ECM approaches to heteroskedastic mixed models with constant variance ratios

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Summary – This paper presents techniques of parameter estimation in heteroskedastic mixed models having constant variance ratios and heterogeneous log residual variances that are described by a linear model. Estimation of dispersion parameters is by standard (ML) and residual (REML) maximum likelihood. Estimating equations are derived using the expectation-conditional maximization (ECM) algorithm and simplified versions of it (gradient ECM). Direct and indirect approaches are proposed with the latter allowing hypothesis testing about the variance ratios. The analysis of a small example is outlined to illustrate the theory.

INTRODUCTION

Heteroskedasticity has recently generated much interest in quantitative genetics and animal breeding. To begin with, there is now a large amount of experimental evidence of heterogeneous variances for most important livestock production traits (Garrick et al, 1989; Visscher et al, 1991; Visscher and Hill, 1992). Second, major theoretical and applied work has been carried out for estimating and testing sources...
of heterogeneous variances arising in univariate mixed models (Foulley et al., 1990; Gianola et al., 1992; Weigel et al., 1993; DeStefano, 1994; Foulley and Quaas, 1995). For many reasons (accuracy of estimation, ease of handling large data sets), a major objective in this area lies in making models as parsimonious as possible. This can be accomplished in at least two ways: i) by modelling variances in the case of potentially numerous sources of heteroskedasticity, and ii) by assuming that some functions of those parameters (eg, intra-class correlation or heritability) are constant. The first aspect corresponds to the so-called structural approach in which the heterogeneity of the log components of variances is described via a linear model structure similar to that used for means (Foulley et al., 1990, 1992; San Cristobal, 1993). Restrictions as in ii) were considered by Meuwissen et al. (1996) and Robert et al. (1995a,b). Meuwissen et al. (1996) introduced a multiplicative mixed model to estimate breeding values and heteroskedasticity factors assuming heritability ($h^2$) constant across herd-years. Robert et al. (1995a,b) developed estimation and testing procedures for homogeneity of heritability within and/or genetic correlations across environments. But Meuwissen's study postulates known $h^2$ and Robert's research applies to only a single classification of heteroskedasticity.

The purpose of this paper is to propose a complete inference approach for parameters having both features i) and ii), ie, for continuous data described by mixed models with constant variance ratios and heteroskedasticity analyzed via a structural approach. For simplicity, the theory will be presented using a one-way random mixed model for data and afterwards it will be generalized to several $u$-components. Inference is based on likelihood procedures (REML and ML) and estimating equations derived from the expectation-maximization (EM) theory, more precisely the expectation/conditional maximization (ECM) algorithm recently introduced by Meng and Rubin (1993).

**THEORY**

**Statistical model**

As usual, it is assumed that the population can be structured into strata ($i = 1, 2, ..., I$) corresponding to potential factors of heterogeneity. Let the one-way random model be written as:

$$y_i = X_i \beta + \sigma_{u_i} Z_i u^* + e_i$$  \[1\]

where $y_i$ is the ($n_i \times 1$) data vector for stratum $i$; $\beta$ is a ($p \times 1$) vector of unknown fixed effects with incidence matrix $X_i$, and $e_i$ is the ($n_i \times 1$) vector of residuals. The contribution of random effects is expressed as in Foulley and Quaas (1995) as $\sigma_{u_i} Z_i u^*$ where $u^*$ is a ($q \times 1$) vector of standardized deviations, $Z_i$ is the corresponding incidence matrix and $\sigma_{u_i}$ is the square root of the $u$-component of variance the value of which depends on stratum $i$. Classical assumptions are made for the distributions of $u^*$ and $e_i$, ie, $u^* \sim N(0, A)$, $e_i \sim N(0, \sigma^2_{e_i} I_{n_i})$, and $E(u^* e_i') = 0$.

The notation in [1] is unusual as compared to that used in the statistical literature on mixed effects (eg, Laird et al., 1987). There are practical motivations for such
an expression of the random part especially in animal breeding. For instance the
between sire variance may vary according to the environment in which the progeny
of the sires are raised. Note also that \( \sigma_{ui} \) can be viewed as a regression coefficient of
any element of \( y_i \) on the corresponding element of \( Z_i u^* \). Thus, in animal breeding,
\( \sigma_{ui} \) acts as a scaling factor of a vector \( u^* \) of standardized sire values on which, for
instance, selection can be based.

A structure is hypothesized on the residual variance so as to model the influence
of factors causing heteroskedasticity. This is carried out along the lines presented
in Foulley et al (1990, 1992) via a linear regression on log-variances:

\[
\ln \sigma^2_{ei} = p'_i \delta
\]

where \( \delta \) is an unknown \((r \times 1)\) real-valued vector of parameters and \( p'_i \)
the corresponding \((1 \times r)\) row incidence vector of qualitative or continuous covariates.

Furthermore, the assumption of a constant intra-class correlation (or heritability)
implies setting

\[
\sigma_{ui} = \tau \sigma_{ei}, \; \forall i
\]

**EM-REML estimation**

Use is made here of the EM algorithm of Dempster et al (1977) to compute
REML estimates of parameters involved in variance components (Patterson and
Thompson, 1971; Searle et al, 1992). The basic procedure proposed by Foulley and
Quaas (1995) is applied here after some adjustment of the M-step taking advantage
of the ECM algorithm of Meng and Rubin (1993).

Letting \( y = (y'_1, y'_2, \ldots, y'_i, \ldots, y'_r)' \), \( e = (e'_1, e'_2, \ldots, e'_i, \ldots, e'_r)' \), \( \gamma = (\delta', \tau)' \),
the ECM algorithm is based on a complete data set defined by \( x = (\beta', u'^*, e')' \)
and its log-likelihood \( L(\gamma; x) \). The iterative process takes place as follows.

The E-step is defined as usual, ie, at iteration \([t]\), calculate the conditional
expectation of \( L(\gamma; x) \) given the data \( y \) and \( \gamma = \gamma^{[t]} \)

\[
Q(\gamma | \gamma^{[t]}) = E \left[ L(\gamma; x) | y, \gamma = \gamma^{[t]} \right]
\]

which, as shown in Foulley and Quaas (1995), reduces to

\[
Q(\gamma | \gamma^{[t]}) = \text{const} - \frac{1}{2} \sum_{i=1}^{l} n_i \ln \sigma^2_{ei} - \frac{1}{2} \sum_{i=1}^{l} \sigma^{-2}_{ei} E^{[t]}_c(e'_i e_i)
\]

where \( E^{[t]}_c(\cdot) \) is a condensed notation for a conditional expectation taken with
respect to the distribution of \( x | y, \gamma = \gamma^{[t]} \).

