

Model specification

We will repress the location notation just to simplify the further expressions. In other words, let $\mathbf{x}_k(\mathbf{s}) = \mathbf{x}_k$, $k \in \{m, g, p\}$ where m relates to the data from the SDM, g from genetic data and p from pollen data.

Level 1 - Data collection specific models

$$\begin{aligned} x_m | x^{(m)} &= \beta_0 + \beta_1 x^{(m)} + \epsilon_m & \epsilon_m &\sim N(0, \sigma_m^2) \\ x_g | x^{(g)} &= \alpha_0 + \alpha_1 x^{(g)} + \epsilon_g & \epsilon_g &\sim N(0, \sigma_g^2) \\ x_p | x^{(p)} &= \gamma_0 + \gamma_1 x^{(p)} + \epsilon_p & \epsilon_p &\sim N(0, \sigma_p^2) \end{aligned}$$

where $x^{(k)}$ indicates the values of \mathbf{x} , the latent spatial process, that relate to model k , $k \in \{m, g, p\}$. Each of these models has a sample size n_k , $k \in \{m, g, p\}$. The sample sizes might be different and they might not add up to n , the length of \mathbf{x} , in the case where one location appears in more than one data collection model.

Since the $\mathbf{x}^{(k)}$ are essentially subsets of \mathbf{x} we can write them in terms of \mathbf{x} and a matrix that chooses the values of \mathbf{x} that appear in $\mathbf{x}^{(k)}$. Let these be the following:

$$\mathbf{x}^{(m)} = \mathbf{M}\mathbf{x} \quad \mathbf{x}^{(g)} = \mathbf{G}\mathbf{x} \quad \mathbf{x}^{(p)} = \mathbf{P}\mathbf{x}$$

Note that $\mathbf{K}\mathbf{K}^T = \mathbf{I}_{n_k}$ but $\mathbf{K}^T\mathbf{K} \neq \mathbf{I}_n$, for $\mathbf{K} \in \{\mathbf{M}, \mathbf{G}, \mathbf{P}\}$ and its corresponding $k \in \{m, g, p\}$.

Level 2 - Latent spatial process

$$\mathbf{x} \sim GP(\mu, \Sigma)$$

where μ is a common mean across all locations and $\Sigma_{i,j} = \sigma^2 e^{-\frac{h_{i,j}}{\phi}}$ is the exponential covariance function where $h_{i,j} = \|\mathbf{s}_i - \mathbf{s}_j\|$ is the distance between the i -th and j -th sites. Note that we can also write the covariance matrix as $\Sigma = \sigma^2 \mathbf{V}$, where \mathbf{V} is a correlation matrix.

Level 3 - Priors

$$\beta_0, \alpha_0, \gamma_0 \sim N(0, \tau)$$

$$\beta_1, \alpha_1, \gamma_1 \sim N(1, \tau)$$

$$\sigma_m^2, \sigma_g^2, \sigma_p^2, \sigma^2 \sim IG(a, b)$$

$$\mu \sim N(0, \omega)$$

$$\phi \sim Unif(A, B)$$

Posterior distributions

Conjugate

The following are the known conditional posterior distributions so that we might use a Gibbs sampler. For notation, *e.e.* will stand for “everything else”. Also, $\bar{x}_k = \frac{1}{n_k} \mathbf{1}_{n_k}^T \mathbf{x}_k$ stands for the average of \mathbf{x}_k , whereas $\bar{x}^{(k)} = \frac{1}{n_k} \mathbf{1}_{n_k}^T \mathbf{x}^{(k)}$ stand for the average of the latent process \mathbf{x} associated to model k , $k \in \{m, g, p\}$.

$$\begin{aligned}\beta_0|e.e. &\sim N\left(\frac{n_m}{\sigma_m^2 \rho_m}(\bar{x}_m - \beta_1 \bar{x}^{(m)}), \frac{1}{\rho_m}\right) \\ \alpha_0|e.e. &\sim N\left(\frac{n_g}{\sigma_g^2 \rho_g}(\bar{x}_g - \alpha_1 \bar{x}^{(g)}), \frac{1}{\rho_g}\right) \\ \gamma_0|e.e. &\sim N\left(\frac{n_p}{\sigma_p^2 \rho_p}(\bar{x}_p - \gamma_1 \bar{x}^{(p)}), \frac{1}{\rho_p}\right)\end{aligned}$$

where $\rho_k = \frac{n_k}{\sigma_k^2} + \frac{1}{\tau}$, $k \in \{m, g, p\}$.

