $R^2 = \frac{SS_M}{SS_T}$ 

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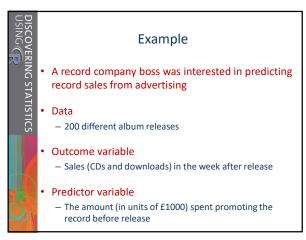


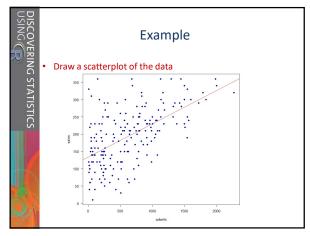
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# Testing the model • However, recall from Chapter 2 - Sums of squares are total values • Good measures of variability, but dependent on sample size - To overcome this, we can calculate Mean Squares (MS) • Compare to how we calculated variance!! • To assess whether our regression model fits the data better than the mean, we can thus perform an analysis of variance (ANOVA): F = MS\_M MS\_R

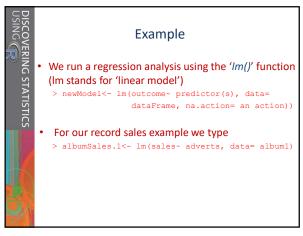
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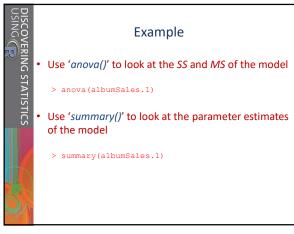
BIO 209: Discovering Statistics using R Erik Willems

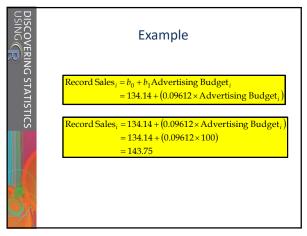




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# Aims and Objectives

- Understand the multiple regression equation
- Understand different methods of regression
  - Hierarchical
  - Forced entry
  - Stepwise
- Know how to perform a multiple regression and interpret the model
- Understand the assumptions of multiple regression and know how to test them

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## What is multiple regression?

- Simple regression is a model to predict the value of one variable from another
- Multiple regression is a natural extension of this:
  - Used to predict values of an outcome from several predictors
  - It is a hypothetical model of the relationship between several variables

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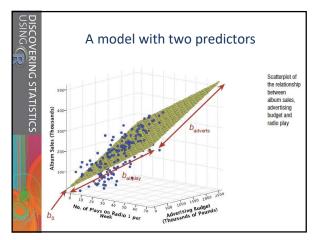


# Example

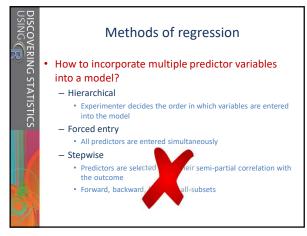
- A record company boss was interested in predicting album sales from advertising and air time
- Data
  - 200 different album releases
- Outcome variable
- $\boldsymbol{\mathsf{-}}$  Sales (CDs and downloads) in the week after release
- Predictor variables
  - The amount (in units of £1000) spent promoting the record
  - $\boldsymbol{-}$  Air time on the radio

Multiple regression as an equation  $y_i = b_0 + b_1 X_1 + b_2 X_2 + ... + b_n X_n + \varepsilon_i$ • In simple regression our model is described by a straight line
• In multiple regression we extend this equation  $-b_0$ • Intercept (value of Y when all Xs= 0)
• Point at which the regression line crosses the Y-axis (vertical)  $-b_{1-n}$ • Regression coefficients for predictors 1 to n
• Gradient (slope) of the regression line
• Direction/strength of relationship

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### Model fit

- Sums of Squares
  - Although the computation of these values is more complex than in simple regression, their use and interpretation is the same
  - **P**2
  - This is a multiple R<sup>2</sup>, i.e. the squared correlation coefficient between observed and fitted values based on all predictors
  - As in simple regression, it is a measure of the proportion of the total variance in the outcome variable accounted for by the model
- Akaike Information Criterion (AIC)
  - Assess model fit, while penalizing for model complexity
  - Only makes sense when comparing different models of the same data, in which case: lower values represent a better fit

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### Model diagnostics

- To assess a model's accuracy within the sample
  - Look for outliers
    - Calculate the standardized residual for each case
    - 95% of data should lie between -1.96 and +1.96
    - 99% of data should lie between -2.58 and + 2.58
    - An absolute value greater than 3 is indicative of an outlier
  - Look for influential cases
    - There are several residual statistics to assess the influence of a particular case on model parameter estimates (Cook's distance, DFBeta, DFFit, leverage and the covariance ratio)
    - "Quick and dirty" rule of thumb: any case with an absolute value of its Cook's distance > 1 may be cause for concern

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# Model generalization

!!Dummy coding!!

