Introduction to statistics: Matrix formulatuon of the linear model

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Matrix form of the linear model

- ▶ This lecture is not crucial for following this course.
- ▶ It is however useful to have seen the matrix form of the linear model once.
- Having see this formulation once will help in understanding contrast coding.

Consider a deterministic function $\phi(\mathbf{x}, \beta)$ which takes as input some variable values x and some fixed values β . A simple (if abstract) example would be

$$y = \beta x \tag{1}$$

Another example with two fixed values β_0 and β_1 is:

$$y = \beta_0 + \beta_1 x \tag{2}$$

We can rewrite the above equation as follows.

$$y = \beta_0 + \beta_1 x$$

$$= \beta_0 \times 1 + \beta_1 x$$

$$= (1 \quad x) \begin{pmatrix} \beta_0 \\ \beta_1 \end{pmatrix}$$

$$y = \phi(x, \beta)$$
(3)

In a statistical model, we don't expect an equation like $y=\phi(x,\beta)$ to fit all the points exactly.

For example, we could come up with an equation that, given a person's weight, gives a prediction of their height:

predicted height =
$$\beta_0 + \beta_1$$
 weight (4)

Given any single value of the weight of a person, we will probably not get a perfectly correct prediction of the height of that person. This leads us to a non-deterministic version of the above function:

$$y = \phi(x, \beta, \varepsilon) = \beta_0 + \beta_1 x + \varepsilon \tag{5}$$

Here, ε is an error random variable which is assumed to have some PDF (the normal distribution) associated with it. It is assumed to have expectation (mean) 0, and some standard deviation (to be estimated from the data) σ .

We can write this statement in compact form as $\varepsilon \sim N(0,\sigma)$.

The **general linear model** is a non-deterministic function like the one above:

$$Y = f(x)^T \beta + \varepsilon \tag{6}$$

The matrix formulation will be written as below. n refers to the number of data points (that is, Y_1, \ldots, Y_n), and the index j ranges from 1 to n.

$$Y = X\beta + \varepsilon \Leftrightarrow y_j = f(x_j)^T \beta + \varepsilon_j, j = 1, \dots, n$$
 (7)

To make this concrete, suppose we have three data points, i.e., n=3. Then, the matrix formulation is

$$\begin{pmatrix} y_1 \\ y_2 \\ y_3 \end{pmatrix} = \begin{pmatrix} 1 & x_1 \\ 1 & x_2 \\ 1 & x_3 \end{pmatrix} \begin{pmatrix} \beta_0 \\ \beta_1 \end{pmatrix} + \varepsilon \tag{8}$$

$$Y = X\beta + \varepsilon \tag{9}$$

Here, $f(x_1)^T = (1 \ x_1)$, and is the first row of the matrix X, $f(x_2)^T = (1 \ x_2)$ is the second row, and $f(x_3)^T = (1 \ x_3)$ is the third row.

Note that $E[Y]=X\beta.$ β is a $p\times 1$ matrix, and X, the **design matrix**, is $n\times p.$

- When we have a deterministic model $y = \phi(f(x)^T, \beta) = \beta_0 + \beta_1 x$, this implies a perfect fit to all data points.
- ▶ This is like solving the equation Ax = b in linear algebra: we solve for β in $X\beta = y$ using, e.g., Gaussian elimination.
- When we have a non-deterministic model $y=\phi(f(x)^T,\beta,\varepsilon)$, there is no solution. Now, the best we can do is to get Ax to be as close an approximation as possible to b in Ax=b.
- ▶ In other words, we try to minimize |b Ax|.

- ▶ The goal is to estimate β ; we want to find a value of β such that the observed Y is as close to its expected value $X\beta$.
- ▶ In order to be able to identify β from $X\beta$, the linear transformation $\beta \to X\beta$ should be one-to-one, so that every possible value of β gives a different $X\beta$.

