# Bayesian Methods for Trait Evolution

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# Bayesian Phylogeography

#### 1 Bayesian for Continuous Traits

Pruning Algorithms

Operation Phylogeography

General Framework BM Gibbs Constrained Spaces

# Outline

#### 1 Bayesian for Continuous Traits

- General Framework
- BM Gibbs
- Constrained Spaces
- Pruning Algorithms
- B Phylogeography

General Framework BM Gibbs Constrained Spaces

#### Maximum Likelihood

Likelihood: Distribution of the data given the parameters.

 $p(\mathbf{Y}, \mathbf{S} \mid \boldsymbol{\theta}, \mathcal{T}, \boldsymbol{\psi})$ 

General Framework BM Gibbs Constrained Spaces

## Maximum Likelihood

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Maximum likelihood: find the "best" possible parameters.

$$(\hat{\boldsymbol{ heta}}, \hat{\mathcal{T}}, \hat{\boldsymbol{\psi}}) = \operatorname*{argmax}_{\substack{\boldsymbol{ heta}, \mathrm{tr}, \boldsymbol{\psi}}} p\left(\mathbf{Y}, \mathbf{S} \mid \boldsymbol{ heta}, \mathcal{T}, \boldsymbol{\psi}\right)$$

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#### Limitations:

No a priori knowledge.
 All parameters are equally probable.

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- No a priori knowledge.
   All parameters are equally probable.
- Point estimate. No notion of uncertainty.

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ight)$$

Limitations:

- No a priori knowledge.
   All parameters are equally probable.
- Point estimate.

No notion of uncertainty.

 $\rightarrow$  Use Bayesian inference.

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#### **Bayesian Inference**

Likelihood: Distribution of the data given the parameters.

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General Framework BM Gibbs Constrained Spaces

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Prior: Distribution of the parameters.

 $p(\theta, \mathcal{T}, \psi)$ 

General Framework BM Gibbs Constrained Spaces

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Posterior: Distribution of the parameters given the data.

 $p(\theta, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S})$ 

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 $p(\theta, T, \psi \mid \mathbf{Y}, \mathbf{S})$ 

Bayesian Inference: Can we learn about the posterior ?

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## **Bayes** Theorem

Bayes Theorem:

$$p(\mathbf{A} \mid \mathbf{B}) = \frac{p(\mathbf{B} \mid \mathbf{A}) p(\mathbf{A})}{p(\mathbf{B})}$$

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•  $p(\mathbf{Y}, \mathbf{S})$  does not depend on the parameters (constant).

General Framework BM Gibbs Constrained Spaces

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We cannot compute the posterior, but we can sample from it.

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# Bayes Inference by Sampling

Assume we can sample from the posterior:

$$(oldsymbol{ heta}_i,\mathcal{T}_i,oldsymbol{\psi}_i)\sim p\left(oldsymbol{ heta},\mathcal{T},oldsymbol{\psi}\mid\mathbf{Y},\mathbf{S}
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 iid

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We can get some moments estimates. E.g. for continuous parameters:

$$\mathbb{E}\left[\boldsymbol{\theta} \mid \mathbf{Y}, \mathbf{S}\right] \approx \frac{1}{n} \sum_{i=1}^{n} \boldsymbol{\theta}_{i}$$
$$\mathbb{V} \text{ar}\left[\boldsymbol{\theta} \mid \mathbf{Y}, \mathbf{S}\right] \approx \frac{1}{n-1} \sum_{i=1}^{n} \left(\boldsymbol{\theta}_{i} - \frac{1}{n} \sum_{i=1}^{n} \boldsymbol{\theta}_{i}\right)^{2}$$

General Framework BM Gibbs Constrained Spaces

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We can estimate the parameters distribution (histogram).

