THE ANALYSIS OF HIERARCHICAL AND OTHER DEPENDENT DATA

Lecture Notes

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(With thanks to Bianca L. De Stavola)

LSHTM MSc in Medical Statistics 2022-23

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Outline

Each session in this module starts with a lecture, followed by a practical based on the material presented. There will be a mixture of face-to-face lectures and practical sessions, recorded lectures accessed via moodle and self guided practical sessions with facilitators to answer questions.

There is a pre-course session to revise linear regression.

There is a face-to-face guest lecture in session 13 which accounts for 10% of the assignment mark and practical 14 is an optional Q&A session.

Session	Date		Title	Lecturers
0	Pre-course		Revision of linear regression modelling Access lecture and practical via moodle	L Sharples
1	20 Feb	am	Introduction and simple treatment of dependent data	L Sharples
2		pm	The random intercept model	J Carpenter
3	21 Feb	am pm	Study time - no lecture The random intercept model with covariates	- L Sharples
4 4 cont	27 Feb	am pm	The random coefficients model Hierarchical models overview	J Carpenter L Sharples
5 6	28 Feb	am pm	Longitudinal data I Longitudinal data II	L Sharples J Carpenter
7 8	6 Mar	am pm	Longitudinal data III Generalized Estimating Equations	L Sharples J Carpenter
9 10	7 Mar	am pm	Further issues and summary Revision and Assignment	L Sharples L Sharples
11 12	13 Mar	am pm	Missing Data I Missing Data II	M Quartagno M Quartagno
13 14	14 Mar	am pm	Guest lecture and discussion Optional Q & A session	Halima Twabi L Sharples & J Carpenter
	21 Mar	am/pm	Study time - assignment preparation	
	22 Mar	$5\mathrm{pm}$	Assignment deadline	

References on which the course is based

Rabe-Hesketh, S. and Skrondal, A. (2012) Multilevel and Longitudinal Modeling Using Stata, 3rd Edition. Stata Press.

Snijders, T. and Bosker, R. (1999) Multilevel Analysis SAGE Publications Ltd.

Verbeke, G. and Molenberghs, G. (2000) Linear Mixed Models for Longitudinal Data. Springer Verlag.

Fitzmaurice, G.M., Laird, N.M., and Ware, J.H. (2011) Applied Longitudinal Analysis. 2nd edition. John Wiley and Sons, New York.

For a simple introductory text

Kreft, I. and de Leeuw, J. (1998) Introducing multilevel modelling. London: SAGE.

Other references

Diggle, P.J., Heagerty, P., Liang, K.-Y. and Zeger, S.L. (2002) Analysis of Longitudinal Data, Second Edition Oxford University Press.

Dwyer, J.H., Feinleib, M., Lippert, P. and Hoffmeister, H. eds (1990) Statistical Methods for Longitudinal Studies of Health. Oxford University Press.

Goldstein, H. (2011) Multilevel Statistical Models, Fourth Edition. Arnold, London.

Jones, B. and Kenward, M.G. (2003) The Design and Analysis of Cross-Over Trials. Second Edition. CRC/Chapman & Hall.

Longford, N.T. (1993) Random Coefficient Models. Oxford University Press.

Molenberghs, G. and Kenward, M.G. (2007) Missing data in Clinical Studies. Wiley.

Chapter 0

Revision of linear regression modelling

In this pre-course session, we will revise linear regression modelling, with emphasis on analysis of variance. In particular, for the module it is important that you have a clear idea of the difference between conditional and marginal models.

0.1 Revision of the linear regression model

A brief revision of the main features and assumptions of the linear regression model is given here using data from a survey of 500 mothers who had singleton births in a large London hospital. The survey included information on the age of the mothers and sex, birth weight and gestational age of their babies. In this session we shall find whether boys and girls differ in terms of mean birth weight and quantify the effect of gestational period on birth weight.

0.1.1 t-test and ANOVA

Let (Y_i, X_i) be the dependent variable and explanatory variable of interest, with Y_i representing birth weight and X_i the sex of baby i, i = 1, ..., N = 500.