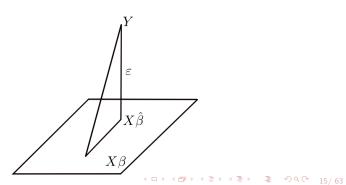
This in turn requires that X be of full rank p.

So, if X is an $n \times p$ matrix, then it is necessary that $n \ge p$. There must be at least as many observations as parameters. If this is not true, then the model is said to be **over-parameterized**.

- Assuming that X is of full rank, and that n > p, Y can be considered a point in n-dimensional space and the set of candidate $X\beta$ is a p-dimensional subspace of this space; see the figure on the next slide.
- There will be one point in this subspace which is closest to Y in terms of Euclidean distance. The unique β that corresponds to this point is the **least squares estimator** of β ; we will call this estimator $\hat{\beta}$.

Lecture 5 Linear modeling theory: Matrix formulation Least squares estimation: Geometric argument

Geometric argument



Notice that $\varepsilon=(Y-X\hat{\beta})$ and $X\beta$ are perpendicular to each other. Because the dot product of two perpendicular (orthogonal) vectors is 0, we get the result:

$$(Y - X\hat{\beta})^T X \beta = 0 \Leftrightarrow (Y - X\hat{\beta})^T X = 0$$
 (10)

Example:

```
x1<-c(1,0)
x2<-c(0,1)
sum(x1*x2)
## [1] 0
```

Multiplying out the terms, we proceed as follows. One result that we use here is that $(AB)^T=B^TA^T.$

$$(Y - X\hat{\beta})^T X = 0$$

$$(Y^T - \hat{\beta}^T X^T) X = 0$$

$$\Leftrightarrow Y^T X - \hat{\beta}^T X^T X = 0$$

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

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

$$\Leftrightarrow X^T Y = X^T X \hat{\beta}$$
(11)

This gives us the important result:

$$\hat{\beta} = (X^T X)^{-1} X^T Y \tag{12}$$

X is of full rank, therefore X^TX is invertible.

- ▶ $X^+ = (X^T X)^{-1} X^T$ is the **generalized matrix inverse** of the design matrix X, and will become very important for us in the **contrast coding** lecture.
- We will see in the contrast coding lecture how X^+ defines hypothesis tests in the linear (mixed) model.

Example:

```
(X < -matrix(c(rep(1,8), rep(c(-1,1), each=4),
         rep(c(-1,1), each=2,2)), ncol=3))
      [,1] [,2] [,3]
##
## [1,] 1 -1 -1
## [2,] 1 -1 -1
## [3,] 1 -1 1
## [4,] 1 -1 1
## [5.]
## [6,]
## [7,]
## [8,]
```

```
library(Matrix)
## full rank:
rankMatrix(X)
## [1] 3
## attr(,"method")
## [1] "tolNorm2"
## attr(,"useGrad")
## [1] FALSE
## attr(,"tol")
## [1] 1.7764e-15
```

Our model is:

$$Y = X\beta + \varepsilon \tag{13}$$

The mean and variance of the parameters beta

Let $\varepsilon \sim N(0, \sigma^2)$.

In other words, we are assuming that each value generated by the random variable ε is independent and it has the same distribution, i.e., it is identically distributed. This is sometimes shortened to the iid assumption.

So we should technically be writing:

$$Y = X\beta + \varepsilon \quad \varepsilon \sim N(0, \sigma) \tag{14}$$

and add that Y are independent and identically distributed. We could also have written:

$$Y \sim Normal(X\beta, \sigma)$$
 (15)

Linear modeling theory: Matrix formulation

The expectation and variance of the parameters beta

The mean and variance of the parameters beta

Some consequences of the above statements:

- 1. $E(\varepsilon) = 0$
- 2. $Var(\varepsilon) = \sigma^2 I_n$ 3. $E[Y] = X\beta = \mu$
- 4. $Var(Y) = \sigma^2 I_n$

 I_n here is the identity matrix, e.g.:

```
I<-matrix(c(1,0,0,1),ncol = 2)
sigma<-10
sigma*I

## [,1] [,2]
## [1,] 10 0
## [2,] 0 10</pre>
```

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The mean and variance of the parameters beta

We can now derive the expectation and variance of the estimators $\hat{\beta}$.