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# Bayes Inference by Sampling

```
set.seed(18300730)
# n=1000 sample from a mystery distribution
sample <- rmystery(10000)
# Histogram
hist(sample, probability = TRUE)</pre>
```



General Framework BM Gibbs Constrained Spaces

# Bayes Inference by Sampling

```
set.seed(18300730)
# n=1000 sample from a mystery distribution
sample <- rmystery(10000)
# Histogram
hist(sample, probability = TRUE)
# distribution
x <- seq(-2, 6, 0.1)
lines(x, dnorm(x, 2, 1), lwd = 2, col = "firebrick3")</pre>
```





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#### **Bayesian Inference**

Goal: Sample from the posterior

$$p(\theta, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S}) = \frac{p(\mathbf{Y}, \mathbf{S} \mid \theta, \mathcal{T}, \psi) p(\theta, \mathcal{T}, \psi)}{p(\mathbf{Y}, \mathbf{S})}$$

That we know only up to a constant.

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#### **Bayesian Phylogenetics**

#### Goal:

#### $p(\theta, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S})$

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#### **Bayesian Phylogenetics**

#### Goal:

#### $p(\theta, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S}) \propto p(\mathbf{Y}, \mathbf{S} \mid \theta, \mathcal{T}, \psi) p(\theta, \mathcal{T}, \psi)$

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## **Bayesian Phylogenetics**

#### Goal:

# $$\begin{split} p(\theta, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S}) &\propto p(\mathbf{Y}, \mathbf{S} \mid \theta, \mathcal{T}, \psi) \, p(\theta, \mathcal{T}, \psi) \\ &\propto p(\mathbf{Y} \mid \theta, \mathcal{T}) \, p(\mathbf{S} \mid \mathcal{T}, \psi) \, p(\theta, \mathcal{T}, \psi) \end{split}$$

#### Assumption: **Y** and **S** independent conditionally on $\mathcal{T}$ .

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### **Bayesian Phylogenetics**

#### Goal:

# $p(\theta, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S}) \propto p(\mathbf{Y}, \mathbf{S} \mid \theta, \mathcal{T}, \psi) p(\theta, \mathcal{T}, \psi)$ $\propto p(\mathbf{Y} \mid \theta, \mathcal{T}) p(\mathbf{S} \mid \mathcal{T}, \psi) p(\theta, \mathcal{T}, \psi)$ $\propto p(\mathbf{Y} \mid \theta, \mathcal{T}) p(\theta) p(\mathbf{S} \mid \mathcal{T}, \psi) p(\mathcal{T}, \psi)$

Assumption: **Y** and **S** independent conditionally on  $\mathcal{T}$ .

For now:  $\mathcal{T}$  fixed.

General Framework BM Gibbs Constrained Spaces

## Markov Chain Monte Carlo

(Larget, 2005)

## Goal: Sample from $p(\theta \mid \mathbf{Y}) \propto p(\mathbf{Y} \mid \theta) p(\theta)$

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Problem: We do not know the normalizing constant.

Metropolis - Hasting: Iterate:

- Draw  $\theta^*$  in  $q(\theta \mid \theta^t)$ .
- Set  $\theta^{(t+1)} = \theta^*$  with probability:

$$r_{t} = \min\left\{1, \frac{p\left(\mathbf{Y} \mid \boldsymbol{\theta}^{*}\right)}{p\left(\mathbf{Y} \mid \boldsymbol{\theta}^{t}\right)} \frac{p\left(\boldsymbol{\theta}^{*}\right)}{p\left(\boldsymbol{\theta}^{t}\right)} \frac{q(\boldsymbol{\theta}^{(t)} \mid \boldsymbol{\theta}^{*})}{q(\boldsymbol{\theta}^{*} \mid \boldsymbol{\theta}^{(t)})}\right\}$$

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- The  $\theta^{(t)}$  is a Markov Chain.
- Its stationary distribution is  $p(\theta | \mathbf{Y})$ .  $\rightarrow$  For "t large enough",  $\theta^{(t)} \sim p(\theta | \mathbf{Y})$ .

(Larget, 2005)

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Animation: Chi Feng's website

(Larget, 2005)

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Gibbs: Rejection free.

