Kenward-Roger modification of the F-statistic for some linear mixed models fitted with Imer

Ulrich Halekoh ¹ Søren Højsgaard ²

¹Department of Molecular Biology and Genetics Aarhus University, Denmark *ulrich.halekoh@agrsci.dk*

²Department of Mathematical Sciences Aalborg University, Denmark sorenh@math.aau.dk

9. november 2011

- Outline
 - Motivation: Sugar beets A split-plot experiment
 - Motivation: A random regression problem
 - Our goal
- 2 The Kenward–Roger modification of the F–statistic
- Parametric bootstrap
- 4 Small simulation study: A random regression problem
- 5 Final remarks

- Dependence of sugar percentage of sugar beets on harvest time and sowing time is investigated.
- Five sowing times (s) and two harvesting times (h).
- Experiment was laid out in three blocks (b).

- For simplicity we assume that there is no interaction between sowing and harvesting times.
- A typical model for such an experiment would be:

$$y_{hbs} = \mu + \alpha_h + \beta_b + \gamma_s + U_{hb} + \epsilon_{hbs}, \tag{1}$$

where $U_{hb} \sim N(0, \omega^2)$ and $\epsilon_{hbs} \sim N(0, \sigma^2)$.

• Notice that U_{hb} describes the random variation between whole–plots (within blocks).

Motivation: Sugar beets - A split-plot experiment

As the design is balanced we may make F-tests for each of the effects as:

```
R-code
> beets$bh <- with(beets, interaction(block, harvest))</pre>
> summary(aov(sugpct~block+sow+harvest+Error(bh), beets))
Error: bh
         Df Sum Sq Mean Sq F value Pr(>F)
block 2 0.0327 0.0163 2.58 0.28
harvest 1 0.0963 0.0963 15.21 0.06
Residuals 2 0.0127 0.0063
Error: Within
         Df Sum Sq Mean Sq F value Pr(>F)
SOW
         4 1.01 0.2525 101 5.7e-13
Residuals 20 0.05 0.0025
```

Notice: the F-statistics are $F_{1,2}$ for harvest time and $F_{4,20}$ for sowing time.

Motivation: Sugar beets - A split-plot experiment

Using lmer() from lme4 we can fit the models and test for no effect of sowing and harvest time as follows:

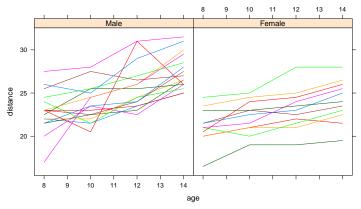
```
R-code
> beetLarge < - lmer(sugpct block+sow+harvest+(1|block:harvest),
             data=beets, REML=FALSE)
> beet_no.harv <- update(beetLarge, .~.-harvest)
> beet_no.sow <- update(beetLarge, .~.-sow)</pre>
> as.data.frame(anova(beetLarge, beet_no.sow))
                 AIC BIC logLik Chisq Chi Df Pr(>Chisq)
beet no.sow 6 -2.795 5.612 7.398 NA
                                             NΑ
beetLarge 10 -79.997 -65.985 49.999 85.2 4 1.374e-17
> as.data.frame(anova(beetLarge, beet_no.harv))
                  AIC BIC logLik Chisq Chi Df Pr(>Chisq)
beet_no.harv 9 -69.08 -56.47 43.54 NA
beetLarge 10 -80.00 -65.99 50.00 12.91 1 0.0003262
```

The LRT based *p*-values are anti-conservative: the effect of harvest appears stronger than it is.

Motivation: A random regression problem

Random coefficient model

The change with age of the distance between two cranial distances was observed for 16 boys and 11 girls from age 8 until age 14.



Random coefficient model

Plot suggests:

$$dist_{[i]} = \alpha_{sex[i]} + \beta_{sex[i]} age_{[i]} + A_{Subj[i]} + B_{Subj[i]} age_{[i]} + e_{[i]}$$
 with $(A, B) \sim N(0, \mathbf{S})$.

ML-test of $\beta_{boy} = \beta_{girl}$:

```
R-code
```

Our goal...

Our goal is to extend the tests provided by lmer().

There are two issues here:

- The choice of test statistic and
- The reference distribution in which the test statistic is evaluated.

