

Supplement to

“Spectral-based noncentral F mixed effect models, with application to otoacoustic emissions”

Lai Wei¹, Peter F. Craigmile^{2,*}, Wayne M. King³

¹ Biostatistics Center, The Ohio State University, Columbus. OH 43210.

² Dept. of Statistics, The Ohio State University, Columbus. OH 43210.

³ Signal Processing and Communications, The Mathworks, Inc., 3 Apple Hill, Natick, MA 01760

*email: pfc@stat.osu.edu

The MCMC algorithm used to fit the Bayesian DPOAE noncentral F mixed model

We use $\boldsymbol{\theta} \setminus \{\kappa\}$ to denote the collection of parameters contained in $\boldsymbol{\theta}$, but excluding the parameter κ . The MCMC algorithm below assumes that $K_j = K$ for all subjects j . (More carefully subsetting and indexing is needed in the unbalanced case.)

Update $\{\log r_{j,k}\}$: Fix a subject $j = 1, \dots, J$, and let $\mathbf{log} \mathbf{r}_{j,\bullet} = (\log r_{j,1}, \dots, \log r_{j,K})^T$.

Then

$$\pi(\mathbf{log} \mathbf{r}_{j,\bullet} | \mathbf{z}, \boldsymbol{\theta} \setminus \{\mathbf{log} \mathbf{r}_{j,\bullet}\}) \propto \prod_{k=1}^K \left[\prod_{l=1}^L f_{Z_{j,k,l}}(z_{j,k,l} | e^{N\Delta r_{j,k}}) \right] n(\log r_{j,k} | \alpha_k + G_j \beta_k, \tau_k^2).$$

We use a Metropolis-Hastings symmetric random walk update. Supposing we are at $\mathbf{log} \mathbf{r}_{j,\bullet}$, we propose $\mathbf{log} \mathbf{r}_{j,\bullet}^{new}$ from a K -variate normal with mean $\mathbf{log} \mathbf{r}_{j,\bullet}$ and covariance Σ_j , for some $K \times K$ positive definite matrix Σ_j (in practice, we base Σ_j on the estimated covariance matrix of the maximum likelihood estimate of $\mathbf{log} \mathbf{r}_{j,\bullet}$, calculated using only the data for

subject j , scaled to obtain an acceptance probability of around 0.4). Then we accept the new value, $\mathbf{log} \mathbf{r}_{j,\bullet}^{new}$, with probability $\min(e^q, 1)$, where

$$q = \left\{ \sum_{k=1}^K \sum_{l=1}^L \log f_{Z_{j,k,l}}(z_{j,k,l} | e^{N\Delta r_{j,k}^{new}}) + \sum_{k=1}^K \log n(\log r_{j,k}^{new} | \alpha_k + G_j \beta_k, \tau_k^2) \right\} - \left\{ \sum_{k=1}^K \sum_{l=1}^L \log f_{Z_{j,k,l}}(z_{j,k,l} | e^{N\Delta r_{j,k}}) + \sum_{k=1}^K \log n(\log r_{j,k} | \alpha_k + G_j \beta_k, \tau_k^2) \right\};$$

otherwise we remain at $\mathbf{log} \mathbf{r}_{j,\bullet}$.

Update $\{\alpha_k\}$ and $\{\beta_k\}$: Fixing a $k = 1, \dots, K$ we sample α_k and β_k jointly, conditional on the data and other parameters. First note that

$$\pi(\alpha_k, \beta_k | \mathbf{z}, \boldsymbol{\theta} \setminus \{\alpha_k, \beta_k\}) \propto \left\{ \prod_{j=1}^J n(\log r_{j,k} | \alpha_k + G_j \beta_k, \tau_k^2) \right\} n(\alpha_k | \mu_\alpha, \sigma_\alpha^2) n(\beta_k | \mu_\beta, \sigma_\beta^2).$$

Let X be a $J \times 2$ design matrix with first column all ones, and second column G_j ($j = 1, \dots, J$). Then $\boldsymbol{\beta}_{\bullet,k} = (\beta_{1,k}, \dots, \log r_{j,k})^T$, conditional on α_k, β_k and τ_k^2 , is J -variate normal with mean $X(\alpha_k, \beta_k)^T$ and covariance $\tau_k^2 I_J$, where I_J is the $J \times J$ identity matrix. This is a Bayesian regression model. Hence, letting

$$V = \begin{bmatrix} \sigma_\alpha^2 & 0 \\ 0 & \sigma_\beta^2 \end{bmatrix},$$

our sample from $(\alpha_k, \beta_k)^T$, conditional on the data and other parameters, is a bivariate normal draw with a mean $\Sigma^{-1} \mathbf{c}$ and covariance Σ^{-1} where $\Sigma = X^T X / \tau_k^2 + V^{-1}$ and $\mathbf{c} = X^T \boldsymbol{\beta}_{\bullet,k} / \tau_k^2 + V^{-1}(\mu_\alpha, \mu_\beta)^T$.

Update $\{\tau_k^2\}$: For each $k = 1, \dots, K$ we have that

$$\pi(\tau_k^2 | \mathbf{z}, \boldsymbol{\theta} \setminus \{\tau_k^2\}) \propto \left\{ \prod_{j=1}^J n(\log r_{j,k} | \alpha_k + G_j \beta_k, \tau_k^2) \right\} ig(\tau_k^2 | s_\tau, r_\tau),$$

leading us to sample τ_k^2 , conditional on the data and other parameters, from an inverse gamma distribution with shape s and rate r , where

$$s = s_{\sigma, \alpha} + J/2, \quad \text{and} \quad r = r_{\sigma, \alpha} + \frac{1}{2} \sum_{j=1}^J (\log r_{j,k} - \alpha_k - G_j \beta_k)^2.$$

Update μ_α : (The update for μ_β is similar.)

$$\pi(\mu_\alpha | \mathbf{z}, \boldsymbol{\theta} \setminus \{\mu_\alpha\}) \propto \left\{ \prod_{k=1}^K n(\alpha_k | \mu_\alpha, \sigma_\alpha^2) \right\} n(\mu_\alpha | m_\alpha, v_\alpha),$$

and hence we sample μ_α , conditional on the data and other parameters, from a normal distribution with mean m/p and variance $1/p$ where

$$m = \sum_{k=1}^K \alpha_k / \sigma_\alpha^2 + m_\alpha / v_\alpha, \quad \text{and} \quad p = K / \sigma_\alpha^2 + 1 / v_\alpha.$$

Update σ_α^2 : (The update for σ_β^2 is similar.)

$$\pi(\sigma_\alpha^2 | \mathbf{z}, \boldsymbol{\theta} \setminus \{\sigma_\alpha^2\}) \propto \left\{ \prod_{k=1}^K n(\alpha_k | \mu_\alpha, \sigma_\alpha^2) \right\} ig(\sigma_\alpha^2 | s_{\sigma, \alpha}, r_{\sigma, \alpha}),$$

and hence we sample σ_α^2 , conditional on the data and other parameters, from an inverse gamma distribution with shape s and rate r , where

$$s = s_\tau + K/2, \quad \text{and} \quad r = r_\tau + \frac{1}{2} \sum_{k=1}^K (\alpha_k - \mu_\alpha)^2.$$