- To assess how well a model generalizes from the sample to the population, several assumptions must hold
  - Pretty straightforward ones:
    - Variable type
      - Outcome must be a continuous variable
    - Predictors can be continuous or dichotomous variables
    - Non-zero variance
      - Predictors must not have zero variance
    - Linearity
      - The relationship we model is, in reality, linear
    - Independence
      - All values of the outcome should come from a different entity

Model generalization

• To assess how well a model generalizes from the sample to the population, several assumptions must hold

- More tricky ones:

• No multicollinearity

- Predictors must not be highly correlated

• Homoscedasticity

- For each value of the predictors the variance of the error term should be constant

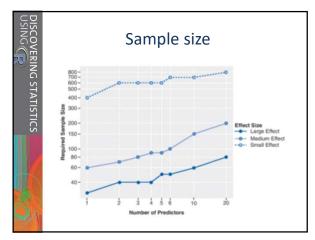
• Independent errors

- For any pair of observations, the error terms should be uncorrelated

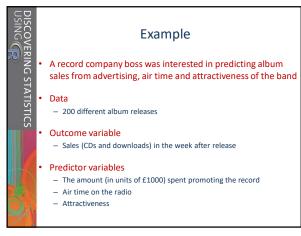
• Normally distributed errors

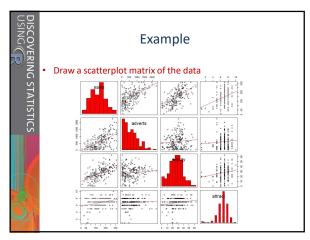
- Visual inspection of histograms and QQ plots

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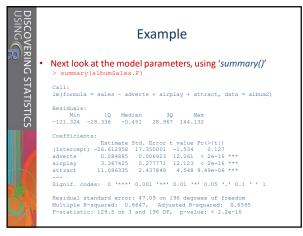
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DISCOVERI USING		Example  • Let's say we want to use the 'forced entry' method (unlike the example in the book)					
RING STATISTICS	•						
- AT	> albumSales.F<- lm(sales~ adverts + airplay +						+ airplay +
TSI.	attract, data= album2)						
SOL	•	Inspect SS	an	d <i>MS</i> , ι	using the	'anova	()' function
		> anova(albumSales.F)					
	Analysis of Variance Table Response: sales						
			Df	Sum Sq	Mean Sq	F value	Pr(>F)
		adverts	1	433688	433688	195.600	<2.2e-16***
A )		airplay	1	381836	381836	172.214	<2.2e-16***
		attract	1	45853	45853	20.681	9.49e-06***
		Residuals	196	434575	2217		

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# Example

- We can now write the equation of our regression model  $sales_i = b_0 + b_1 \ adverts + b_2 \ airplay + b_3 \ attract + \varepsilon_i$   $sales_i = -26.61 + 0.08*adverts + 3.37*airplay + 11.09*attract + \varepsilon_i$
- To be able to directly compare the influence of different predictors, we can calculate standardized parameters (β<sub>i</sub> instead of b<sub>i</sub>) using 'lm.beta()'
   lm.beta(albumSales.F)
  - adverts airplay attract 0.5108462 0.5119881 0.1916834

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DISCOVERING STATISTICS

### Outliers and influential cases

- Add residual statistics to the original dataframe, e.g.
  - > album2\$res<- resid(albumSales.F)
  - > album2\$stz.r<- rstandard(albumSales.F)
  - > album2\$stu.r<- rstudent(albumSales.F)
    > album2\$cd<- cooks.distance(albumSales.F)</pre>
  - > album2\$dfbeta<- dfbeta(albumSales.F)
  - > album2\$dffit<- dffits(albumSales.F)</pre>
  - > album2\$leverage<- hatvalues(albumSales.F)</pre>
  - > album2\$stz.cvr<- covratio(albumSales.F)</pre>
- Evaluate these against the criteria detailed in book

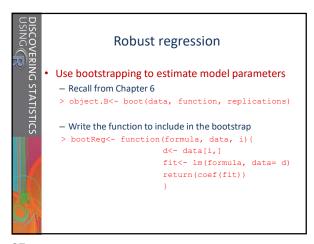
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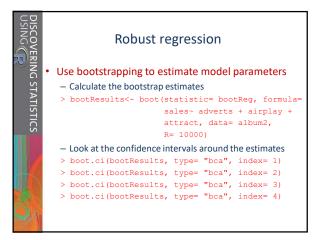


### Model assumptions

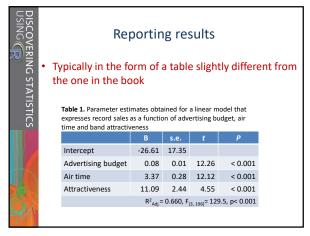
(the tricky ones)

- Independent errors
  - Durban-Watson test must not be significant
    - > dwt(albumSales.F)
- No multicollinearity
  - Value Inflation Factors (VIF)< 10</li>
  - Mean VIF not substantially greater than 1
    > vif(albumSales.F)
- Normally distributed errors and homoscedasticity
  - Inspect plots
    - > par(mfrow= c(2,2)); plot(albumSales.F)





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# Where do we go from here...

- Most of the remaining models we'll cover in this course are variations and generalizations of the regression (or linear model) framework
- We will explore ways in which to deal with:
  - categorical predictor variables that are not dichotomies (dummy coding -this chapter- and Chapters 10-12)
  - dependencies in the data, e.g. PGLS-models, repeated measures designs (Karin, Chapter 13)
- Biological data typically suffer from both non-normality and dependencies, and at the end of the course you should be sufficiently equipped to deal with these complications

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DISCOVERING STATISTICS

### Rest of today...

- Practical Chapter 7
  - Read § 7.1, 7.2, "Cramming Sam's Tips" and "What Have I Discovered about Statistics?"
  - Skip sections on R commander: §7.4.1, § 7.8.2.1, §7.8.4.1, §7.9.1
  - Work through self-tests as you see fit (but skip self-tests in §7.12)
  - Solve Smart Alex's Tasks 1-2

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### Errata

• p. 299

As we did for correlations, we need to write a function (R's Souls' Tip 6.2) we want to bootstrap. We'll write one called bootReg() – this function is a little more complex than the function we wrote for the correlation, because we are interested in more than one statistic (we have an intercept and three slope parameters to bootstrap). The function we need to execute is:

bootReg <- function (formula, data, indices) { d <- data [i,]