We need a fact about variances: Var(aB), where a is a constant, is $a^2Var(B)$. In the matrix setting, Var(AB), where A is a constant, is $AVar(B)A^T$.

$$E[\hat{\beta}] = E[(X^T X)^{-1} X^T Y] = (X^T X)^{-1} X^T X \beta = \beta$$
 (16)

Notice that the above shows that $\hat{\beta}$ is an unbiased estimator of β .

Next, we compute the variance:

$$Var(\hat{\beta}) = Var([(X^T X)^{-1} X^T]Y)$$
(17)

Expanding the right hand side out:

$$Var([(X^TX)^{-1}X^T]Y) = [(X^TX)^{-1}X^T]Var(Y)[(X^TX)^{-1}X^T]^T$$
(18)

The mean and variance of the parameters beta

Replacing Var(Y) with its variance $\sigma^2 I$, and unpacking the transpose on the right-most expression $[(X^TX)^{-1}X^T]^T$:

$$Var(\hat{\beta}) = [(X^T X)^{-1} X^T] \sigma^2 IX [(X^T X)^{-1}]^T$$
 (19)

Since σ^2 is a scalar we can move it to the left, and any matrix multiplied by I is the matrix itself, so we ignore I, getting:

$$Var(\hat{\beta}) = \sigma^2[(X^T X)^{-1} X^T X [(X^T X)^{-1}]^T$$
 (20)

The mean and variance of the parameters beta

Since $(X^TX)^{-1}X^TX = I$, we can simplify to

$$Var(\hat{\beta}) = \sigma^2 [(X^T X)^{-1}]^T$$
 (21)

Now, $(X^TX)^{-1}$ is symmetric, so $[(X^TX)^{-1}]^T=(X^TX)^{-1}.$ This gives us:

$$Var(\hat{\beta}) = \sigma^2 (X^T X)^{-1} \tag{22}$$

The mean and variance of the parameters beta

An example:

```
hindi<-read.table("data/hindiJEMR.txt",header=TRUE)
y<-as.matrix(hindi$TFT)
x<-log(hindi$word_len)
m0<-lm(y~x)</pre>
```

The mean and variance of the parameters beta

```
## (X^TX)^{-1}
invXTX<-solve(t(X)%*%X)
## estimate of beta:
(beta<-invXTX%*%t(X)%*%y)

## [,1]
## (Intercept) 30.872
## x 189.429</pre>
```

```
## estimated SD (se) of the estimate of beta:
(hat_sigma<-summary(m0)$sigma)

## [1] 239.88

(hat_var<-hat_sigma^2*invXTX)

## (Intercept) x

## (Intercept) 4.0592 -2.9284

## x -2.9284 2.5679</pre>
```

Linear modeling theory: Matrix formulation

The expectation and variance of the parameters beta

The mean and variance of the parameters beta

What we have here is a bivariate normal distribution as an estimate of the β parameters:

$$\begin{pmatrix} \hat{\beta}_0 \\ \hat{\beta}_1 \end{pmatrix} \sim N(\begin{pmatrix} 210.7777 \\ 129.4064 \end{pmatrix}, \begin{pmatrix} 31.35647 & -21.61119 \\ -21.61119 & 16.88379 \end{pmatrix})$$
 (23)

The mean and variance of the parameters beta

The variance of a bivariate distribution has the variances along the diagonal, and the covariance between β_0 and β_1 on the off-diagonals. Covariance is defined as:

$$Cov(\hat{\beta}_0, \hat{\beta}_1) = \rho \sigma_{\hat{\beta}_0} \sigma_{\hat{\beta}_1}$$
 (24)

where ρ is the correlation between β_0 and β_1 .