We can formally compare mean birth weight in the two sexes with a t-test (in the dataset sex is coded 1 for boys and 2 for girls), assuming that the two groups have the same **population variance**:

- . use births, clear
- . ttest bweight, by(sex)

Two-sample t test with equal variances

Group	Obs	Mean	Std. Err.	Std. Dev.
1 2	264 236	3229.902 3032.831	38.99802 40.80225	633.6428 626.816
combined	500	3136.884	28.5077	637.4515
diff		197.071	56.47605	

<OMITTED OUTPUT>

The sample mean birth weights in boys and girls, $\hat{\mu_1}$ and $\hat{\mu_2}$ are, respectively, 3230g and 3033g and

N=500. The t-test of whether the true means μ_1 and μ_2 are the same is

$$t = \frac{\hat{\mu}_1 - \hat{\mu}_2}{\hat{SE}(\hat{\mu}_1 - \hat{\mu}_2)} = \frac{197.071}{56.476} = 3.4895$$

where SE is the standard error of the difference in estimated means. The relevant part of the Stata output is:

Under the null hypothesis of no difference between the means, the statistic has a t distribution with df = N - J = 500 - 2 = 498 degrees of freedom, where J = 2 is the number of estimated means (i.e. the number of groups being compared).

The model underlying this t-test is also called the one-way analysis of variance (ANOVA) model. Analysis of variance involves partitioning the total sum of squares (TSS) (i.e. the sum of squared deviations of the Y_i from their overall mean) into the model sum of squares (MSS) and the residual sum of squares (RSS):

$$TSS = \sum_{i=1}^{N} (Y_i - \overline{Y})^2 = MSS + RSS$$

where

MSS =
$$\sum_{i=1}^{N} (\hat{Y}_i - \overline{Y})^2$$
=
$$\sum_{i \in I} (\overline{Y}_1 - \overline{Y})^2 + \sum_{i \in 2} (\overline{Y}_2 - \overline{Y})^2$$
=
$$n_1 (\overline{Y}_1 - \overline{Y})^2 + n_2 (\overline{Y}_2 - \overline{Y})^2$$

and

RSS =
$$\sum_{i=1}^{N} (Y_i - \hat{Y}_i)^2$$

= $\sum_{i \in 1} (Y_i - \overline{Y}_1)^2 + \sum_{i \in 2} (Y_i - \overline{Y}_2)^2$

The model mean square (MMS) and the mean square error (MSE) can be obtained from the corresponding sums of squares by dividing them by the appropriate degrees of freedom as described in the table below:

Source	Sum of Squares	DF	Mean Square
Model	MSS	J-1	MMS
Residual	RSS	N-J	MSE
Total	TSS	N-1	

J = number of groups; N = Total sample size; DF: degrees of freedom

The MSE is the pooled within-group sample variance, and is an estimate of the residual population variance σ^2 .

The F statistic for the null hypothesis that the population means are the same is then

$$F(J-1, N-J) = \frac{\text{MMS}}{\text{MSE}}$$

Under the null hypothesis this statistic has an F distribution with (J - 1, N - J) degrees of freedom. When J=2 the F statistic is the square of the t statistic (assuming equal variances).

We can perform an ANOVA of birth weight (measured in kg) in terms of sex using Stata, either with the command anova (or equivalently oneway) and obtain $F = 12.18 = t^2 = (3.4895)^2$:

. anova bweight sex

Number of obs	s = 500	R-squared =	0.0239
Root MSE	= 630.431	Adj R-squared =	0.0219

	Partial SS		MS	F	Prob F
Model sex Residual	4839398.61 4839398.61	1 1 498	4839398.61 4839398.61 397442.68	12.18 12.18	0.0005 0.0005
Total			406344.395		

The entries for the rows headed Model and sex are identical because only one explanatory variable is considered here.

0.1.2 Simple Linear Regression

Now we will use linear regression to model birth weight (measured in kg) in terms of a continuous explanatory variable, gestational period (measured in weeks), where Y_i is again birth weight and X_i is gestational period of baby i (Figure 0.1.2).

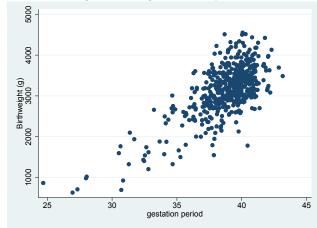


Figure 1: Birth weight versus gestational period of London babies

A simple linear regression model of Y on X can be written:

$$Y_i = \beta_0 + \beta_1 X_i + \epsilon_i$$

where i = 1, ..., N and the residuals ϵ_i are independent with

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

Alternatively we can say that the random variable Y_i has the conditional distribution

$$Y_i \mid X_i \sim N(\beta_0 + \beta_1 X_i, \sigma^2).$$

Here β_0 is the intercept and represents the conditional expectation of Y_i when $X_i = 0$

$$E(Y_i \mid X_i = 0) = \beta_0$$

and β_1 is the slope and represents the difference in conditional expectations when X_i increases by one unit, for example from a to a + 1:

$$E(Y_i \mid X_i = a+1) - E(Y_i \mid X_i = a) = (\beta_0 + \beta_1(a+1)) - (\beta_0 + \beta_1a) = \beta_1$$

Hence the implicit assumption of this model is that the conditional expectations of every baby fall on a straight line: $E(Y_i \mid X_i) = \beta_0 + \beta_1 X_i$ (this is the assumption of linearity). It implies that residuals have mean zero, conditional on X_i (i.e. $E(\epsilon_i \mid X_i) = 0$) and also that $Cor(\epsilon_i, X_i) = 0$ (assumption of 'exogeneity' of the covariate). In addition the model assumes that residuals have constant variance σ^2 (assumption of 'homoscedasticity').