$$\begin{aligned}\beta_1|e.e. &\sim N\left(\frac{\frac{n_m}{\sigma_m^2}(c_m - \beta_0 \bar{x}^{(m)}) + \frac{1}{\tau}}{\eta_m}, \frac{1}{\eta_m}\right) \\ \alpha_1|e.e. &\sim N\left(\frac{\frac{n_g}{\sigma_g^2}(c_g - \alpha_0 \bar{x}^{(g)}) + \frac{1}{\tau}}{\eta_g}, \frac{1}{\eta_g}\right) \\ \gamma_1|e.e. &\sim N\left(\frac{\frac{n_p}{\sigma_p^2}(c_p - \gamma_0 \bar{x}^{(p)}) + \frac{1}{\tau}}{\eta_p}, \frac{1}{\eta_p}\right)\end{aligned}$$

where $c_k = \frac{1}{n_k} \mathbf{x}_k^T \mathbf{x}^{(k)}$ is the average of the cross-products and $\eta_k = \frac{n_k s_k}{\sigma_k^2} + \frac{1}{\tau}$, where

$s_k = \frac{1}{n_k} \sum_{i=1}^{n_k} [x_i^{(k)}]^2 = \frac{1}{n_k} \mathbf{x}^T \mathbf{K}^T \mathbf{K} \mathbf{x}$ is the mean squares of the latent process associated to model k , $\mathbf{K} \in \{\mathbf{M}, \mathbf{G}, \mathbf{P}\}$ and its corresponding $k \in \{m, g, p\}$.

$$\begin{aligned}\sigma_m^2|e.e. &\sim IG\left(\frac{n_m}{2} + a, b + \frac{1}{2} \sum_{i=1}^{n_m} (x_{m,i} - \beta_0 - \beta_1 x_i^{(m)})^2\right) \\ \sigma_g^2|e.e. &\sim IG\left(\frac{n_g}{2} + a, b + \frac{1}{2} \sum_{i=1}^{n_g} (x_{g,i} - \alpha_0 - \alpha_1 x_i^{(g)})^2\right) \\ \sigma_p^2|e.e. &\sim IG\left(\frac{n_p}{2} + a, b + \frac{1}{2} \sum_{i=1}^{n_p} (x_{p,i} - \gamma_0 - \gamma_1 x_i^{(p)})^2\right)\end{aligned}$$

Finally, for the latent process:

$$\begin{aligned}\mu|e.e. &\sim N\left(\frac{\mathbf{1}_n^T \Sigma^{-1} \mathbf{x}}{\mathbf{1}_n^T \Sigma^{-1} \mathbf{1}_n + \frac{1}{\omega}}, \frac{1}{\mathbf{1}_n^T \Sigma^{-1} \mathbf{1}_n + \frac{1}{\omega}}\right) \\ \sigma^2|e.e. &\sim IG\left(\frac{n}{2} + a, \frac{1}{2} (\mathbf{x} - \mu \mathbf{1}_n)^T \mathbf{V}^{-1} (\mathbf{x} - \mu \mathbf{1}_n) + b\right)\end{aligned}$$

$$\mathbf{x}|e.e. \sim N_n\left(\tilde{\Sigma}\tilde{\boldsymbol{\mu}}, \tilde{\Sigma}\right)$$

where,

$$\begin{aligned}\tilde{\boldsymbol{\mu}} &= \mu\boldsymbol{\Sigma}^{-1}\mathbf{1}_n + \frac{\beta_1}{\sigma_m^2}\mathbf{M}^T(\mathbf{x}_m - \beta_0\mathbf{1}_{n_m}) + \frac{\alpha_1}{\sigma_g^2}\mathbf{G}^T(\mathbf{x}_g - \alpha_0\mathbf{1}_{n_g}) + \frac{\gamma_1}{\sigma_p^2}\mathbf{P}^T(\mathbf{x}_p - \gamma_0\mathbf{1}_{n_p}) \\ \tilde{\Sigma} &= \left[\boldsymbol{\Sigma}^{-1} + \frac{\beta_1^2}{\sigma_m^2}\mathbf{M}^T\mathbf{M} + \frac{\alpha_1^2}{\sigma_g^2}\mathbf{G}^T\mathbf{G} + \frac{\gamma_1^2}{\sigma_p^2}\mathbf{P}^T\mathbf{P}\right]^{-1}\end{aligned}$$

Non-Conjugate

ϕ has no know posterior distribution, however we can state what the density is proportional to so that we can use Metropolis-Hastings. Note that \mathbf{V} is a function of ϕ so for this last part we will use the notation $\mathbf{V}(\phi)$ to indicate this.

$$\pi(\phi|e.e.) \propto |\mathbf{V}(\phi)|^{-1/2} \cdot \exp\left\{\frac{1}{2\sigma^2}(\mathbf{x} - \mu\mathbf{1}_n)^T\mathbf{V}(\phi)^{-1}(\mathbf{x} - \mu\mathbf{1}_n)\right\} \cdot 1_{\{A < \phi < B\}}$$

where $|\mathbf{V}(\phi)|$ is the determinant of the matrix.