# Fitting Linear Models

Requires assumptions about  $\epsilon_i$ s. Usual assumptions:

- 1.  $\epsilon_1, \ldots, \epsilon_n$  are independent. (Sometimes we assume only that  $Cov(\epsilon_i, \epsilon_j) = 0$  for  $i \neq j$ ; that is we assume the errors are **uncorrelated**.)
- 2. Homoscedastic errors; all variances are equal:

$$\operatorname{Var}(\epsilon_1) = \operatorname{Var}(\epsilon_2) = \cdots = \sigma^2$$

3. Normal errors:  $\epsilon_i \sim N(0, \sigma^2)$ .

Remember: we already have assumed  $E(\epsilon_i) = 0$ .



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## Notes

- Assumptions 1, 2 and 3 permit Maximum Likelihood
   Estimation.
- Assumptions 1 and 2 justify least squares.
- Assumption 3 can be replaced by other error distributions, but not in this course.
- With normal errors maximum likelihood estimates are the same as least squares estimates.
- Assumption 2 Homoscedastic errors can be relaxed; see STAT 402 "Generalized Linear Models" or "weighted least square".
- Assumption 1 can be relaxed; see STAT 804 for Time Series models.



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# Motivation for Least Squares

Choose  $\hat{\beta}$  to make **fitted values**  $\hat{\mu} = X\hat{\beta}$  as close to Ys as possible.

There are many possible choices for "close":

Mean Absolute Deviation: minimize

$$\frac{1}{n}\sum|Y_i-\hat{\mu}_i|$$

Least Median of Squares: minimize

$$\mathrm{median}\{|Y_i - \hat{\mu}_i|^2\}$$

Least squares: minimize

$$\sum (Y_i - \hat{\mu}_i)^2$$

WE DO LS = least squares.



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To minimize  $\sum (Y_i - \hat{\mu}_i)^2$  take derivatives with respect to each  $\hat{\beta}_j$  and set them equal to 0:

$$\frac{\partial}{\partial \hat{\beta}_j} \sum_{i=1}^n (Y_i - \hat{\mu}_i)^2 = \sum_{i=1}^n \frac{\partial}{\partial \hat{\beta}_j} (Y_i - \hat{\mu}_i)^2$$
$$= \sum_{i=1}^n \left[ \frac{\partial}{\partial \hat{\mu}_i} (Y_i - \hat{\mu}_i)^2 \right] \frac{\partial \hat{\mu}_i}{\partial \hat{\beta}_j}$$

But

$$\frac{\partial}{\partial \hat{\mu}_i} (Y_i - \hat{\mu}_i)^2 = -2(Y_i - \hat{\mu}_i)$$

and

$$\hat{\mu}_i = \sum_{j=1}^p x_{ij}\hat{\beta}_j$$

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#### Thus





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### Normal Equations

Set this equal to 0; Get so-called **normal equations**:

$$\sum_{i=1}^{n} Y_{i} x_{ij} = \sum_{i=1}^{n} \hat{\mu}_{i} x_{ij} \qquad j = 1, \dots, p$$

Finally remember that  $\hat{\mu}_i = \sum_{k=1}^p x_{ik} \hat{\beta}_k$  to get

$$\sum Y_i x_{ij} = \sum_{i=1}^n \sum_{k=1}^p x_{ij} x_{ik} \hat{\beta}_k \tag{1}$$



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### Formula looks dreadful

- but it's just a bunch of matrix-vector multiplications written out in summation notation.
- Note that it is a set of *p* linear equations in *p* unknowns  $\hat{\beta}_1, \ldots, \hat{\beta}_p$ .



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Normal equations in vector matrix form

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is the dot product between the *j*th and *k*th columns of X. Another way to view this is as the *jk*th entry in the matrix  $X^T X$ :

$$X^{T}X = \begin{bmatrix} x_{11} & x_{21} & \cdots & x_{n1} \\ x_{12} & x_{22} & \cdots & x_{n2} \\ \vdots & \vdots & \cdots & \vdots \\ x_{1p} & x_{2p} & \cdots & x_{np} \end{bmatrix} \begin{bmatrix} x_{11} & \cdots & x_{1p} \\ \vdots & \cdots & \vdots \\ x_{n1} & \cdots & x_{np} \end{bmatrix}$$



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The *jk*th entry in this matrix product is clearly

$$x_{1j}x_{1k} + x_{2j}x_{2k} + \cdots + x_{nj}x_{nk}$$

so that the right hand side of (1) is

$$\sum_{k=1}^{p} (X^{\mathsf{T}} X)_{jk} \hat{\beta}_{k}$$

which is just the matrix product

 $((X^{\mathsf{T}}X)\hat{\beta})_j$ 



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Now look at the left hand side of (1), namely,  $\sum_{i=1}^{n} Y_i x_{ij}$  which is just the dot product of Y and the *j*th column of X or the *j*th entry of  $X^T Y$ :

$$\begin{bmatrix} x_{11} & x_{21} & \cdots & x_{n1} \\ x_{12} & x_{22} & \cdots & x_{n2} \\ \vdots & \vdots & \cdots & \vdots \\ x_{1p} & x_{2p} & \cdots & x_{np} \end{bmatrix} \begin{bmatrix} Y_1 \\ Y_2 \\ \vdots \\ Y_n \end{bmatrix} = \begin{bmatrix} x_{11}Y_1 + x_{21}Y_2 + \cdots + x_{n1}Y_n \\ \vdots \\ x_{1p}Y_1 + x_{2p}Y_2 + \cdots + x_{np}Y_n \end{bmatrix}$$