Setting the scene

For multivariate normal data

$$Y_{n\times 1} \sim N(\mathbf{X}_{n\times p}\beta_{p\times 1}, \mathbf{\Sigma})$$

we consider the test of the hypothesis

$$\mathbf{L}_{I \times p} \boldsymbol{\beta} = \boldsymbol{\beta}_0$$

where **L** is a regular matrix of estimable functions of β .

The linear hypothesis can be tested via the Wald-type statistic

$$F = rac{1}{l}(\hat{oldsymbol{eta}} - oldsymbol{eta}_0)^{ op} \mathbf{\mathsf{L}}^{ op} (\mathbf{\mathsf{L}}^{ op} \mathbf{\Phi}(\hat{oldsymbol{\sigma}}) \mathbf{\mathsf{L}})^{-1} \mathbf{\mathsf{L}} (\hat{oldsymbol{eta}} - oldsymbol{eta}_0)$$

- ullet $\Phi(\sigma) = (\mathbf{X}^{ op} \mathbf{\Sigma}(\sigma) \mathbf{X})^{-1} pprox \mathbb{C}$ ov (\hat{eta}) , \hat{eta} REML estimate of eta
- $\hat{\sigma}$: vector of REML estimates of the elements of Σ

Kenward and Roger's modification

Kenward and Roger (1997) modify the test statistic

ullet Φ is replaced by an improved small sample approximation $\Phi_{\mathcal{A}}$

Furthermore

- the statistic F is scaled by a factor λ ,
- denominator degrees of freedom m are determined

such that the approximate expectation and variance are those of a $F_{l,m}$ distribution.

Restriction on covariance

Consider only situations where

$$\Sigma = \sum_{i} \sigma_{i} \mathbf{G}_{i}, \quad \mathbf{G}_{i}$$
 known matrices

- Variance component and random coefficient models satisfy this restriction.
- ullet $\Phi_A(\hat{oldsymbol{\sigma}})$ depends now only on the first partial derivatives of Σ^{-1} :

$$\frac{\partial \Sigma^{-1}}{\partial \sigma_i} = -\Sigma^{-1} \frac{\partial \Sigma}{\partial \sigma_i} \Sigma^{-1}.$$

- $\Phi_A(\hat{\sigma})$ depends also on $\mathbb{V}ar(\hat{\sigma})$.
- Kenward and Roger propose to estimate $Var(\hat{\sigma})$ via the inverse expected information matrix.

Properties of the Kenward-Roger adjustment

The modification of the F-statistic by Kenward and Roger

- yields the exact F-statistic for balanced mixed classification nested models or balanced split plot models (Alnosaier, 2007).
- Simulation studies (e.g. Spilke, J. et al.(2003)) indicate that the Kenward-Roger approach perform mostly better than alternatives (like Satterthwaite or containment method) for blocked experiments even with missing data.

R package 1me4

The R package Ime4 (Bates, D., Maechler, M, Bolker, B., 2011) provides efficient estimation of linear mixed models.

The package provides all necessary matrices and estimates to implement the Kenward-Roger approach.

- The implementation uses a straightforward transcription of the description in the article of Kenward and Roger, 1997.
- Matrix operations use sparse matrices representation.
- Matrices are extracted from lmer objects via their slots (using @).

Kenward-Roger: split-plot (sugar-beets)

The Kenward–Roger approach yields the same results as the anova-test:

R-code

```
> beetLarge <- update(beetLarge, REML=TRUE)
```

> beet_no.harv <- update(beet_no.harv, REML=TRUE)</pre>

Test for harvest effect:

R-code

```
> KRmodcomp(beetLarge,beet_no.harv)
F-test with Kenward-Roger approximation
large : sugpct ~ block + sow + harvest + (1 | block:harvest)
small : sugpct ~ block + sow + (1 | block:harvest)
Fstat df1 df2 p.value F.scaling
15.21 1 2 0.0599 1
```

Kenward-Roger: random regression (cranial change)

For the cranial distances data the Kenward and Roger modified F-test yields

```
R-code

> ort1<- update(ort1ML, .~., REML = TRUE)
> ort2<- update(ort2ML, .~., REML = TRUE)
> KRmodcomp(ort1, ort2)

F-test with Kenward-Roger approximation
large: distance ~ age + Sex + (1 + age | Subject) + age:Sex
small: distance ~ age + Sex + (1 + age | Subject)

Fstat df1 df2 p.value F.scaling
5.118 1 25 0.0326 1
```

The p-value form the ML-test was 0.0249.

