Lecture 10

September 25, 2018

A common model for when you have categorical predictors is

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We could rewrite this to look slightly more familiar,

$$y_i = \mu_1 I(i \in 1st) + \ldots + \mu_J I(i \in Jth) + \epsilon_i$$

= $\mu_1 x_{1i} + \ldots + \mu_J x_{Ji} + \epsilon_i$

where x_{ji} is an indicator variable as to whether observation i is in the jth group.

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where x_{jj} is an indicator variable as to whether observation i is in the jth group. We can call the predictor variable X_j a **dummy variable**, in that it gives 0/1 to whether it is in a group or not.

► We then get an **X** matrix

Why don't we get multiple model for each group?

If all groups come from the same population, then variance are the same for each group.

But if we have multiple models, then we assume each model has different vairance $\mathbf{X} = \begin{bmatrix}
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where within each group j, we have the same vector of predictors \mathbf{x}_i^T repeated n_i times.

Note that the $\hat{\mu}_j$ that solve the least-squares solution is just the mean of the observations in the group, which is intuitive.

```
Call:
lm(formula = coag ~ diet - 1, data = coagulation)
Residuals:
Min 1Q Median 3Q Max
-5.00 -1.25 0.00 1.25 5.00
Coefficients:
      Estimate Std. Error t value Pr(>|t|)
dietA 61.0000 1.1832 51.55 <2e-16 ***
dietB 66.0000 0.9661 68.32 <2e-16 ***
dietC 68.0000 0.9661 70.39 <2e-16 ***
dietD 61.0000 0.8367 72.91 <2e-16 ***
Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0
Residual standard error: 2.366 on 20 degrees of freedom
Multiple R-squared: 0.9989, Adjusted R-squared:
F-statistic: 4399 on 4 and 20 DF, p-value: < 2.2e-16
```

A B C D 61 66 68 61

[1] 2.366432

▶ Separate analysis If all we are going to do is estimate the mean, we could ask why put them in a linear model. We could just take the mean per group, which as we've seen gives the same answer, and then calculate SE for them.

A B C D
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 0.9128709 1.1547005 0.6831301 0.9258201
- Note that this is not the same estimate of SE that we got from the linear model. In particular, this allows each group to have a separate variance. Our linear model finds the same estimate of σ^2 for all observations. And then the variance of an estimate is given by $\hat{\sigma}^2(\mathbf{X}^T\mathbf{X})^{-1}$.

X'X

dietA dietB dietC dietD

dietA 4 0 0 0

dietB 0 6 0 0

dietC 0 0 6 0

dietD 0 0 0 8

•	Diagonal entries are numbers of samples in each group				
	dietA	dietB	dietC	dietD	
	dietA	4	0	0	0
	dietB	0	6	0	0
	dietC	0	0	6	0
	dietD	0	0	0	8

Then we see that our estimate of SE in both cases is $\hat{\sigma}_j/n_j$, but when we use the linear model, then we assume that σ_j is the same for all groups.

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where e_{ij} are i.i.d normal random variables with mean zero and variance σ^2 . Let $\sum_{i=1}^t n_i = n$.

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Consider different school as subjects, and students as treatment for each school. We want to know the performance for each school taking an Standard Test

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 - 2. We are looking at some performance of n subjects who can naturally be divided into t groups. We would like to see if the performance difference between the subjects can be explained by the fact that there in these different groups. y_{i1},..., y_{ini} denote the performance of the subjects in the ith group.

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where μ is called the baseline score and τ_i is the difference between the average score for the *i*th treatment and the baseline score.

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- Because of this lack of estimability, people often impose the condition $\sum_{i=1}^{t} \tau_i = 0$. This condition ensures that all parameters μ and τ_1, \ldots, τ_t are estimable.
- Moreover, it provides a nice interpretation. μ denotes the baseline response value and τ_i is the value by which the response value needs to be adjusted from the baseline μ for the group i. Because $\sum_i \tau_i = 0$, some adjustments will be positive and some negative but the overall adjustment averaged across all groups is zero.

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- what is the RSS in the reduced model (*m*). What is the RSS in the full model? Let $\bar{y}_i = \sum_{j=1}^{n_i} y_{ij}/n_i$ and $\bar{y} = \sum_{j=1}^{t} \sum_{i=1}^{n_i} y_{ij}/n$.