Since the parameters to be estimated are heterogeneous, the estimating equations
are derived at the maximization stage from a slightly different version of the
EM algorithm, the so-called ECM algorithm. As explained in detail in Meng and
Rubin (1993), a CM stage replaces the M-step by a sequence of several conditional
maximization steps. This is basically the same principle as that employed in a cyclic
ascent maximization procedure (Zangwill, 1969). We suggest here the following procedure:

- maximize $Q$ over $\gamma$ to get $\delta^{[t+1]}$ with $\tau$ set at its last value $\tau^{[t]}$, ie

$$\delta^{[t+1]} = \text{Argmax}_\delta Q(\delta, \tau^{[t]}|\gamma^{[t]})$$  \[5\]

- then, maximize $Q$ over $\tau$ to obtain $\tau^{[t+1]}$ with $\delta$ in $\gamma$ of $Q(\gamma|\gamma^{[t]})$ set to $\delta^{[t+1]}$, ie,

$$\tau^{[t+1]} = \text{Argmax}_\tau Q(\delta^{[t+1]}, \tau|\gamma^{[t]})$$  \[6\]

Thus, the maximization step consists of two CM-steps within the same E-step in order to reduce the need to compute the conditional expectation of $e_i^j e_i^j$, and its components more than once. The algebra of differentiation is given in Appendix A. The iterative system for computing formulae $\delta$ can be written as

$$P'W_{\delta}^{[t,\ell]} P \cdot \Delta \delta^{[t,\ell+1]} = P' v^{[t,\ell]}_\delta$$  \[7\]

where

$$\Delta \delta^{[t,\ell+1]} = \delta^{[t,\ell+1]} - \delta^{[t,\ell]}$$

$$P'_{(r \times I)} = (p_1, p_2, \ldots, p_i, \ldots, p_I)$$

$$v^{[t,\ell]}_\delta (I \times 1) = \left \{ v^{[t,\ell]}_{\delta, i} \right \}$$

with the elements of the right-hand side being

$$\nu^{[t,\ell]}_{\delta, i} = \frac{E^{[t]} (y_i - X_i \beta)' e_i^{[t,\ell]} e_i^{[t,\ell]}}{2 e_i^{2[t,\ell]}} - n_i$$  \[8a\]

based on $e_i^{[t,\ell]} = y_i - X_i \beta - \tau^{[t]} e_i^{[t,\ell]} Z_i u^*$, and $e_i^{2[t,\ell]} = \exp(p_i e_i^{[t,\ell]})$

$$W_{\delta}^{[t,\ell]} (r \times r) = \text{Diag} \left \{ w^{[t,\ell]}_{\delta, ii} \right \}$$

with

$$w^{[t]}_{\delta, ii} = \frac{E^{[t]} [(y_i - X_i \beta)' (y_i - X_i \beta)]}{\sigma_{e_i}^{2[t,\ell]}} - \frac{\tau^{[t]} E^{[t]} [(y_i - X_i \beta)' Z_i u^*]}{2 \sigma_{e_i}^{[t,\ell]}}$$  \[8b\]

Note that for this algorithm to be a true ECM, one would have to iterate the NR algorithm in [7] within an inner cycle (index $\ell$) until convergence to the conditional maximizer $\gamma^{[t+\ell]} = \gamma^{[t,c]}$ at each M-step $[t]$. In practice it may be advantageous to reduce the number of inner iterations, even up to only one, ie, by solving just once

$$P'W_{\delta}^{[t]} P \cdot \Delta \delta^{[t]} = P' v^{[t]}_\delta$$  \[9\]

However, caution should be exercised when applying such a hybrid algorithm that no longer guarantees the monotonic convergence in likelihood values (Lange, 1995).
The formula to update $\tau$ reduces to

$$
\tau^{[t+1]} = \frac{\sum_{i=1}^{I} E_{c[i]}^{[t]} \left( u^{*'} Z_i' (y_i - X_i \beta) \right) / \sigma_{e_i}^{[t+1]}}{\sum_{i=1}^{I} E_{c[i]}^{[t]} (u^{*'} Z_i Z_i u^*)}
$$

mimicking the form of a scaled regression coefficient pooled over strata.

The elements to compute at the E-step can be expressed as functions of the sums $X_i' y_i$, $Z_i' y_i$, the sums of squares $y_i'y_i$ within strata, and GLS-BLUP solutions of Henderson's mixed model equations and of their accuracy (Henderson, 1984), ie

$$
\left[ \begin{array}{c}
\sum_{i=1}^{I} X_i' X_i / \sigma_{e_i}^{2[t]} \\
\sum_{i=1}^{I} X_i' Z_i \tau^{[t]} / \sigma_{e_i}^{2[t]} \\
\sum_{i=1}^{I} Z_i' X_i \tau^{[t]} / \sigma_{e_i}^{2[t]} \\
\sum_{i=1}^{I} Z_i' Z_i \tau^{[t]} + A^{-1}
\end{array} \right] \begin{bmatrix}
\hat{\beta} \\
\hat{u}^{*[t]}
\end{bmatrix} = \left[ \begin{array}{c}
\sum_{i=1}^{I} X_i' y_i / \sigma_{e_i}^{2[t]} \\
\sum_{i=1}^{I} Z_i' y_i / \sigma_{e_i}^{2[t]}
\end{array} \right]
$$

where $\hat{u}^{*[t]} = E(u^*|y, \delta^{[t]}, \tau^{[t]}$).

Thus, deleting $[t]$ for the sake of simplicity, one has:

E: $E_c [(y_i - X_i \beta)'(y_i - X_i \beta)] = (y_i - X_i \beta)'(y_i - X_i \beta) + tr(X_i' X_i C_{\beta\beta})$ \[12a\]

E: $E \left[ u^{*'} Z_i' (y_i - X_i \beta) \right] = \tilde{u}^{*'} Z_i' (y_i - X_i \tilde{\beta}) - tr(Z_i' X_i C_{\beta u})$ \[12b\]

E: $E_c (u^{*'} Z_i' Z_i u^*) = \tilde{u}^{*'} Z_i' Z_i \tilde{u}^* + tr(Z_i' Z_i C_{uu})$ \[12c\]

where $\hat{\beta}$ and $\hat{u}^*$ are mixed model equations for $\beta$ and $u^*$, and $C = \begin{bmatrix} C_{\beta\beta} & C_{\beta u} \\ C_{u\beta} & C_{uu} \end{bmatrix}$ is the partitioned inverse of the coefficient matrix.

Expressions in \[12a-c\] can easily accommodate grouped data (see Appendix B).

The close connection between the system of equations \[7\] for residual parameters and formula \[12\] given in Foulley et al (1990) can be observed. There is also a remarkable similarity between formula \[9\] for the ratio and formula \[7\] in Foulley and Quaas (1995). This means that the computations can be implemented with very little change in the code used previously. True or gradient EM could also have been applied (see Appendix A). The advantage of ECM will be more substantial for the next situations considered, and especially in the case of the indirect approach.
Extension to several u-components

Formulae \([7, 8ab] and [9]\) can easily be generalized to a mixed model including several \((k = 1, 2, \ldots, K)\) independent \(u\)-components

\[
y_i = X_i \beta + \sum_{k=1}^{K} \tau_k \sigma_{e_i} Z_{ik} u_k^* + e_i
\]

with \(\tau_k = \sigma_{u_{ik}} / \sigma_{e_i}\) constant over strata \(i\).