Using parametric bootstrap

- The Kenward-Roger approach is no panacea.
- Additionally, we provide the parametric bootstrap p-value $P_{\hat{\theta}_0}(T \geq t_{obs})$ based on the log-LR statistic T.

We draw B parametric bootstrap samples t^1,\ldots,t^B under the estimated null model $\hat{\theta}_0$ and provide three choices to calculate the p-value.

- **1** directly via the proportion of sampled t_i exceeding t_{obs} ,
- ② approximating the distribution of the scaled statistic $\frac{f}{t} \cdot T$ by a χ_f^2 distribution (Bartlett type correction) (\bar{t} is the sample average and f the difference in the number of parameters between the null and the alternative model)
- **3** approximating the bootstrap distribution by a $\Gamma(\alpha, \beta)$ distribution which mean and variance match the moments of the bootstrap sample.

```
> PBmodcomp(beetLarge,beet_no.harv)
```

R-code

```
large: sugpct ~ block + sow + harvest + (1 | block:harvest)
small: sugpct ~ block + sow + (1 | block:harvest)
Number of parametric bootstrap samples: 200
stat df p.value
LRT 11.815 1 0.0006
PBtest 11.815 NA 0.0550
Bartlett 2.447 1 0.1178
Gamma 11.815 NA 0.0782
```

Results from sugar beets:

Tabel: p-values (\times 100) for removing the harvest or sow effect.

	LRT	KR	ParmBoot	Bartlett	Gamma
harvest	0.03	6	4.1	8.3	4.9
sow	< 0.001	< 0.001	< 0.001	< 0.001	< 0.001

Results for cranial distance data:

Tabel: p-values (\times 100) testing $\beta_{boy} = \beta_{girl}$.

LRT	KR	ParmBoot	Bartlett	Gamma
2.5	3.3	4.2	4.0	4.2

Random coefficient model

We consider the simulation from a simple random coefficient model (cf. Kenward and Roger (1997, table 4)):

$$y_{it} = \beta_0 + \beta_1 \cdot t_i + A_i + B_i \cdot t_i + \epsilon_{it}$$

with
$$cov(A_i, B_i) = \begin{bmatrix} 0.250 & -0.133 \\ -0.133 & 0.250 \end{bmatrix}$$
 and $var(\epsilon_{it}) = 0.25$.

There are observed $i=1,\ldots,24$ subjects divided in groups of 8. For each group observations are at the non overlapping times t=0,1,2; t=3,4,5 and t=6,7,8.

Results from random coefficient model

Tabel: Observed test sizes ($\times 100$) for H_0 : $\beta_k = 0$ for random coefficient model.

	LR	Wald	ParmBoot	Bartlett	Gamma	KR(R)	KR(SAS)
β_0	6.8	8.8	5.6	5.4	5.8	4.0	4.8
β_1	7.1	6.6	5.6	5.4	5.7	5.4	5.0

Summary

- The functions KRmodcomp() and PBmodcomp() described here are available in the pbkrtest package.
- The Kenward–Roger approach requires fitting by REML; the parametric bootstrap approaches requires fitting by ML.
- The required fitting scheme is set by the relevant functions, so the user needs not worry about this.
- Parametric bootstrap is parallelized using the snow package.

Literature

- Alnosaier, W. (2007) Kenward-Roger Approximate F Test for Fixed Effects in Mixed Linear Models, Dissertation, Oregon State University
- Bates, D., Maechler, M. and Bolker, B. (2011) Ime4: Linear mixed-effects models using S4 classes, R package version 0.999375-39.
- Kenward, M. G. and Roger, J. H. (1997) Small Sample Inference for Fixed Effects from Restricted Maximum Likelihood, Biometrics, Vol. 53, pp. 983–997
- Spilke J., Piepho, H.-P. and Hu, X. Hu (2005) A Simulation Study on Tests of Hypotheses and Confidence Intervals for Fixed Effects in Mixed Models for Blocked Experiments With Missing Data Journal of Agricultural, Biological, and Environmental Statistics, Vol. 10,p. 374-389