• Split 
$$\theta = (\theta_{[1]}, \dots, \theta_{[K]})$$
.  
• Draw  $\theta^*$  in  $p(\theta_{[k]} \mid \theta_{[-k]}^{(t)}, \mathbf{Y})$  so that  $r_t = 1$ .

General Framework BM Gibbs Constrained Spaces

# BM: Gibbs with Conjugate Priors

Likelihood:

 $\mathbf{Y}|\mathbf{R}, \boldsymbol{\mu} \sim \mathcal{MN}(\mathbf{1}_n \boldsymbol{\mu}^T, \mathbf{V}, \mathbf{R})$ 

General Framework BM Gibbs Constrained Spaces

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Conjugate Priors:

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u) \ egin{aligned} oldsymbol{\mu} & | \mathbf{\mathsf{R}} &\sim \mathcal{N}(oldsymbol{\mu}_0, \kappa_0^{-1}\mathbf{\mathsf{R}}) \end{aligned}$ 

General Framework BM Gibbs Constrained Spaces

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Gibbs:

$$\mathbf{R}|\mathbf{Y}, \boldsymbol{\mu} \sim \mathcal{IW}(\mathbf{R}_n, \nu_n)$$
 with  $\mathbf{R}_n = f(\mathbf{Y}, \boldsymbol{\mu}, \mathbf{V})$
General Framework BM Gibbs Constrained Spaces

### BM: Gibbs with Conjugate Priors

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Gibbs:

$$\mathbf{R}|\mathbf{Y}, \boldsymbol{\mu} \sim \mathcal{IW}(\mathbf{R}_n, \nu_n) \quad \text{with} \quad \mathbf{R}_n = f(\mathbf{Y}, \boldsymbol{\mu}, \mathbf{V})$$

 $\hookrightarrow$  Automatic sampling in the space of variance matrices.

General Framework BM Gibbs Constrained Spaces

### Wishart Distribution

Wishart: generalization of the gamma distribution to matrices.

$$\mathbf{P} \sim \mathcal{W}_{p}(\mathbf{P}_{0}, 
u), \quad 
u \geq p$$

General Framework BM Gibbs Constrained Spaces

### Wishart Distribution

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$$\mathbf{P} \sim \mathcal{W}_{p}(\mathbf{P}_{0}, \nu), \quad \nu \geq p$$

Moments:

$$\mathbb{E}[\mathbf{P}] = \nu \mathbf{P}_0 \qquad \mathbb{V} \text{ar}[\mathbf{P}_{ij}] = \nu [(P_0)_{ij}^2 + (P_0)_{ii}(P_0)_{jj}]$$

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Sample Variance: If  $\mathbf{Y}_i \sim \mathcal{N}(\boldsymbol{\mu}, \boldsymbol{\Sigma})$  are *n* iid vectors of size *p*, then:

$$\hat{\boldsymbol{\Sigma}} = \sum_{i=1}^{n} (\mathbf{Y}_i - \boldsymbol{\mu}) (\mathbf{Y}_i - \boldsymbol{\mu})^T \sim \mathcal{W}_p(\boldsymbol{\Sigma}, n)$$

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Inverse Wishart:

$$\mathbf{R} \sim \mathcal{IW}(\mathbf{R}_0, 
u) \iff \mathbf{R}^{-1} \sim \mathcal{W}_p(\mathbf{R}_0^{-1}, 
u), \quad 
u > p-1$$

Bayesian for Continuous Traits

General Framework BM Gibbs Constrained Spaces

### Wishart Distribution



Diagonal and off-diagonal elements are related.