Letting \(\mathbf{y} = (\mathbf{d}'', \mathbf{T}')'\) as previously but now with \(\tau = \{\tau_k\}\) being a vector of ratios of standard deviations, the \(Q\) function to be maximized has the same form as in \([4]\) with \(e_i\) expressed from \([13]\). One can perform the CM-steps using either i) the sequence \(\delta, \tau_1, \tau_2, \ldots, \tau_k, \ldots, \tau_K\), ie, each \(\tau_k\) one by one, the remaining ones being held constant, or ii) the sequence \(\delta\), and \(\tau\) as a whole with all the \(\tau_k\)'s maximized jointly. In both cases, the algorithm for computing \(\delta\) is formally the same as in \([7]\) with only a slight change in the definition of the elements of \(\mathbf{W}_{\delta \delta}, \mathbf{v}_{\delta}\) being unchanged

\[
w_{\delta \delta, ii}^{[t, \ell]} = \frac{E_c^{[t]} (y_i - X_i \beta)' (y_i - X_i \beta)}{\sigma_{e_i}^{2[t, \ell]}} - \frac{\sum_{k=1}^{K} E_{c_k}^{[t]} [(y_i - X_i \beta)' Z_{ik} u_k^*]}{2 \sigma_{e_i}^{[t, \ell]}}
\]

If the conditional maximization of the \(\tau_k\)'s takes place one by one (case i), formula \([9]\) still applies for each of them. Otherwise (case ii), one has to solve the following system:

\[
F^{[t]} \tau^{[t+1]} = \mathbf{g}^{[t]}
\]

where

\[
F^{[t]}_{(K \times K)} = \left\{ f_{k \ell}^{[t]} \right\} = \left\{ \sum_{i=1}^{I} E_{c_i}^{[t]} (u_k^* Z_{ik} Z_{i \ell} u_{\ell}^*) \right\}
\]

and

\[
g^{[t]}_{(K \times 1)} = \left\{ g_k^{[t]} \right\} = \left\{ \sum_{i=1}^{I} (\sigma_{e_i}^{[t]})^{-1} E_{c_i}^{[t]} \left[ u_k^* Z_{ik} (y_i - X_i \beta) \right] \right\}
\]

An indirect approach

The original model with a constant \(\tau\) ratio specified in \([1-3]\) can be viewed as a special case of a more general model

\[
y_i = X_i \beta + \tau_i \sigma_{e_i} Z_i u^* + e_i
\]

with, as previously, \(\ell n \sigma_{e_i}^2 = \mathbf{p}_i' \delta\), but also with a linear structure on log-ratios

\[
\ell n \tau_i = \mathbf{h}_i' \lambda
\]

involving either the same \((\mathbf{h}_i = \mathbf{p}_i)\) or possibly different covariates.
Letting $\gamma = (\delta', \lambda')'$ here, the sequence of the CM-steps are
\[
\delta^{[t+1]} = \text{Argmax}_Q(\delta, \lambda^{[t]} | \gamma^{[t]}),
\]
and
\[
\lambda^{[t+1]} = \text{Argmax}_Q(\delta^{[t+1]}, \lambda | \gamma^{[t]}).
\]

The algorithm for $\delta$ is the same as in [7]. The algebra for $\lambda$ is shown in the Appendix, and leads to a system that can be written under a similar form as that of $\delta$,
\[
\mathbf{H}'\mathbf{W}_{\lambda\lambda}^{[t,\ell]} \mathbf{H} \cdot \Delta \lambda^{[t,\ell+1]} = \mathbf{H}'\mathbf{v}_{\lambda}^{[t,\ell]},
\]
where
\[
\Delta \lambda^{[t,\ell+1]} = \lambda^{[t,\ell+1]} - \lambda^{[t,\ell]},
\]
\[
\mathbf{H}'_{(s \times I)} = (h_1, h_2, \ldots, h_i, \ldots, h_I),
\]
\[
\mathbf{v}_{\lambda}^{[t,\ell]} (I \times 1) = \left\{ \tau_{i,\ell}^{[t,\ell]} \right\} = \left\{ \frac{\tau_{i,\ell}^{[t,\ell]}}{\sigma_{e_i}^{[t]}} \mathbf{E}_c^{[t]} \left( \mathbf{u}' \mathbf{Z}' \mathbf{e}_i^{[t,\ell]} \right) \right\}.
\]

Based on $\mathbf{e}_{i}^{[t,\ell]} = \mathbf{y}_i - \mathbf{X}_i \mathbf{\beta} - \tau_{i,\ell}^{[t,\ell]} \sigma_{e_i}^{[t]} \mathbf{Z}_i \mathbf{u}^*$, and $\tau_{i,\ell}^{[t,\ell]} = \exp(h'_i \lambda^{[t,\ell]})$,
\[
\mathbf{W}_{\lambda\lambda}^{[t,\ell]} (s \times s) = \text{diag} \left\{ \tau_{i,\ell}^{[t,\ell]} \mathbf{E}_c^{[t]} (\mathbf{u}' \mathbf{Z}' \mathbf{Z}_i \mathbf{u}^*) - \left( \sigma_{e_i}^{[t]} \right)^{-1} \mathbf{E}_c^{[t]} \left( \mathbf{u}' \mathbf{Z}' \mathbf{e}_i^{[t,\ell]} \right) \right\}
\]

For practical reasons, one may also wish to limit the number of inner iterations (index $\ell$) even to only one in order to reduce the volume of computation but the application of this ECM gradient algorithm should be performed carefully. Further empirical simplifications for the elements of [22] can be proposed along the same lines as in Foulley et al (1990).

Again, these results can be extended to a model with several random independent factors ($k = 1, 2, \ldots, K$) by setting
\[
\ell n \tau_{i,k} = h'_{ik} \lambda_k.
\]

Actually, if the CM-steps are performed for each vector $\lambda_k$ separately, the same formulae as in [20], [21] and [22] apply: just replace $\tau_i, \mathbf{Z}_i, \mathbf{u}^*$ by $\tau_{i,k}, \mathbf{Z}_{ik}, \mathbf{u}_k^*$ and set $\mathbf{e}_i$ equal to $\mathbf{y}_i - \mathbf{X}_i \mathbf{\beta} - \sum_{k=1}^{K} \tau_{i,k} \sigma_{e_i} \mathbf{Z}_{ik} \mathbf{u}_k^*$.

**ML estimation**

It may be interesting in some instances to use ML rather than REML for estimating variance components (see Discussion). The ECM procedure developed in this paper can be easily adapted to obtain ML parameter estimates. $\mathbf{\beta}$ is now part of the parameter vector instead of being a vector of random effects with infinite variance included in missing data. The $Q$ function to be maximized has the same formal expression as in [4] but here at the E-step, expectations have to be taken with
respect to the distribution of $u^*$ given $y, \gamma = \gamma^{[t]}$, and $\beta = \beta^{[t]}$. Maximization with respect to $\beta$ can be based on the equation $\partial Q/\partial \beta = 0$, ie

$$\sum_{i=1}^{I} \sigma_{e_i}^{-2} X_i^t \left[ y_i - X_i \beta - \tau \sigma_{e_i} Z_i \mathbb{E}(u^*|y, \beta, \gamma) \right] = 0 \quad [23]$$

One can proceed as previously, ie, run two CM-steps for the dispersion parameters based on the same E-step so as to obtain $\delta^{[t+1]}$ and $\tau^{[t+1]}$ (or $\lambda^{[t+1]}$), and then perform an additional CM-step for computing $\beta^{[t+1]}$ based on $[23]$, ie

$$\sum_{i=1}^{I} \sigma_{e_i}^{-2[t+1]} X_i^t X_i \beta^{[t+1]} = \sum_{i=1}^{I} \sigma_{e_i}^{-2[t+1]} X_i^t \left[ y_i - \tau^{[t+1]} \sigma_{e_i}^{[t+1]} Z_i \mathbb{E}(u^*|y, \beta^{[t]}, \delta^{[t]}, \tau^{[t]}) \right] \quad [24]$$

Alternatively, it may be advantageous to perform the CM-step for $\beta$ and the next E-step jointly by solving Henderson’s mixed model equations in $\beta^{[t+1]}$ and $u^{*[t+1]} = \mathbb{E}(u^*|y, \delta^{[t+1]}, \tau^{[t+1]})$ based on $\delta^{[t+1]}$ and $\tau^{[t+1]}$.