So $\hat{\beta}_0 \sim N(210.78, 31.36)$ and $\hat{\beta}_1 \sim N(129.41, 16.88)$, and $Cov(\beta_0, \beta_1) = -21.61$. So the correlation between the $\hat{\beta}$ is

```
## hat rho:
-21.61/(sqrt(31.36)*sqrt(16.88))
## [1] -0.93925
```

The likelihood ratio test

Define the likelihood ratio statistic as:

$$\Lambda = \frac{\max_{\theta \in \omega_0}(lik(\theta))}{\max_{\theta \in \omega_1}(lik(\theta))}$$
 (25)

where, $\omega_0 = \{\mu_0\}$ and $\omega_1 = \{\forall \mu \mid \mu \neq \mu_0\}$.

Suppose that X_1, \ldots, X_n are iid and normally distributed with mean μ and standard deviation σ (assume for simplicity that σ is known).

The likelihood ratio test

Let the null hypothesis be $H_0: \mu=\mu_0$ and the alternative be $H_1: \mu \neq \mu_0$. Here, μ_0 is a number, such as 0. Now, the numerator of the likelihood ratio statistic is:

$$\frac{1}{(\sigma\sqrt{2\pi})^n}exp\left(-\frac{1}{2\sigma^2}\sum_{i=1}^n(X_i-\mu_0)^2\right)$$
 (26)

For the denominator, the maximum likelihood can be achieved by specifying the MLE \bar{X} as μ :

$$\frac{1}{(\sigma\sqrt{2\pi})^n}exp\left(-\frac{1}{2\sigma^2}\sum_{i=1}^n(X_i-\bar{X})^2\right)$$
 (27)

The likelihood ratio statistic is then:

$$\Lambda = \frac{\frac{1}{(\sigma\sqrt{2\pi})^n} exp\left(-\frac{1}{2\sigma^2} \sum_{i=1}^n (X_i - \mu_0)^2\right)}{\frac{1}{(\sigma\sqrt{2\pi})^n} exp\left(-\frac{1}{2\sigma^2} \sum_{i=1}^n (X_i - \bar{X})^2\right)}$$
(28)

Canceling out common terms:

$$\Lambda = \frac{exp\left(-\frac{1}{2\sigma^2}\sum_{i=1}^n(X_i - \mu_0)^2\right)}{exp\left(-\frac{1}{2\sigma^2}\sum_{i=1}^n(X_i - \bar{X})^2\right)}$$
(29)

Taking logs:

$$\log \Lambda = \left(-\frac{1}{2\sigma^2} \sum_{i=1}^n (X_i - \mu_0)^2 \right) - \left(-\frac{1}{2\sigma^2} \sum_{i=1}^n (X_i - \bar{X})^2 \right)$$

$$= -\frac{1}{2\sigma^2} \left(\sum_{i=1}^n (X_i - \mu_0)^2 - \sum_{i=1}^n (X_i - \bar{X})^2 \right)$$
(30)

Now, note that

$$\sum_{i=1}^{n} (X_i - \mu_0)^2 = \sum_{i=1}^{n} (X_i - \bar{X})^2 + n(\bar{X} - \mu_0)^2$$
 (31)

This means that

$$\sum_{i=1}^{n} (X_i - \mu_0)^2 - \sum_{i=1}^{n} (X_i - \bar{X})^2 = n(\bar{X} - \mu_0)^2$$
 (32)

So, we can write Λ as:

$$\Lambda = -\frac{1}{2\sigma^2}n(\bar{X} - \mu_0)^2 \tag{33}$$

Rearranging terms:

$$-2\Lambda = \frac{n(\bar{X} - \mu_0)^2}{\sigma^2} \tag{34}$$

Or maybe even more transparently:

$$-2\Lambda = \frac{(\bar{X} - \mu_0)^2}{\frac{\sigma^2}{n}} \tag{35}$$

This should remind you of the Wald statistic we saw earlier—the t-test! Basically, all this is saying is that we reject the null when $\mid \bar{X} - \mu_0 \mid$ is large.