So the normal equations are

$$(X^{\mathsf{T}}Y)_j = (X^{\mathsf{T}}X\hat{\beta})_j$$

or just

$$X^T Y = X^T X \hat{\beta}$$

Last formula is usual way to write the normal equations.



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# Solving Normal Equations for $\hat{\beta}$

Let's look at the dimensions of the matrices first.

- $X^T$  is  $p \times n$ ,
- ➤ Y is n × 1,
- X<sup>T</sup>X is a p × n matrix multiplied by a n × p matrix which just produces a p × p matrix.
- If the matrix X<sup>T</sup>X has rank p then X<sup>T</sup>X is not singular and its inverse (X<sup>T</sup>X)<sup>-1</sup> exists. So solve

$$X^T Y = X^T X \hat{\beta}$$

for  $\hat{\beta}$  by multiplying both sides by  $(X^T X)^{-1}$  to get

$$\hat{\beta} = (X^T X)^{-1} X^T Y$$

This is the ordinary least squares estimator. See assignment 1 for an example with rank(X) < p. See chapter 5 in the text for a review of matrices and vectors.



Normal Equations for Simple Linear Regression

Thermoluminescence Example

See Introduction for the framework. Here I consider two models:

a straight-line model,

$$Y_i = \beta_1 + \beta_2 D_i + \epsilon_i$$

► a quadratic model,

$$Y_i = \beta_1 + \beta_2 D_i + \beta_3 D_i^2 + \epsilon_i \,.$$



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First, general theoretical formulas, then numbers and arithmetic:

$$X = \begin{bmatrix} 1 & D_1 \\ \vdots & \vdots \\ 1 & D_n \end{bmatrix}$$
$$X^{\mathsf{T}}X = \begin{bmatrix} 1 & \cdots & 1 \\ D_1 & \cdots & D_n \end{bmatrix} \begin{bmatrix} 1 & D_1 \\ \vdots & \vdots \\ 1 & D_n \end{bmatrix} = \begin{bmatrix} n & \sum_{i=1}^n D_i \\ \sum_{i=1}^n D_i & \sum_{i=1}^n D_i^2 \end{bmatrix}$$
$$(X^{\mathsf{T}}X)^{-1} = \frac{1}{n \sum D_i^2 - (\sum D_i)^2} \begin{bmatrix} \sum_{i=1}^n D_i^2 & -\sum_{i=1}^n D_i \\ -\sum_{i=1}^n D_i & n \end{bmatrix}$$



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$$X^{T}Y = \begin{bmatrix} 1 & \cdots & 1 \\ D_{1} & \cdots & D_{n} \end{bmatrix} \begin{bmatrix} Y_{1} \\ \vdots \\ Y_{n} \end{bmatrix} = \begin{bmatrix} \sum_{i=1}^{n} Y_{i} \\ \sum_{i=1}^{n} D_{i}Y_{i} \end{bmatrix}$$

$$(X^T X)^{-1} X^T Y = \begin{bmatrix} \frac{\sum Y_i \sum D_i^2 - \sum D_i \sum D_i Y_i}{n \sum D_i^2 - (\sum D_i)^2} \\ \frac{n \sum D_i Y_i - (\sum D_i)(\sum Y_i)}{n \sum D_i^2 - (\sum D_i)^2} \end{bmatrix}$$



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$$(X^T X)^{-1} X^T Y = \begin{bmatrix} \frac{\bar{Y} \sum D_i^2 - \bar{D} \sum D_i Y_i}{\sum (D_i - \bar{D})^2} \\ \frac{\sum (D_i - \bar{D})(Y_i - \bar{Y})}{\sum (D_i - \bar{D})^2} \end{bmatrix}$$





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### The data are

Dose	Count
0	27043
0	26902
0	25959
150	27700
150	27530
150	27460
420	30650
420	30150
420	29480
900	34790
900	32020
1800	42280
1800	39370
1800	36200
3600	53230
3600	49260
3600	53030



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The design matrix for the linear model is

X

[ 1	27043
1	26902
1	25959
1	27700
1	27530
1	27460
1	30650
1	30150
1	29480
1	34790
1	32020
1	42280
1	39370
1	36200
1	53230
1	49260
$\lfloor 1$	53030
	$   \begin{bmatrix}     1 \\$



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- Compute  $X^T X$  in Minitab or Splus or R.
- That matrix has 4 numbers each of which is computed as the dot product of 2 columns of X.
- For instance the first column dotted with itself produces  $1^2 + \cdots + 1^2 = 17$ .
- Here is an example S session which reads in the data, produces the design matrices for the two models and computes X<sup>T</sup>X.