Formulae for the two CM-steps do not change. The only additional modification results from taking the conditional expectation of components of $e_i^t e_i$ given $y, \gamma = \gamma^{[t]}, \beta = \beta^{[t]}$ instead of $y, \gamma = \gamma^{[t]}$. Formulae in [12] reduce to

$$E_c^t \left[ (y_i - X_i \beta)'(y_i - X_i \beta) \right] = (y_i - X_i \widehat{\beta})'(y_i - X_i \widehat{\beta}) \quad [25a]$$

$$E_c^t \left[ u^* Z_i'(y_i - X_i \beta) \right] = \widehat{u}^* Z_i'(y_i - X_i \widehat{\beta}) \quad [25b]$$

$$E_c^t (u^* Z_i' Z_i u^*) = \widehat{u}^* Z_i' Z_i \widehat{u}^* + \text{tr} (Z_i' Z_i M_{uu}^{-1}) \quad [25c]$$

where $M_{uu}$ is the $u$ by $u$ block of the coefficient matrix [11].

Note that the trace terms inside those formulae have disappeared or have been greatly simplified owing to conditioning with respect to $\beta = \beta^{[t]}$. More generally, for models [13] involving several $u$-components, [25c] becomes

$$E_c^t (u_k^* Z_{ik} Z_{ik} u_k^*) = \widehat{u}_k^* Z_{ik} Z_{ik} \widehat{u}_k^* + \text{tr} (Z_{ik} Z_{ik} (M_{uu}^{-1})_{k\ell}) \quad [26]$$

where $(M_{uu}^{-1})_{k\ell}$ is the block pertaining to random factors $k$ and $\ell$ in the inverse of the random part of the coefficient matrix.

**Numerical example**

The procedures presented in this paper are illustrated with a small data set obtained from simulation. Data were generated according to a cross-classified model having two (environmental) fixed factors ($A = 2$ levels; $B = 3$ levels) and one (genetic) random factor ($S = 9$ levels). The genetic contribution consists of sire and maternal grand sire effects, the latter being assumed to have half the value of the first one. The model to generate the records was

$$y_{ijklm} = \mu + a_i + b_j + \tau \sigma_{eij} (s_k^* + 1/2s_k^*) + e_{ijklm} \quad [27]$$
where \( \mu \) is a general mean, \( a_i \) the effect of environmental factor \( A \) (\( i = 1, 2 \)), \( b_j \) the effect of environmental factor \( B \) (\( j = 1, 2, 3 \)), \( s_k^* \) the standardized contribution of male \( k \) as a sire, and \( 1/2s_{\ell}^* \) the standardized contribution of male \( \ell \) as a maternal grand sire, and \( e_{ijklm} \) the residual term.

Values chosen for the fixed effects were (using a full-rank parameterization):
\[
\mu + a_1 + b_1 = 100; a_2 - a_1 = 20; b_2 - b_1 = -10; b_3 - b_1 = -20. \]
The vector \( s^* = \{ s_k^* \} \) of sire effects is assumed to be \( N(0, A) \) with elements of the relationship matrix \( A \) shown at the bottom of table I.

**Table I.** Structure of the data set, number \(( n)\), sum of observations \(( \sum y)\) and sum of squares \(( \sum y^2)\) per cell.

| No | A | B | S | T | n | \( \sum y \) | \( \sum y^2 \) |
|----|---|---|---|---|---|-------------|-------------|
| 1  | 1 | 1 | 1 | 4 | 21| 2287        | 258771      |
| 2  | 1 | 2 | 1 | 4 | 19| 1739        | 162655      |
| 3  | 1 | 1 | 1 | 7 | 14| 1536        | 174610      |
| 4  | 1 | 3 | 1 | 7 | 7 | 628         | 57224       |
| 5  | 1 | 2 | 1 | 8 | 6 | 598         | 60398       |
| 6  | 1 | 3 | 2 | 6 | 12| 1045        | 95517       |
| 7  | 1 | 2 | 2 | 7 | 7 | 770         | 86554       |
| 8  | 1 | 2 | 2 | 8 | 18| 1726        | 169736      |
| 9  | 2 | 2 | 3 | 5 | 7 | 786         | 92030       |
| 10 | 2 | 2 | 3 | 5 | 10| 2523        | 350873      |
| 11 | 2 | 3 | 3 | 5 | 19| 1280        | 168166      |
| 12 | 2 | 3 | 3 | 2 | 37| 4305        | 536409      |
| 13 | 2 | 3 | 3 | 2 | 2 | 1460        | 168160      |
| 14 | 2 | 3 | 4 | 7 | 13| 1231        | 137599      |
| 15 | 2 | 4 | 4 | 8 | 6 | 243         | 14887       |
| 16 | 2 | 4 | 4 | 9 | 19| 1930        | 204578      |
| 17 | 2 | 4 | 5 | 12| 1141| 111575     |             |
| 18 | 2 | 4 | 5 | 12| 1141| 111575     |             |

\( A, B \): environmental factors treated as fixed; \( S \): sire and \( T \): maternal grand sire treated as random. Elements of the \( A \) matrix are the following \( \forall i, (i, i) = 1; \forall i \neq j, (i, j) = (j, i) = 0 \) except for \((1, 5) = (2, 5) = (3, 7) = (4, 6) = 1/2, and (1, 2) = (8, 9) = 1/4.\)

Residual variances were obtained from
\[
\sigma^2_{e_{ij}} = \exp(\mu^* + a_i^* + b_j^*)
\]
with a base line value \( \sigma^2_{e_{11}} = \exp(\mu^* + a_1^* + b_1^*) = 400 \), and multiplicative adjustment factors: \( \exp(a_2^* - a_1^*) = 2 \); \( \exp(b_2^* - b_1^*) = 1/2 \) and \( \exp(b_3^* - b_1^*) = 3/2 \). The ratio \( \tau = \sigma_{s_{ij}}/\sigma_{e_{ij}} \) of the square root of the sire to the residual variance was taken as constant over \( A \times B \) cells and set to \( 8.75^{-1/2} \) (heritability equal to 0.41).

There were 267 observations distributed among 18 different \( A \times B \times \) sire \times maternal grand sire subclasses. The data structure is displayed in table I as well as cell size \(( n)\), sum \(( \sum y)\) and sum of squares \(( \sum y^2)\) in each subclass.
Tests of hypotheses about the location parameters $\beta$, the residual dispersion parameters $\delta$ and the ratios $\tau_{ij}$ were carried out via the likelihood ratio statistic as described in previous studies (Foulley et al, 1990; San Cristobal et al, 1993; Meyer et al, 1993; Foulley and Quaas, 1995). Formulae by Quaas (1992) were used to compute maximized likelihood functions ($L_{\text{max}}$).