LKJ

Likelihood Ancestral State Reconstruction Gradient

## Outline

#### Bayesian for Continuous Traits

#### Pruning Algorithms

- Likelihood
- Ancestral State Reconstruction
- Gradient

#### B Phylogeography

### General Gaussian Model: Likelihood



Likelihood:  $\log p(\mathbf{Y})$  in one post-order traversal.  $\hookrightarrow O(N)$ . Bayesian for Continuous Traits Likelihood Pruning Algorithms Ancestral State Reconstruction Phylogeography Gradient

### General Gaussian Model: Likelihood



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 $\hookrightarrow O(N).$ 

 $\hookrightarrow$  "Pruning", "Gaussian elimination", "Phylogenetic Kalman filter", ...

Bayesian for Continuous Traits Likelihood Pruning Algorithms Ancestral State Reconstruction Phylogeography Gradient

### General Gaussian Model: Likelihood



Likelihood:  $\log p(\mathbf{Y})$  in one post-order traversal.

 $\hookrightarrow O(N).$ 

 $\hookrightarrow$  "Pruning", "Gaussian elimination", "Phylogenetic Kalman filter", ...

Difficulty: Numerical robustness.

Likelihood Ancestral State Reconstruction Gradient

### Likelihood Computation

Likelihood function:

$$f_{\mathbf{Y}^{j}|X_{j}}(\mathbf{Y}^{j};a) = A_{j}(\mathbf{Y}^{j})\Phi_{M_{j}(\mathbf{Y}^{j}),S_{j}^{2}(\mathbf{Y}^{j})}(a)$$



Likelihood Ancestral State Reconstruction Gradient

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Initialization: for tips

$$f_{Y_i|Y_i}(Y_i;a) = \Phi_{Y_i,0}(a)$$



Likelihood Ancestral State Reconstruction Gradient

## Likelihood Computation

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Initialization: for tips

$$f_{Y_i|Y_i}(Y_i;a) = \Phi_{Y_i,0}(a)$$

Propagation:

$$f_{\mathbf{Y}^{j}|X_{j}}(\mathbf{Y}^{j};a) = \prod_{l=1}^{L} f_{\mathbf{Y}^{j_{l}}|X_{j}}(\mathbf{Y}^{j_{l}};a)$$
 conditional independence



Likelihood Ancestral State Reconstruction Gradient

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Likelihood function:

$$f_{\mathbf{Y}^{j}|X_{j}}(\mathbf{Y}^{j};a) = A_{j}(\mathbf{Y}^{j})\Phi_{M_{j}(\mathbf{Y}^{j}),S_{j}^{2}(\mathbf{Y}^{j})}(a)$$

Initialization: for tips

$$f_{Y_i|Y_i}(Y_i;a) = \Phi_{Y_i,0}(a)$$

Propagation:

 $f_{\mathbf{Y}^{j}|X_{j}}(\mathbf{Y}^{j};a) = \prod_{l=1}^{L} f_{\mathbf{Y}^{j}|X_{j}}(\mathbf{Y}^{j};a) \text{ conditional independence}$ 

$$f_{\mathbf{Y}^{j_l}|X_j}(\mathbf{Y}^{j_l}; a) = \int_{\mathbb{R}} f_{\mathbf{Y}^{j_l}|X_{j_l}}(\mathbf{Y}^{j_l}; b) f_{X_{j_l}|X_j}(b; a) db \quad \text{explicit (Gaussian)}$$



#### Ancestral State Reconstruction



#### Ancestral State Reconstruction: compute $\mathbb{E}[\mathbf{Z} \mid \mathbf{Y}]$

### Ancestral State Reconstruction: Naive Gaussian

$$\begin{pmatrix} \textbf{Z} \\ \textbf{Y} \end{pmatrix} \sim \mathcal{N} \left( \begin{pmatrix} \boldsymbol{\mu}_{\textbf{Z}} \\ \boldsymbol{\mu}_{\textbf{Y}} \end{pmatrix}; \quad \begin{pmatrix} \textbf{V}_{\textbf{Z}\textbf{Z}} & \textbf{V}_{\textbf{Z}\textbf{Y}} \\ \textbf{V}_{\textbf{Y}\textbf{Z}} & \textbf{V}_{\textbf{Y}\textbf{Y}} \end{pmatrix} \right)$$