More generally, we will define the **likelihood ratio test statistic** as follows. Here, $L(\theta)$ refers to the likelihood given some value of θ , and $\ell(\theta)$ refers to the log likelihood.

$$L = -2 \times \log(L(\theta_0)/L(\theta_1))$$

$$\log L = -2 \times \{\ell(\theta_0) - \ell(\theta_1)\}$$

$$\log L = -2 \times \{\ell(\theta_0) - \ell(\theta_1)\}$$
(36)

where θ_1 and θ_0 are the estimates of θ under the alternative and null hypotheses, respectively. The likelihood ratio test rejects H_0 if $\log L$ is sufficiently large. As the sample size approaches infinity:

$$\log L \to \chi_r^2 \text{ as } n \to \infty \tag{37}$$

where r is called the degrees of freedom and is the difference in the number of parameters under H_1 and H_0 . 44 / 63

This is called Wilks' theorem. The proof of Wilks' theorem is fairly involved but you can find it on the internet if you are interested, or in Lehmann's *Testing Statistical Hypotheses*.

Note that sometimes you will see the form:

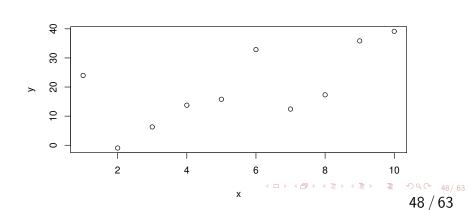
$$\log L = 2\{\ell(\theta_1) - \ell(\theta_0)\} \tag{38}$$

It should be clear that both statements are saying the same thing; in the second case, we are just subtracting the null hypothesis log likelihood from the alternative hypothesis log likelihood.

A practical example will make the usage of this test clear. Let's just simulate a linear model:

```
x<-1:10
y<- 10 + 2*x+rnorm(10,sd=10)
```

plot(x,y)



```
## null hypothesis model:
m0<-lm(y~1)
## alternative hypothesis model:
m1<-lm(y~x)</pre>
```

```
lambda<- -2*(logLik(m0)-logLik(m1))
## observed value:
lambda[1]
## [1] 5.4378</pre>
```

critical value:
qchisq(0.95,df=1)

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Here, we fit the null hypothesis model which only has an intercept term β_0 , and the alternative model that has β_1 as well.

Finally, we compare the λ with the critical chi-squared value for degrees of freedom 1.

We also computed the probability of getting a λ as extreme as we got assuming that the null is true:

Note that in the likelihood test above, we are comparing one nested model against another: the null hypothesis model is nested inside the alternative hypothesis model.

What this means is that the alternative hypothesis model contains all the parameters in the null hypothesis model plus some others.

We can compare two models, one nested inside another, as follows:

```
anova(m0,m1)
## Analysis of Variance Table
##
## Model 1: y ~ 1
## Model 2: y ~ x
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 9 1545
## 2 8 897 1 648 5.78 0.043
```

The F-score you get here is actually the square of the t-value you get in the linear model summary:

```
sqrt(anova(m0,m1)$F[2])
## [1] 2.4042
summary(m1)$coefficients[2,3]
## [1] 2.4042
```

This is because $t^2=F.$ The proof is discussed on page 9 of the Dobson and Barnett book.

The ANOVA works as follows. First define the residual as:1

$$e = Y - X\hat{\beta}$$

(39)

The square of this is:

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

(40)

Note that I do not use ε here, but e, to refer to the residual. This is because these are the estimates of ε derived from the estimates $\hat{\beta}$.