Results can be arranged as an analysis of variance (or deviance) table: see table II for hypothesis testing about $\beta$, and table III for residual ($\delta$) and ratio ($\lambda$) parameters. Note also that the test statistic for $\beta$ relies on $-2L_{\text{max}}$ evaluated from the ML estimates of all parameters, whereas a maximized residual likelihood can be better employed for $\delta$ and $\lambda$.

Interaction effects on location parameters are constantly rejected under different assumptions for the other parameters. The hypothesis of residual variance homogeneity is strongly rejected as well as single factor descriptions of heterogeneity. The assumption of a constant ratio $\tau$ turns out to be a reasonable one. The test results eventually agree with the simulation model; they support the practical conclusion that the $\mu + A + B$ model is the most appropriate to account for variation both in location and in log-residual variances, the ratio $\tau$ being constant.

The estimation procedure for $\delta$ and $\tau$ (or $\lambda$) is illustrated in table IV for this model and an alternative one using both standard and residual maximum likelihood methods of estimation. ML and REML estimates of residual variances do not differ very much; on the contrary, the ML estimates of the ratio $\tau$ turns out to be, as expected, lower than the REML ones, the values of the latter being close to the true value.

DISCUSSION AND CONCLUSION

The main purpose of this paper was to extend the general structural approach to heteroskedasticity in mixed models proposed by Foulley et al (1990, 1992) to the case of homogeneous ratios of $u$ to $e$ variance components.

In a sire by environment interaction, this is equivalent to postulating homogeneous intra-class correlations or heritabilities. This seems to be a reasonable assumption in practice, or at least serves as a suitable compromise between the existence of heteroskedasticity and parsimony of models. Less restrictive assumptions might also be investigated (Quaas, 1995, pers comm). This paper also provides a generalization of LR tests of this assumption to unbalanced data and complex model structures: see the previous work of Visscher (1992) on a one-way random balanced design, and that by Robert et al (1995a,b) for heterogeneous variances due to a single classification.

The EM algorithm turns out to be a convenient and powerful tool for solving variance component estimation problems. The ECM algorithm allows us to simplify the estimating equations, in particular the ECM gradient version. The advantage of this algorithm was especially clear here in the case of the indirect approach. A few examples of this for the mixed model have been already mentioned (Meng and Rubin, 1993 example 1; Walker, 1996). It offers great flexibility in defining the sequence of the conditional maximization steps, all the alternatives of which have not been investigated here. In the case addressed in this paper, the basic statistics
Table III. Likelihood statistics for testing location and dispersion parameters in heteroskedastic mixed models: residual components and ratios of variance

| No | Location | Model<sup>a</sup>  | Ratio  | Likelihood<sup>b</sup> | Covariate | Test<sup>c</sup> |
|----|----------|---------------------|--------|-----------------------|-----------|-----------------|
|    |          | Residual            |        |                       | par -2L   | Comp df Statistic | P-value |
| 1  | μ + A + B | μ* + A* + B* + AB*  | μ"    | 11 2420.9841          | AB*       | 2-1 2 3.5517 0.169 |
| 2  | μ + A + B | μ* + A* + B*        | μ"    | 9 2424.5359          | A*        | 3-2 1 19.5522 10^-5 |
| 3  | μ + A + B | μ* + B*             | μ"    | 8 2444.0881          | B*        | 4-2 2 21.6501 2.10^-5 |
| 4  | μ + A + B | μ* + A*             | μ"    | 7 2446.1860          | A"        | 5-1 5 54.5049 0   |
| 5  | μ + A + B | μ*                 | μ"    | 6 2475.4890          | AB* + B* + AB* | 5-1 5 54.5049 0   |

<sup>a</sup> Covariates: μ: intercept; A: 1st factor (eg, row); B: 2nd factor (eg, column); AB: interaction. Covariates coded as: location (eg, A), residual (A*), u to residual variance log-ratio (A")<sup>b</sup> par: number of parameters: L = maximum of the log-likelihood (REML version). <sup>c</sup> Likelihood ratio (LR) test based on difference in -2L between the full and reduced models; P-value based on the asymptotic chi-square distribution of the LR statistic under the null hypothesis.
Table II. Likelihood statistics for testing location and dispersion parameters in heteroskedastic mixed models: location parameters

| No | Location            | Model<sup>a</sup> | Residual       | u-or Ratio | Likelihood<sup>b</sup> | Covariate | Test<sup>c</sup> | Comp df | Statistic | P-value |
|----|---------------------|-------------------|----------------|-------------|-----------------------|-----------|------------------|---------|-----------|---------|
| 1  | $\mu + A + B + AB$  | $\mu^* + A^* + B^* + AB^*$ | $\mu^i$      | 13          | 2431.7779            | AB        | 2-1              | 2       | 3.7610    | 0.153   |
| 2  | $\mu + A + B$       | $\mu^* + A^* + B^* + AB^*$ | $\mu^i$      | 11          | 2435.5389            | AB        | 2-1              | 2       | 3.7896    | 0.150   |
| 3  | $\mu + B$           | $\mu^* + A^* + B^* + AB^*$ | $\mu^i$      | 10          | 2445.0835            | A         | 3-2              | 1       | 9.5446    | 0.0020  |
| 4  | $\mu + A$           | $\mu^* + A^* + B^* + AB^*$ | $\mu^i$      | 9           | 2467.1045            | B         | 4-2              | 2       | 31.5656   | 1.4.10^-7|
| 1  | $\mu + A + B + AB$  | $\mu^* + A^* + B^* + AB^*$ | $\mu^{''}$   | 13          | 2427.8476            | AB        | 2-1              | 2       | 3.7896    | 0.150   |
| 2  | $\mu + A + B$       | $\mu^* + A^* + B^* + AB^*$ | $\mu^{''}$   | 11          | 2431.6372            | AB        | 2-1              | 2       | 3.7896    | 0.150   |
| 3  | $\mu + B$           | $\mu^* + A^* + B^* + AB^*$ | $\mu^{''}$   | 10          | 2442.2131            | A         | 3-2              | 1       | 10.5759   | 0.0011  |
| 4  | $\mu + A$           | $\mu^* + A^* + B^* + AB^*$ | $\mu^{''}$   | 9           | 2457.5191            | B         | 4-2              | 2       | 25.8819   | 2.4.10^-6|

<sup>a</sup> Covariates: $\mu$: intercept; A: 1st factor (eg, row); B: 2nd factor (eg, column); AB: interaction. Covariates coded as: (eg, A), residual variance ($A^*$), $u$-component ($A^i$) and $u$ to residual variance log-ratio ($A^{''}$).<sup>b</sup> par: number of parameters; $L =$ maximum of the loglikelihood (ML version). <sup>c</sup> Likelihood ratio (LR) test based on difference in $-2L$ between the full and reduced models; $P$-value based on the asymptotic chi-square distribution of the LR statistic under the null hypothesis.
generated by the EM algorithm are strikingly natural (see Appendix B) thus giving a flavour of simplicity to the whole procedure.

It also makes it possible to switch from REML to ML or vice versa with very little change in implementing computations (Foulley et al, 1994). Some authors such as Leonard (1975), Denis (1983) and Anderson (1984) in statistics and Shaw (1987) in quantitative genetics have advocated the use of ML rather than REML procedures to estimate variance components. Although the interest of ML versus REML in that case remains questionable, ML estimates remain mandatory for hypothesis testing about $\beta$ via the LR statistic (see the numerical example). Bayesian point estimators can also be envisioned via EM (Foulley et al, 1992; Gianola et al, 1992; San Cristobal et al, 1993; Weigel and Gianola, 1993; Foulley and Quaas, 1995).