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Ancestral State Reconstruction: Naive Gaussian

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# $\mathbf{Z} \mid \mathbf{Y} ~\sim \mathcal{N} \left( \boldsymbol{\mu}_{\mathbf{Z}} + \mathbf{V}_{\mathbf{Z}\mathbf{Y}}\mathbf{V}_{\mathbf{Y}\mathbf{Y}}^{-1}\boldsymbol{\mu}_{\mathbf{Y}}; ~ \mathbf{V}_{\mathbf{Z}\mathbf{Z}} - \mathbf{V}_{\mathbf{Z}\mathbf{Y}}\mathbf{V}_{\mathbf{Y}\mathbf{Y}}^{-1}\mathbf{V}_{\mathbf{Y}\mathbf{Z}} \right)$

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Ancestral State Reconstruction: Naive Gaussian

$$\begin{pmatrix} \mathbf{Z} \\ \mathbf{Y} \end{pmatrix} \sim \mathcal{N} \left( \begin{pmatrix} \boldsymbol{\mu}_{\mathbf{Z}} \\ \boldsymbol{\mu}_{\mathbf{Y}} \end{pmatrix}; \quad \begin{pmatrix} \mathbf{V}_{\mathbf{Z}\mathbf{Z}} & \mathbf{V}_{\mathbf{Z}\mathbf{Y}} \\ \mathbf{V}_{\mathbf{Y}\mathbf{Z}} & \mathbf{V}_{\mathbf{Y}\mathbf{Y}} \end{pmatrix} \right)$$

$$\mathbf{Z} \mid \mathbf{Y} \sim \mathcal{N} \left( \boldsymbol{\mu}_{\mathbf{Z}} + \mathbf{V}_{\mathbf{Z}\mathbf{Y}} \mathbf{V}_{\mathbf{Y}\mathbf{Y}}^{-1} \boldsymbol{\mu}_{\mathbf{Y}}; \ \mathbf{V}_{\mathbf{Z}\mathbf{Z}} - \mathbf{V}_{\mathbf{Z}\mathbf{Y}} \mathbf{V}_{\mathbf{Y}\mathbf{Y}}^{-1} \mathbf{V}_{\mathbf{Y}\mathbf{Z}} \right)$$

 $\rightarrow$  requires a matrix inversion  $O(n^3)$ .

ayesian for Continuous Traits Likelihood Pruning Algorithms Ancestral State Reconstruction Phylogeography Gradient

### Ancestral State Reconstruction: Efficient Computations



Conditional Moments: in one pre-order traversal.

$$\hookrightarrow \mathbb{E} \, [\, \mathbf{Z}_j \mid \mathbf{Y} \,], \, \mathbb{V} \text{ar} \, [\, \mathbf{Z}_j \mid \mathbf{Y} \,]. \\ \hookrightarrow O(N).$$

Likelihood Ancestral State Reconstruction Gradient

### Conditional Moments Computation

Conditional Likelihood function:

$$f_{X_j \mid \mathbf{Y}}(a; \mathbf{Y}) \propto \Phi_{N_j, T_j^2}(a)$$



Likelihood Ancestral State Reconstruction Gradient

### Conditional Moments Computation

Conditional Likelihood function:

$$f_{X_j \mid \mathbf{Y}}(a; \mathbf{Y}) \propto \Phi_{N_j, \mathcal{T}_j^2}(a)$$

Initialization: for the root

$$f_{X_r|\mathbf{Y}}(a;\mathbf{Y}) \propto f_{\mathbf{Y}|X_r}(\mathbf{Y};a)f_{X_r}(a)$$



Likelihood Ancestral State Reconstruction Gradient

## Conditional Moments Computation

Conditional Likelihood function:

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Initialization: for the root

$$f_{X_r|\mathbf{Y}}(a;\mathbf{Y}) \propto f_{\mathbf{Y}|X_r}(\mathbf{Y};a)f_{X_r}(a)$$