As before we can partition the TSS into MSS and RSS, with

$$\hat{Y}_i = \hat{\beta}_0 + \hat{\beta}_1 X_i$$

The OLS estimates for β_0 and β_1 are found by minimizing the RSS while again the estimate of σ^2 is found by dividing the RSS by N-2, 2 being the number of estimated parameters.

ML estimation is achieved via the log-likelihood function:

$$\ell(\beta_0, \beta_1, \sigma^2 \mid \mathbf{Y}, \mathbf{X}) = -\frac{n}{2} \ln(2\pi\sigma^2) - \frac{1}{2\sigma^2} \sum_{i=1}^{N} (Y_i - \beta_0 - \beta_1 X_i)^2.$$

where **Y** is the $(N \times 1)$ dependent variable vector and **X** the $(N \times 1)$ explanatory variable vector for the N babies.

The MLE's of β_0 and β_1 can be obtained from the **score equations**:

$$U(\beta_0) = \ell'(\beta_0) = \frac{1}{\sigma^2} \sum_{i=1}^{N} (Y_i - \beta_0 - \beta_1 X_i)$$

$$U(\beta_1) = \ell'(\beta_1) = \frac{1}{\sigma^2} \sum_{i=1}^N X_i (Y_i - \beta_0 - \beta_1 X_i).$$

Solving these, i.e. setting $U(\hat{\beta}_0) = 0$ and $U(\hat{\beta}_1) = 0$ imply:

$$\hat{\beta}_0 = \overline{Y} - \hat{\beta}_1 \overline{X}$$

$$\hat{\beta}_1 = \frac{\sum X_i Y_i - N \overline{XY}}{\sum X_i^2 - N \overline{X}^2}$$

They are the same as the OLS estimates. The MLE for σ^2 instead is the RSS divided by N, not N-2. In Stata we can fit a linear regression model with:

. regress bweight gestwks

Source	SS	df	MS	Num	ber of	obs =	490
				F(1, 4	188) =	502.36
Model	101603845	1	101603845	Pro	b > F	=	0.0000
Residual	98698697.8	488	202251.43	R-s	quared	=	0.5073
				Adj	R-squa	ared =	0.5062
Total	200302543	489	409616.652	J	-		449.72
bweight	Coef.	Std. E	Err. t	P> t	 [95%	Conf.	Interval

bweight	Coef.	Std. Err.	t	P> t	[95% Conf.	Interval]
gestwks	196.9726 -4489.14	8.788133			179.7054 -5158.95	

The regression parameter labelled gstwks is the estimated slope coefficient, and represents the increase in birth weight predicted by the fitted model for each unit increase in gestational period, in this case one week. The other parameter, _cons, is the intercept, the average weight at period 0 given by the fitted model. This is an impossible negative value, as it gives the expected birth weight in a baby whose gestation is 0 weeks!

gestational period: Inted regression line with point with point gestation period and gestation period special p

Figure 2: Birth weight and gestational period: fitted regression line with pointwise confidence interval

The ANOVA table, reproduced below for clarity, provides a breakdown of the variability of the data. The Model Mean Square represents the variability in the data that is explained by the fitted line, the Residual Mean Square (also called *Mean Square Error*, MSE) quantifies the remaining variability that is NOT explained by the model. The estimated population residual SD σ is $\sqrt{98698697.8/488} = \sqrt{202251.43} = 449.72$, i.e. around 450g.

Source	Sum of Squares	DF	Mean Square	F
Model	101,603,845	1	101,603,845	502.36
Residual	9,869,869,7.8	488	202,251.43	
Total	200,302,543	489		

The fitted model (a straight line) is plotted together with the pointwise confidence interval in Figure 2 with the command:

. twoway (scatter bweight gestwks) (lfitci bweight gestwks)

Note how wide the confidence intervals become at the extremes of the range of X values. Indeed at X=0 the confidence interval goes from -5158.95 to -3819.329. Thus, when the data on the explanatory variable do not include the value 0 as in this case, it is best to centre the explanatory variable around its mean value (or some other sensible value) and refit the model:

$$Y_i = \beta_0 + \beta_1 (X_i - \hat{\mu}_X) + \epsilon_i$$

In Stata:

. su gestwks

Variable	0bs	Mean	Std.	Dev.	Min	Max

gestwks | 490 38.72186 2.314167 24.69 43.16

- . gen c_gest=gestwks-r(mean)
- . regress bweight c_gest

<EDITED OUTPUT>

bweight		Std. Err.				Interval]
c_gest	196.9726	8.788133 20.31645	22.41	0.000	179.7054 3098.088	214.2399 3177.925

The ANOVA table and the estimated slope (and corresponding SE) obtained when fitting this last model have not changed (beside minimal differences in the ANOVA table due to rounding error). The only change is in the estimated intercept that now refers to the expected weight of a baby who was 38.7 weeks at birth and which is now more precisely estimated.

0.1.3 Adding covariates and interactions

Now we consider whether the effect of gestational period on birth weight is in anyway confounded or modified by the sex of the baby. We first add sex to the linear model to see whether the coefficient for gestational age c_gest changes.

. regress bweight c_gest i.sex <OMITTED OUTPUT>

bweight	Coef.	Std. Err.	t	P> t	[95% Conf.	Interval]
c_gest	196.3436	8.599464	22.83	0.000	179.4469	213.2402
2.sex	-189.9113	39.8007	-4.77	0.000	-268.1136	-111.709
_cons	3228.698	27.50261	117.40	0.000	3174.66	3282.737

The estimate of the slope obtained from the simple model indicated a 197g increase in birth weight per gestational week. This does not appear to be confounded by sex as the sex-adjusted estimate of the slope is 196g per week of gestation. The intercepts in boys and girls are however significantly different, with girls being on average 190g lighter than boys for any given gestational period.

Now add the interaction between sex and gestational age, then test for modification as follows:

. regress bweight i.sex##c.c_gest

bweight		Std. Err.	t	P> t	[95% Conf.	Interval]
2.sex c_gest 	-189.8624	39.82193 13.26672	-4.77 15.33	0.000	-268.1068 177.2909	-111.618 229.4253
sex#c.c_gest 2	-12.10667	17.42919	-0.69	0.488	-46.35255	22.1392
_cons	3228.461	27.51936	117.32	0.000	3174.39	3282.533

. testparm i.sex#c.cgest

(1)
$$2.sex\#c.cgest = 0$$

F(1, 486) = 0.48

$$Prob > F = 0.4876$$

The coefficient c_gest is now interpreted as the effect of gestational age on birth weight for boys. The interaction term estimates that the girls' growth rate is 12g lower than boys, but according to the interaction test, there is no evidence of effect modification (Partial F-test: P=0.49 for the interaction between sex and gestational age). Hence it is likely that the birth weight of boys and girls has the same relation with gestational period.

0.1.4 Non linear effects

It may be unreasonable to assume that birth weight increases linearly with gestational age. To test whether the relationship is quadratic instead of linear we generate a new variable equal to the square of X and add it to the model. Because we are centering gestational age, the model we are considering is:

$$Y_i = \beta_0 + \beta_1 (X_i - \hat{\mu}_X) + \beta_2 (X_i - \hat{\mu}_X)^2 + \beta_3 \text{ female} + \epsilon_i$$
 (1)

In Stata:

- . gen cgestsq=c_gest^2
- . regress bweight c_gest cgestsq i.sex

<EDITED OUTPUT>

bweight		Coef.	Std. Err.	t	P> t
c_gest		184.9798	12.20059	15.16	0.000
cgestsq		-2.260747	1.723046	-1.31	0.190
2.sex		-185.7864	39.89531	-4.66	0.000
_cons		3238.811	28.54257	113.47	0.000

There does not seem to be evidence of departure from linearity of the effect of gestational period.

0.1.5 Residual diagnostics

We can estimate residuals from a fitted model as the differences between observed and predicted values of Y. For the last fitted model these are:

$$\hat{\epsilon}_i = Y_i - (\hat{\beta}_0 + \hat{\beta}_1 (X_i - \hat{\mu}_X) + \hat{\beta}_2 (X_i - \hat{\mu}_X)^2)$$

The estimated standardized residuals are defined as

$$\hat{r}_i = \frac{\hat{\epsilon}_i}{\sqrt{\hat{\sigma}^2}}$$

They are often referred to as 'Pearson residuals'.

If the model is correct, these residuals are normally distributed with constant variance and therefore can be used to assess the model's assumptions. In Stata:

- . predict r, rst
- . hist r, normal
- . count if r>3 | r<-3

The histogram of the standardized residuals show no departure from the normality assumption, nor is there evidence of too many outliers (out of 500):

Figure 3: Birth weight data: histogram of standardized residuals from model (1)