In addition, as already pointed out by Denis et al (1996), a LR test about $\beta$ requires $\delta$ and $\tau$ (or $\lambda$) being the same for the null hypothesis and its alternative; the same rule holds in hypothesis testing about $\delta$ (or $\lambda$) by keeping the other parameters the same over the models to be compared. This is part of the general

### Table IV. ML and REML estimates of dispersion parameters (a) and subclass variances (b) under two different models.

(a)

| No | Model   | Method of estimation | Parameter                          |
|----|---------|----------------------|------------------------------------|
|    |         |                      | Intercept $a_2-a_1$ | $b_2-b_1$ | $b_3-b_1$ |
| I  | Residual $\mu^* + A^* + B^*$ | REML                  | 5.955404 0.82921 | -0.67086 0.27739 |
|    |         | ML                   | 5.93191 0.85011 | -0.67577 0.28717 |
|    | Ratio$^a$ $\mu'' + A''$ | REML                  | -2.76809 2.05948 |
|    |         | ML                   | -3.43898 2.49856 |
| II | Residual $\mu^* + A^* + B^*$ | REML                  | 5.94316 0.85746 | -0.67391 0.30203 |
|    |         | ML                   | 5.92066 0.87583 | -0.67257 0.30706 |
|    | Ratio   $\mu''$ | REML                  | -1.11978 |
|    |         | ML                   | -1.31294 |

$^a$ $\tau_i = \sigma_{\mu_i}/\sigma_{e_i}$: estimates of parameters are expressed on a log-scale.

(b)

| No | Model   | Method | Subclass |
|----|---------|--------|----------|
|    |         |        | 11  | 12  | 13  | 21  | 22  | 23  |
| I  | Resid   $\mu^* + A^* + B^*$ | REML   | 385.305 196.995 508.480 882.932 441.516 1165.187 |
|    |         | ML     | 376.874 191.740 502.242 881.848 448.652 1175.196 |
|    | Ratio$^b$ $\mu'' + A''$ | REML   | 0.0628 0.0628 0.0628 0.4923 0.4923 0.4923 |
|    |         | ML     | 0.0321 0.0321 0.0321 0.3905 0.3905 0.3905 |
| II | Resid   $\mu^* + A^* + B^*$ | REML   | 381.135 194.269 515.521 898.398 457.923 1215.168 |
|    |         | ML     | 372.657 190.201 506.600 894.701 456.649 1216.282 |
|    | Ratio$^b$ $\mu''$ | REML   | 0.3264 0.3264 0.3264 0.3264 0.3264 0.3264 |
|    |         | ML     | 0.2690 0.2690 0.2690 0.2690 0.2690 0.2690 |

In both instances I and II, mode for location parameters is $\mu + A + B$. $^b$ True value = 0.338.
and difficult problem of joint modelling of means and variances, which is related to such issues as the Behrens–Fisher problem and multi-stage hypothesis testing, and which needs further consideration.

Another area that deserves caution and further development is that of estimability. Difficulties are expected when all the cells contributing to an element \( j \) of \( \delta \) or \( \lambda \) (or to a linear combination of them) have a weight tending towards zero. This may arise due to i) purely overparameterization problems, or due to ii) parameter values becoming extreme (eg, ratios \( \tau \), tending to zero implying elements of \( \lambda \) being infinite). This last phenomenon is similar to what happens in the analysis of binary and ordinal data with latent variable models (Misztal et al, 1989; Fahrmeir and Tutz, 1994). Such difficulties can be avoided by reparameterization (i), or by setting lower bounds to the diagonal elements of the system of equations to solve (ii).

Finally, asymptotic accuracy can also be worked out numerically within the EM framework via the so-called SEM algorithm (Meng and Rubin, 1991). Although it only requires the code of the complete data variance covariance matrix and of the EM or ECM outputs, the burden of calculations is then heavier, so that we suggest that it is restricted to the simplest models. Other computing alternatives should also be considered, eg, the average information-restricted maximum likelihood algorithm (AI-REML) of Gilmour et al (1995).

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APPENDIX A: Algebra for the estimating equations

The $Q$ function to be maximized is (in condensed notation)

$$Q(y) = \text{const} - \frac{1}{2} \sum_{i=1}^{I} n_i \ln \sigma^2_{e_i} - \frac{1}{2} \sum_{i=1}^{I} \sigma^{-2}_{e_i} E_c(e'_i e_i) \quad [A1]$$

with

$$\ln \sigma^2_{e_i} = p'_i \delta \quad [A2]$$

and

$$e_i = y_i - X_i \beta - \tau \sigma_{e_i} Z_i u^* \quad [A3]$$

Derivatives with respect to $\delta$ (residual dispersion parameters)

First derivatives: according to the chaining rule, one has

$$\frac{\partial (2Q)}{\partial \delta} = \sum_{i=1}^{I} \frac{\partial (2Q)}{\partial \ln \sigma^2_{e_i}} \frac{\partial \ln \sigma^2_{e_i}}{\partial \delta}$$

Now

$$\frac{\partial (2Q)}{\partial \ln \sigma^2_{e_i}} = \sigma^2_{e_i} \frac{\partial (2Q)}{\partial \sigma^2_{e_i}}$$