Propagation:

$$f_{X_j \mid \mathbf{Y}}(a; \mathbf{Y}) = \int_{\mathbb{R}} f_{X_j \mid X_{pa(j)}}(a; b) f_{X_{pa(j)} \mid \mathbf{Y}}(b; \mathbf{Y}) db$$
 Gaussians



Aayesian for Continuous Traits Likelihood Pruning Algorithms Ancestral State Reconstruction Phylogeography Gradient

### Efficient Computations: Gradient



Branch-specific Gradient: in one pre-order traversal.

$$\begin{aligned} \nabla_{\boldsymbol{\theta}_{k}}\left[\log p\left(\mathbf{Y} \mid \boldsymbol{\theta}_{k}\right)\right] &= \mathbb{E}\left[\nabla_{\boldsymbol{\theta}_{k}}\left[\log p\left(\mathbf{Z}^{k}, \mathbf{Y} \mid \boldsymbol{\theta}_{k}\right)\right] \mid \mathbf{Y}\right] \\ &= \mathbb{E}\left[\nabla_{\boldsymbol{\theta}_{k}}\left[\log p\left(\mathbf{Z}^{k} \mid \mathbf{Y}_{\lceil k \rceil}, \boldsymbol{\theta}_{k}\right)\right] \mid \mathbf{Y}\right]. \end{aligned}$$

ayesian for Continuous Traits Likelihood Pruning Algorithms Ancestral State Reconstruction Phylogeography Gradient

### Efficient Computations: Gradient



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Chain rule: Gradient w.r.t. any parameter in O(N).

ayesian for Continuous Traits Likelihood Pruning Algorithms Ancestral State Reconstruction Phylogeography Gradient

### Efficient Computations: Gradient



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Chain rule: Gradient w.r.t. any parameter in O(N).  $\rightarrow \blacksquare$ 

Phylogeography Relaxed Random Walk

### Outline

#### Bayesian for Continuous Traits

Pruning Algorithms

#### O Phylogeography

- Phylogeography
- Relaxed Random Walk

Phylogeography Relaxed Random Walk

# Phylogeography

Trait: Geographical coordinates: latitude and longitude.

Phylogeography Relaxed Random Walk

## Phylogeography

Trait: Geographical coordinates: latitude and longitude.



Phylogeography Relaxed Random Walk

## Phylogeography



Spread on an heterogeneous geographical region.

### Relaxed Random Walk

(Lemey et al., 2010)

Model: BM with varying ("relaxed") variance.

$$\mathbf{Z}^{j} \mid \mathbf{Z}^{\mathsf{pa}(j)} \sim \mathcal{N}\left(\mathbf{Z}^{\mathsf{pa}(j)}, \phi_{j}\mathbf{R}\right)$$

Idea: Each branch has its own propagation speed.

 $\rightarrow$  Allow for varying speed of propagation.

### Relaxed Random Walk

(Lemey et al., 2010)

Model: BM with varying ("relaxed") variance.

$$\mathbf{Z}^{j} \mid \mathbf{Z}^{\mathsf{pa}(j)} \sim \mathcal{N}\left(\mathbf{Z}^{\mathsf{pa}(j)}, \phi_{j}\mathbf{R}\right)$$

Idea: Each branch has its own propagation speed.

 $\rightarrow$  Allow for varying speed of propagation.