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[36] skekoowahts% S # The data are in a file called linear. The ! # tells S that what follows is not an S command but a standard # UNIX (or DOS) command # > !more linear Dose Count 0 27043 0 26902 0 25959 150 27700 150 27530 150 27460 420 30650 420 30150 420 29480 900 34790 900 32020 1800 42280 1800 39370 1800 36200 3600 53230 3600 49260 3600 53030 #



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```
# The function help(function) produces help for
# a function such as
# > help(read.table)
#
# Read in the data from a file. The file has 18 lines:
# 17 lines of data and a first line which has the names
# of the variables. The function read.table reads such
# data and header=T warns S that the first line is
# variable names. The first argument of read.table is
# a character string containing the name of the file
# to read from.
#
```

> dat <- read.table("linear",header=T)</pre>



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> dat

Dose Count

- 1 0 27043
- 2 0 26902
- 3 0 25959
- 4 150 27700
- 5 150 27530
- 6 150 27460
- 7 420 30650
- 8 420 30150
- 9 420 29480
- 10 900 34790
- 11 900 32020
- 12 1800 42280
- 13 1800 39370
- 14 1800 36200
- 15 3600 53230
- 16 3600 49260
- 17 3600 53030
- #

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```
# the design matrix has a column of 1s and also
# a column consisting of the first column of dat
# which is just the list of covariate values
# The notation dat[,1] picks out the first column of dat
#
> design.mat <- cbind(rep(1,17),dat[,1])
#
# To print out an object you type its name!
#</pre>
```



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>	design.mat
---	------------

	[,1]	[,2]
[1,]	1	0
[2,]	1	0
[3,]	1	0
[4,]	1	150
[5,]	1	150
[6,]	1	150
[7,]	1	420
[8,]	1	420
[9,]	1	420
[10,]	1	900
[11,]	1	900
[12,]	1	1800
[13,]	1	1800
[14,]	1	1800
[15,]	1	3600
[16,]	1	3600
[17,]	1	3600
#		



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```
# Compute X^T X -- uses %*% to multiply matrices
# and t(x) to compute the transpose of a matrix x.
#
> xprimex <- t(design.mat)%*% design.mat</pre>
> xprimex
     [,1] [,2]
[1,] 17 19710
[2,] 19710 50816700
#
# Compute X^T Y
#
> xprimey <- t(design.mat)%*% dat[,2]</pre>
> xprimey
          [,1]
[1,]
        593054
[2,] 882452100
```



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#
# Next compute least squares estimates by solving
# normal equations
#
> solve(xprimex,xprimey)
       [,1]
[1,] 26806.734691
[2,] 6.968012
#
# solve(A,b) computes solution of Ax=b for A a
# square matrix and b a vector. Note x=A^{-1}b.
#
```



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#
# The next piece of code regresses the variable
# Count on Dose taking the data from dat.
#
> lm( Count~Dose,data=dat)
Call:
lm(formula = Count ~ Dose, data = dat)
Coefficients:
 (Intercept) Dose
    26806.73 6.968012
Degrees of freedom: 17 total; 15 residual
Residual standard error: 1521.238
#
# Notice the estimates agree with our calculations
# Residual standard error is usual estimate of sigma
# namely the square root of the Mean Square for Error.
#
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#
 Now add a third column to fit the quadratic model
#
#
 design.mat2_cbind(design.mat,design.mat[,2]^2)
>
#
# Here is X^T X
#
> t(design.mat2)%*% design.mat2
         [,1] [,2]
                                  [,3]
[1,]
          17
                   19710 5.081670e+07
[2,]
                 50816700 1.591544e+11
       19710
[3,] 50816700 159154389000 5.367847e+14
```



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#
# Here is X^T Y
#
> t(design.mat2)%*% dat[,2]
             [.1]
[1,] 5.930540e+05
[2,] 8.824521e+08
[3,] 2.469275e+12
#
# But the following illustrates the dangers
# of doing computations blindly on the computer.
# The trouble is that the design matrix has a
# third column which is so much larger that
# the first two.
#
> solve(t(design.mat2)%*% design.mat2,
        t(design.mat2)%*% dat[,2])
Error in solve.qr(a, b): apparently singular matrix
Dumped
```



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#
# However, good packages know numerical techniques
# which avoid the danger.
#
> lm(Count ~ Dose+Dose^2,data=dat)
Call:
lm(formula = Count ~ Dose + Dose^2, data = dat)
```

```
Coefficients:
(Intercept) Dose I(Dose<sup>2</sup>)
26718.11 7.240314 -7.596867e-05
```

```
Degrees of freedom: 17 total; 14 residual
Residual standard error: 1571.277
```



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```
#
# WARNING: you can't tell from the size of the
# estimate of an estimate such as that of beta_3
# whether or not it is important -- you have to
# compare it to values of the corresponding
# covariate and to its standard error
#
> q()
# Used to quit S: pay attention to () --
# that part is essential!
```



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