$$\frac{\partial \ln \sigma^2_{e_i}}{\partial \delta} = p_i$$
That is
\[
\frac{\partial (2Q)}{\partial \sigma_{e_i}^2} = -\frac{n_i}{\sigma_{e_i}^2} - \frac{\partial}{\partial \sigma_{e_i}^2} \left[ \sigma_{e_i}^{-2} E_c(e_i'e_i) \right]
\]
\[
\frac{\partial}{\partial \sigma_{e_i}^2} \left[ \sigma_{e_i}^{-2} E_c(e_i'e_i) \right] = \frac{E_c(e_i'e_i)}{\sigma_{e_i}^4} + \frac{1}{\sigma_{e_i}^2} \frac{\partial E_c(e_i'e_i)}{\partial \sigma_{e_i}^2}
\]
\[
\frac{\partial E_c(e_i'e_i)}{\partial \sigma_{e_i}^2} = \frac{1}{2\sigma_{e_i}} \frac{\partial E_c(e_i'e_i)}{\sigma_{e_i}} = \frac{1}{\sigma_{e_i}^2} E_c \left[ \left( \frac{\partial e_i'}{\partial \sigma_{e_i}} \right) e_i \right]
\]
and
\[
\frac{\partial e_i'}{\partial \sigma_{e_i}} = -\tau u'Z_i'
\]
Thus
\[
\frac{\partial E_c(e_i'e_i)}{\partial \sigma_{e_i}^2} = -\frac{\tau E_c(u'Z_i'e_i)}{\sigma_{e_i}}
\]
and
\[
\frac{\partial (2Q)}{\partial \sigma_{e_i}^2} = -\frac{n_i}{\sigma_{e_i}^2} + \frac{E_c(e_i'e_i)}{\sigma_{e_i}^4} - \frac{\tau E_c(u'Z_i'e_i)}{\sigma_{e_i}^3}
\]
Letting \( \nu_{\delta,i} = \frac{\partial (2Q)}{\partial \ln \sigma_{e_i}^2} \)
\[
\nu_{\delta,i} = \frac{E_c(e_i'e_i)}{\sigma_{e_i}^2} - n_i + \frac{\tau}{\sigma_{e_i}} E_c(u'Z_i'e_i)
\]
\[
= \frac{1}{\sigma_{e_i}^2} E_c \left[ (e_i + \tau \sigma_{e_i} Z_i u')' e_i \right] - n_i
\]
so that
\[
\frac{\partial (2Q)}{\partial \delta} = \sum_{i=1}^{I} \nu_{\delta,i} p_i = P'v_{\delta}
\]
with
\[
v_{\delta} = \left\{ \frac{E_c \left[ (y_i - X_i \beta)' e_i \right]}{\sigma_{e_i}^2} - n_i \right\}
\]
Second derivatives: from [A4a], one has
\[
\frac{\partial (2Q)}{\partial \delta \partial \delta'} = \sum_{i=1}^{I} p_i' \frac{\partial \nu_{\delta,i}}{\partial \ln \sigma_{e_i}^2} p_i
\]
Now, using [A4b]
\[
\frac{\partial \nu_{\delta,i}}{\partial \sigma_{e_i}^2} = -\frac{1}{\sigma_{e_i}^4} E_c \left[ (y_i - X_i \beta)' e_i \right] + \frac{1}{\sigma_{e_i}^2} \frac{\partial E_c \left[ (y_i - X_i \beta)' e_i \right]}{\partial \sigma_{e_i}^2}
\]
\[
\frac{\partial E_c}{\partial \sigma^2_{e_i}} \left[ (y_i - X_i\beta)' e_i \right] = \frac{1}{2\sigma_{e_i}} E_c \left[ (y_i - X_i\beta)' \frac{\partial e_i}{\partial \sigma_{e_i}} \right] = -\frac{\tau}{2\sigma_{e_i}} E_c \left[ (y_i - X_i\beta)' Z_i u^* \right]
\]

Thus, letting

\[
\frac{\partial (2Q)}{\partial \delta \delta'} = -\sum_{i=1}^{I} w_{\delta \delta,ii} p'_i p_i = -P'WP \tag{A5a}
\]

\[
w_{\delta \delta,ii} = \frac{1}{\sigma^2_{e_i}} E_c \left[ (y_i - X_i\beta)' e_i \right] \left[ \left( e_i + \frac{1}{2} \tau \sigma_{e_i} Z_i u^* \right) \right] \tag{A5b}
\]

or, alternatively

\[
w_{\delta \delta,ii} = \frac{1}{\sigma^2_{e_i}} E_c \left[ (y_i - X_i\beta)' (y_i - X_i\beta) \right] - \frac{\tau}{2\sigma_{e_i}} E_c \left[ (y_i - X_i\beta)' Z_i u^* \right] \tag{A5c}
\]

Finally, the non-linear system to solve reduces to

\[
P'W_{\delta \delta} P \cdot \Delta \delta = P'v_0 \tag{A6}
\]

**Derivatives with respect to \( \tau \) (ratio)**

\[
\frac{\partial Q}{\partial \tau} = -\frac{1}{2} \sum_{i=1}^{I} \frac{1}{\sigma^2_{e_i}} \frac{\partial [E_c (e'_i e_i) ]}{\partial \tau}
\]

Now

\[
\frac{\partial E_c (e'_i e_i)}{\partial \tau} = 2 E_c \left[ \left( \frac{\partial e'_i}{\partial \sigma_{e_i}} \right) e_i \right]
\]

and

\[
\frac{\partial e'_i}{\partial \tau} = -\sigma_{e_i} u^{*'} Z'_i
\]

Thus

\[
\frac{\partial Q}{\partial \tau} = \sum_{i=1}^{I} \frac{1}{\sigma_{e_i}} E_c (u^{*'} Z'_i e_i)
\]

\[
= \sum_{i=1}^{I} \frac{1}{\sigma_{e_i}} E_c \left[ u^{*'} Z'_i (y_i - X_i\beta) \right] - \tau \sum_{i=1}^{I} E_c (u^{*'} Z'_i Z_i u^*) \tag{A7}
\]

The equation \( \partial Q/\partial \tau = 0 \) results in

\[
\tau = \frac{\sum_{i=1}^{I} \sigma_{e_i}^{-1} E_c \left[ u^{*'} Z'_i (y_i - X_i\beta) \right]}{\sum_{i=1}^{I} E_c (u^{*'} Z'_i Z_i u^*)} \tag{A8}
\]
**Additional derivatives for the exact EM**

From the second expression in [A7] it follows immediately

\[ W_{\tau \tau} = - \frac{\partial^2 (2Q)}{\partial \tau^2} = 2 \sum_{i=1}^{l} E_c(\text{u}^* Z_i Z_i^t \text{u}^*) \quad [A9] \]

One has also to express

\[ \frac{\partial^2 (2Q)}{\partial \tau \partial \delta} = \sum_{i=1}^{l} \frac{\partial^2 Q}{\partial \tau \partial \sigma_{e_i}} \sigma_{e_i} p_i \]

Now, from [A7]

\[ \frac{\partial^2 Q}{\partial \tau \partial \sigma_{e_i}} = - \frac{1}{\sigma_{e_i}^2} E_c \left[ \text{u}^* Z_i^t (y_i - X_i \beta) \right] \]

so that

\[ - \frac{\partial^2 (2Q)}{\partial \tau \partial \delta} = \sum_{i=1}^{l} w_{\tau \delta, i} p_i = P' W_{\tau \delta} \quad [A10a] \]

where \( W_{\tau \delta} \) is a \((I \times 1)\) defined by

\[ W_{\tau \delta} = \{ w_{\tau \delta, i} \} = \left\{ \frac{1}{\sigma_{e_i}} E_c \left[ \text{u}^* Z_i^t (y_i - X_i \beta) \right] \right\} \quad [A10b] \]

The system for true EM (or gradient EM without inner iterations) is then

\[ \begin{bmatrix} P' W_{\delta \delta} P & P' W_{\tau \delta} \\ W_{\tau \delta} P & W_{\tau \tau} \end{bmatrix} \begin{bmatrix} \Delta \delta \\ \Delta \tau \end{bmatrix} = \begin{bmatrix} P' v_{\delta} \\ \nu_{\tau} \end{bmatrix} \quad [A11] \]

where \( \nu_{\tau} = 2 \frac{\partial Q}{\partial \tau} \), see [A7]

**Extension to \( K \) independent random factors**

\( Q \) in [A1] remains formally unchanged with \( e_i \) in [A3] being now

\[ e_i = y_i - X_i \beta - \sigma_{e_i} \sum_{k=1}^{K} \tau_k Z_{ik} u_k^* \quad [A12] \]

Based on this expression of the residual, formulae [A4ab] still hold.