Define the **deviance** as:

$$D = \frac{1}{\sigma^2} (Y - X\hat{\beta})^T (Y - X\hat{\beta})$$

$$= \frac{1}{\sigma^2} (Y^T - \hat{\beta}^T X^T) (Y - X\hat{\beta})$$

$$= \frac{1}{\sigma^2} (Y^T Y - Y^T X\hat{\beta} - \hat{\beta}^T X^T Y + \hat{\beta}^T X^T X\hat{\beta})$$

$$(41)$$

Linear modeling theory tells us $\hat{\beta}=(X^TX)^{-1}X^TY$. Premultiplying both sides with (X^TX) , we get

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

It follows that we can rewrite the last line in equation 41 as follows: We can replace $(X^TX)\hat{\beta}$ with X^TY .

$$D = \frac{1}{\sigma^2} (Y^T Y - Y^T X \hat{\beta} - \hat{\beta}^T X^T Y + \hat{\beta}^T \underline{X^T X} \hat{\beta})$$

$$= \frac{1}{\sigma^2} (Y^T Y - Y^T X \hat{\beta} - \hat{\beta}^T X^T Y + \hat{\beta}^T \underline{X^T Y})$$

$$= \frac{1}{\sigma^2} (Y^T Y - Y^T X \hat{\beta})$$

$$(42)$$

Notice that $Y^TX\hat{\beta}$ is a scalar (1 \times 1) and is identical to β^TX^TY (check this), so we could write:

$$D = \frac{1}{\sigma^2} (Y^T Y - \hat{\beta}^T X^T Y)$$

Assume now that we have data of size n.

Suppose we have a null hypothesis $H_0: \beta = \beta_0$ and an alternative hypothesis $H_1: \beta = \beta_1$.

Let the null hypothesis have q parameters, and the alternative p, where q .

Let X_0 be the design matrix for H_0 , and X_1 the design matrix for H_1 .

Compute the deviances D_0 and D_1 for each hypothesis, and compute ΔD :

$$\Delta D = D_0 - D_1 = \frac{1}{\sigma^2} [(Y^T Y - \hat{\beta}_0 X_0^T Y) - (Y^T Y - \hat{\beta}_1 X_1^T Y)]$$

$$= \frac{1}{\sigma^2} [\hat{\beta}_1 X_1^T Y - \hat{\beta}_0 X_0^T Y]$$
(43)

It turns out that the F-statistic has the following distribution if the null hypothesis is true:

$$F = \frac{\Delta D/(p-q)}{D_1/(n-p)} \sim F(p-q, n-p)$$
(44)

So, an extreme value of F is inconsistent with the null and we reject it.

The F-statistic is:

$$F = \frac{\Delta D/(p-q)}{D_1/(n-p)}$$

$$= \frac{\hat{\beta}_1 X_1^T Y - \hat{\beta}_0^T X_0^T Y}{p-q} / \frac{Y^T Y - \hat{\beta}_1^T X_1^T Y}{n-p}$$
(45)

Traditionally, the way the F-test is summarized is:

Table: ANOVA table

Source of variance	df	Sum of squares	Mean square
Model with eta_0	q	$\beta_0^T X_0^T Y$	
Improvement due to	p-q	$\hat{\beta}_1 X_1^T Y - \hat{\beta}_0^T X_0^T Y$	$\frac{\hat{\beta}_1 X_1^T Y - \hat{\beta}_0^T X_0^T Y}{p - q}$
eta_1			
Residual	n-p	$Y^TY - \hat{\beta}_1^T X_1^T Y$	$\frac{Y^TY - \hat{\beta}_1^T X_1^T Y}{n - p}$
Total	n	y^Ty	

There is much more to say here about ANOVA, but this is the basic idea.

Practically speaking, the usage is simple:

```
anova(m0,m1)
## Analysis of Variance Table
##
## Model 1: y ~ 1
## Model 2: y ~ x
## Res.Df RSS Df Sum of Sq F Pr(>F)
## 1 9 1545
## 2 8 897 1 648 5.78 0.043
```

Here, the F-statistic tells us that the model m1 is "better" (there is a significant effect of the predictor x).

Whether that is a meaningful result depends on the power properties of the design (to be discussed later).