Regularisation: Set a prior on the  $\phi_j \sim \mathcal{L}(s)$ 

Examples:

$$\phi_j \sim \mathsf{Gamma}(
u/2, 
u/2)$$
  
 $\phi_j \sim \mathsf{LogNormal}(1, \sigma)$ 

Phylogeography Relaxed Random Walk

#### Relaxed Random Walk

(Lemey et al., 2010)

$$\mathbf{Z}^{j} \mid \mathbf{Z}^{\mathsf{pa}(j)} \sim \mathcal{N}\left(\mathbf{Z}^{\mathsf{pa}(j)}, \phi_{j}\mathbf{R}\right) \qquad \phi_{j} \sim \mathcal{L}(s)$$

 $p(\mathbf{R}, \phi, s, \mathcal{T}, \psi \mid \mathbf{Y}, \mathbf{S}) \propto p(\mathbf{Y}, \mathbf{S} \mid \mathbf{R}, \phi, s, \mathcal{T}, \psi) p(\mathbf{R}, \phi, s, \mathcal{T}, \psi)$ 

Phylogeography Relaxed Random Walk

### Relaxed Random Walk

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$$\mathbf{Z}^{j} \mid \mathbf{Z}^{\mathsf{pa}(j)} \sim \mathcal{N}\left(\mathbf{Z}^{\mathsf{pa}(j)}, \phi_{j}\mathbf{R}\right) \qquad \phi_{j} \sim \mathcal{L}(s)$$

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 $\propto p(\mathbf{Y} \mid \mathbf{R}, \phi, s, T) p(\mathbf{S} \mid T, \psi) p(\mathbf{R}, \phi, s, T, \psi)$ 

Phylogeography Relaxed Random Walk

### Relaxed Random Walk

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 $\propto$   $p(\mathbf{Y} \mid \mathbf{R}, \phi, s, T) p(\mathbf{S} \mid T, \psi) p(\mathbf{R}, \phi, s, T, \psi)$ 

$$\propto p(\mathbf{Y} \mid \mathbf{R}, \phi, \mathcal{T}) p(\phi \mid s) p(s) p(\mathbf{R}) \\ \times p(\mathbf{S} \mid \mathcal{T}, \psi) p(\mathcal{T}, \psi)$$

Phylogeography Relaxed Random Walk

# Phylogeography



#### Relaxed Random Walk

- Simple model.
- Flexible: accomodate for heterogeneous landscape.
- Ancestral state reconstruction: geopraphical spread of the epidemy.

Phylogeography Relaxed Random Walk

### Phylogeography: post-hoc analyses



Paul Bastide
Betancourt. 2017. arXiv e-print. p. 170102434.

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# Thank you for listening



Institut Montpelliérain Alexander Grothendieck

# Appendices

General Framework BM Gibbs Constrained Spaces

# LKJ Distribution

(Lewandowski, Kurowicka, and Joe, 2009)

Idea:

Use two different distributions for the variance and the correlation.

General Framework BM Gibbs Constrained Spaces

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Use two different distributions for the variance and the correlation.

Decomposition: Use correlation matrix C

$$\mathbf{R} = \begin{pmatrix} \sigma_1 & 0 \\ & \ddots & \\ 0 & & \sigma_p \end{pmatrix} \begin{pmatrix} 1 & C_{kl} \\ & \ddots & \\ C_{kl} & 1 \end{pmatrix} \begin{pmatrix} \sigma_1 & 0 \\ & \ddots & \\ 0 & & \sigma_p \end{pmatrix}$$

General Framework BM Gibbs Constrained Spaces

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 $\sigma$ : Real positive ightarrow log-normal, gamma, half Gaussian, ...

General Framework BM Gibbs Constrained Spaces

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 $\textbf{C}: \mbox{ Correlation matrix} \rightarrow "LKJ" \mbox{ distribution}.$ 

General Framework BM Gibbs Constrained Spaces

# LKJ Distribution

(Lewandowski, Kurowicka, and Joe, 2009)

LKJ distribution:

$$\mathsf{LKJ}(\mathsf{C} \mid \eta) = c_{\mathsf{p}}(\eta) \left| \mathsf{C} \right|^{\eta-1}$$

- $\eta = 1$ : Uniform.
- $\eta > 1$ : Peak around identity matrix.
- $0 < \eta < 1$ : Trough around identity matrix.

General Framework BM Gibbs Constrained Spaces

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 $\label{eq:problem: Problem: We cannot sample each coefficient of $\mathbf{C}$ independently.} \\ \hookrightarrow $\mathbf{C}$ has a complex structure.}$ 

Solution: Metropolis-Hasting in a constrained space.