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- $y = \sum_{i=1}^{n} \sum_{j=1}^{n} y_{ij}/r$ Write

$$\sum_{i=1}^{t} \sum_{i=1}^{n_i} (y_{ij} - \tilde{\mu}_i)^2 = \sum_{i=1}^{t} \sum_{i=1}^{n_i} (y_{ij} - \bar{y}_i + \bar{y}_i - \tilde{\mu}_i)^2$$

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Write

$$\begin{split} \sum_{i=1}^{t} \sum_{j=1}^{n_{i}} \left(y_{ij} - \bar{\mu}_{i} \right)^{2} &= \sum_{i=1}^{t} \sum_{j=1}^{n_{i}} \left(y_{ij} - \bar{y}_{i} + \bar{y}_{i} - \bar{\mu}_{i} \right)^{2} \\ &= \sum_{i=1}^{t} \sum_{j=1}^{n_{i}} \left(y_{ij} - \bar{y}_{i} \right)^{2} + 2 \sum_{i=1}^{t} (\bar{y}_{i} - \bar{\mu}_{i}) \sum_{j=1}^{n_{i}} \left(y_{ij} - \bar{y}_{i} \right) + \sum_{i=1}^{t} n_{i} (\bar{y}_{i} - \bar{\mu}_{i})^{2} \end{split}$$

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- $\bar{y} = \sum_{i=1}^{t} \sum_{j=1}^{n_i} y_{ij}/n.$ Write

$$\sum_{i=1}^{t} \sum_{j=1}^{n_{i}} (y_{ij} - \tilde{\mu}_{i})^{2} = \sum_{i=1}^{t} \sum_{j=1}^{n_{i}} (y_{ij} - \bar{y}_{i} + \bar{y}_{i} - \tilde{\mu}_{i})^{2}$$

$$= \sum_{i=1}^{t} \sum_{j=1}^{n_{i}} (y_{ij} - \bar{y}_{i})^{2} + 2 \sum_{i=1}^{t} (\bar{y}_{i} - \tilde{\mu}_{i}) \sum_{j=1}^{n_{i}} (y_{ij} - \bar{y}_{i}) + \sum_{i=1}^{t} n_{i} (\bar{y}_{i} - \tilde{\mu}_{i})^{2}$$

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▶ Thus the *F*-statistic for testing $H_0: \mu_1 = \cdots = \mu_t$ is

$$T = \frac{\sum_{i=1}^{t} n_i (\bar{y}_i - \bar{y})^2 / (t-1)}{\sum_{i=1}^{t} \sum_{i=1}^{n_i} (y_{ii} - \bar{y})^2 / (n-t)}$$

which has the F-distribution with t-1 and n-t degrees of freedom under H_0 .

Confidence Intervals for β_i

▶ Because $\hat{\beta} \sim N(\beta, \sigma^2(X^TX)^{-1})$, we have $\hat{\beta}_j \sim N(\beta_j, \sigma^2v_j)$ where v_j is the corresponding diagonal entry of $(X^TX)^{-1}$.

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b But σ is not known, so we use the fact that

$$\frac{\hat{\beta}_j - \beta_j}{\hat{\sigma}\sqrt{\mathbf{v}_j}} \sim t_{n-p-1}$$

to construct the following 100(1 $-\alpha$) % C.I for β_i :

$$\hat{\beta}_j \pm t_{n-p-1}^{\alpha/2} \hat{\sigma} \sqrt{v_j}.$$

Confidence Intervals for β_j

▶ Because $\hat{\sigma}\sqrt{v_j}$ is the standard error for $\hat{\beta}_j$, we can write this C.I as

$$\hat{\beta}_j \pm t_{n-p-1}^{\alpha/2} s.e(\hat{\beta}_j).$$

If this interval contains the value 0, it means that the hypothesis $H_0: \beta_i = 0$ will not be rejected at the α level.

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- Our linear model says that the response for this new subject will be $y_0 = \beta_0 + \beta_1 x_{01} + \cdots + \beta_p x_{0p} + e_0$.

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 - 1. Interval for the mean response without noise
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Chi-square sigma hat

Therefore

$$x_0^T \hat{\beta} \pm t_{n-p-1}^{(\alpha/2)} \hat{\sigma} \sqrt{x_0^T (X^T X)^{-1} x_0}$$
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- ▶ It is easy to see that $y_0 \hat{y_0}$ has a normal distribution with mean zero and variance $\sigma^2 \left(1 + x_0^T (X^T X)^{-1} x_0\right)$. Therefore

$$\frac{y_0 - \hat{y_0}}{\hat{\sigma}\sqrt{1 + x_0^T(X^TX)^{-1}x_0}} \sim t_{n-p-1}$$

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- We can hope to reduce the estimation error by using a lot of data, but we still have to allow for the variablility in the observations while constructing the prediction interval. The latter component does not depend on *n*.