Similarly, the expression of \( w_{\delta \delta, ii} \) in [A5b] becomes

\[ w_{\delta \delta, ii} = \frac{1}{\sigma_{e_i}^2} E_c \left[ (y_i - X_i \beta)' \left( e_i + \frac{\sigma_{e_i}}{2} \sum_{k=1}^{K} \tau_k Z_{ik} u_k^* \right) \right] \quad [A13a] \]
or, alternatively

\[
\frac{1}{\sigma_{e_i}^2} E_c [(y_i - X_i \beta)'(y_i - X_i \beta)] - \frac{1}{2\sigma_{e_i}} \sum_{k=1}^{K} \tau_k E_c [(y_i - X_i \beta)'Z_{ik}u^*_k]
\]

As far as \( \tau \) is concerned, [A6] becomes

\[
\frac{\partial Q}{\partial \tau_k} = \sum_{i=1}^{I} \frac{1}{\sigma_{e_i}} E_c (u^*_k Z'_{ik} e_i) \tag{A14}
\]

leading to the following system

\[
\begin{align*}
\tau_k \sum_{i=1}^{I} E_c (u^*_k Z'_{ik} Z_{ik} u^*_k) + \sum_{\ell \neq k} \tau_\ell \sum_{i=1}^{I} E_c (u^*_k Z'_{ik} Z_{i\ell} u^*_\ell) \\
= \sum_{i=1}^{I} \sigma_{e_i}^{-1} E_c (u^*_k Z'_{ik} Z_{i\ell} u^*_\ell)
\end{align*}
\]

\[
\text{for } k = 1, 2, \ldots, K \tag{A15}
\]

**Derivatives with respect to \( \lambda \) (parameters of the log-ratio)**

\( Q \) and the model for \( \ln \sigma_{e_i}^2 \) are the same as in [A1] and [A2] but the vector of residuals is defined as

\[
e_i = y_i - X_i \beta - \tau_i \sigma_{e_i} Z_i u^* \tag{A16}
\]

with

\[
\ln \tau_i = h'_i \lambda \tag{A17}
\]

Using condensed notation, the iterative system to solve can be written in the same way as previously, ie

\[
H'W_{\lambda \lambda}H \cdot \Delta \lambda = H'v_{\lambda} \tag{A18}
\]

where

\[
H'_{(s \times 1)} = (h_1, h_2, \ldots, h_i, \ldots, h_f)
\]

\[
v_{\lambda} = \{\nu_{\lambda,i} = \partial Q / \partial \ln \tau_i\} \tag{A19}
\]

\[
W_{\lambda \lambda} = \text{Diag} \left\{ w_{\lambda \lambda,ii} = -\frac{\partial \nu_{\lambda,i}}{\partial \ln \tau_i} \right\} \tag{A20}
\]

Now

\[
\nu_{\lambda,i} = \frac{\tau_i}{\sigma_{e_i}} E_c (u'^* Z'_{i} e_i) \tag{A21a}
\]
or, after some algebra
\[
\nu_{\lambda,i} = \frac{\tau_i}{\sigma_{e_i}} \text{E}_c \left[ u^* Z'_i(y_i - X_i \beta) \right] - \tau_i^2 \text{E}_c (u^* Z'_i Z_i u^*) \]

Furthermore,
\[
\frac{\partial \nu_{\lambda,i}}{\partial \tau_i} = \frac{1}{\sigma_{e_i}} \text{E}_c (u^* Z'_i e_i) + \frac{\tau_i}{\sigma_{e_i}} \text{E}_c \left( u^* Z'_i \frac{\partial e_i}{\partial \tau_i} \right)
\]
\[
\frac{\partial e_i}{\partial \tau_i} = -\sigma_{e_i} Z_i u^*
\]

Then
\[
w_{\lambda\lambda,ii} = -\tau_i \frac{\partial \nu_{\lambda,i}}{\partial \tau_i}
\]
\[
w_{\lambda\lambda,ii} = -\frac{\tau_i}{\sigma_{e_i}} \text{E}_c (u^* Z'_i e_i) + \tau_i^2 \text{E}_c (u^* Z'_i Z_i u^*)
\]

or
\[
w_{\lambda\lambda,ii} = 2\tau_i^2 \text{E}_c (u^* Z'_i Z_i u^*) - \frac{\tau_i}{\sigma_{e_i}} \text{E}_c \left[ u^* Z'_i (y_i - X_i \beta) \right]
\]

This can be easily generalized to several independent random factors
\[
\ln \tau_{ik} = h_{ik}' \lambda_k \quad \text{for} \quad k = 1, 2, \ldots, K
\]
if conditional maximization is performed factor by factor. The system [A18] applies to each factor \( k \) with
\[
\nu_{\lambda,i}^{(k)} = \frac{\tau_{ik}}{\sigma_{e_i}} \text{E}_c (u_{ik}' Z'_i k e_i)
\]
\[
w_{\lambda\lambda,ii}^{(k)} = -\frac{\tau_{ik}}{\sigma_{e_i}} \text{E}_c (u_{ik}' Z'_i k e_i) + \tau_{ik}^2 \text{E}_c (u_{ik}' Z'_i k Z_i k u_{ik})
\]

APPENDIX B: Formulae [12] for grouped data

In some instances (see, eg, the example in table I) data can be grouped so that the \( n_i \) observations within a stratum \( i \) share the same covariates
\[
\left\{ \begin{array}{l}
X_i = 1_{n_i} x'_i \\
Z_i = 1_{n_i} z'_i
\end{array} \right. \quad [B1]
\]

where \( x'_i \) and \( z'_i \) are the common row vectors \((1 \times p)\) and \((1 \times q)\) pertaining to fixed and random effects, respectively.
\[
\mu_i = x'_i \beta, \quad u'^*_i = z'_i u^*
\]

\[
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\]
and
\[
\begin{bmatrix}
\hat{S}_{\varepsilon\varepsilon,i} = E_c [(y_i - X_i\beta)'(y_i - X_i\beta)] \\
\hat{S}_{\varepsilon u,i} = E_c [u^*Z_i'y_i - X_i\beta] \\
\hat{S}_{uu,i} = E_c (u^*Z_i'Z_iu^*)
\end{bmatrix}
\tag{B3}
\]

one has
\[
\hat{S}_{\varepsilon\varepsilon,i} = y_i'y_i - 2\hat{\beta}'X_i'y_i + \hat{\beta}'X_i'X_i\hat{\beta} + \text{tr}(X_i'X_iC_{\beta\beta})
\]

Substituting $X_i$ by its expression in [B1] gives
\[
\hat{S}_{\varepsilon\varepsilon,i} = \sum_{j=1}^{n_i} (y_{ij} - \hat{\mu}_i)^2 + n_i \text{tr}(x_i'x_i' C_{\beta\beta})
\]
\[
= \sum_{j=1}^{n_i} y_{ij}^2 + n_i\hat{\mu}_i^2 - 2\hat{\mu}_iy_{i+} + n_i \text{tr}(x_i'x_i' C_{\beta\beta})
\tag{B4}
\]

where $y_{ij}$ is the $j$th element of $y_i$, and $y_{i+} = \sum_{j=1}^{n_i} y_{ij}$

Similarly
\[
u_i^* = z_i'z_i
\]
\[
\hat{S}_{\varepsilon u,i} = \hat{u}_i^*(y_{i+} - n_i\hat{\mu}_i) - n_i \text{tr}(z_i'X_i' C_{\beta u})
\tag{B5}
\]

and
\[
\hat{S}_{uu,i} = n_i \{\hat{u}_i^2 + \text{tr}(z_i'z_i' C_{uu})\}
\tag{B6}
\]