General Framework BM Gibbs Constrained Spaces

### MH in constrained space

Transformation:

$$f:egin{cases} \mathcal{C}_q o \mathbb{R}^q \ oldsymbol{ heta} \mapsto oldsymbol{
u} = f(oldsymbol{ heta}) \end{cases}$$

General Framework BM Gibbs Constrained Spaces

## MH in constrained space

Transformation:

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Distribution: For a distribution  $\pi$ :

$$\pi_{\boldsymbol{ heta}}(\boldsymbol{ heta}) = \pi_{\boldsymbol{
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General Framework BM Gibbs Constrained Spaces

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Metropolis - Hasting: Iterate:

- Draw  $\boldsymbol{\nu}^*$  in  $q(\boldsymbol{\nu} \mid \boldsymbol{\nu}^t)$ .
- Set  $heta^{(t+1)}= heta^*=f^{-1}(
  u^*)$  with probability

$$r_{t} = \min\left\{1, \frac{p\left(\mathbf{Y} \mid \boldsymbol{\theta}^{*}\right)}{p\left(\mathbf{Y} \mid \boldsymbol{\theta}^{t}\right)} \frac{p\left(\boldsymbol{\theta}^{*}\right)}{p\left(\boldsymbol{\theta}^{t}\right)} \frac{q(\boldsymbol{\nu}^{(t)} \mid \boldsymbol{\nu}^{*})}{q(\boldsymbol{\nu}^{*} \mid \boldsymbol{\nu}^{(t)})} \frac{|J_{f}(\boldsymbol{\theta}^{(t)})|}{|J_{f}(\boldsymbol{\theta}^{*})|}\right\}$$

General Framework BM Gibbs Constrained Spaces

Variance: LKJ transformation (Lewandowski, Kurowicka, and Joe, 2009)

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 $\sigma$ : Real positive

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- $\sigma$ : Real positive
- C: Correlation matrix
- LKJ: Transformation on the space of correlation matrices.
  - $\hookrightarrow$  Use "vine" theory.
  - $\hookrightarrow$  Easier and more efficient: Cholesky representation.

Bayesian for Continuous Traits Pruning Algorithms Phylogeography General Framework BM Gibbs Constrained Spaces

Variance: LKJ transformation (Lewandowski, Kurowicka, and Joe, 2009)

$$\mathbf{C} = \mathbf{W}^{T} \mathbf{W} = \begin{pmatrix} 1 & W_{12} & \cdots & W_{1p} \\ & W_{22} & & \vdots \\ & & \ddots & \vdots \\ 0 & & & W_{pp} \end{pmatrix}^{T} \begin{pmatrix} 1 & W_{12} & \cdots & W_{1p} \\ & W_{22} & & \vdots \\ & & \ddots & \vdots \\ 0 & & & & W_{pp} \end{pmatrix}$$

With:

$$W_{1k}^2 + \dots + W_{kk}^2 = 1$$
 (correlation)  

$$W_{kk} > 0$$
 (identifiability)

Bayesian for Continuous Traits Pruning Algorithms Phylogeography General Framework BM Gibbs Constrained Spaces

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With:

$$\label{eq:w1k} \begin{array}{ll} \bullet & W_{1k}^2 + \cdots + W_{kk}^2 = 1 \\ \bullet & W_{kk} > 0 \end{array} \hspace{1.5cm} ( \mbox{correlation} ) \\ \mbox{(identifiability)} \end{array}$$

 $\hookrightarrow$  Each column k is in the half euclidian unit sphere  $\mathcal{S}_k^h(\mathbb{R})$ .

General Framework BM Gibbs Constrained Spaces

## Sampling in the Sphere

 $\rightarrow$  Sphere  $\mathcal{S}_k^{\mathsf{h}}(\mathbb{R})$  to euclidean ball  $\mathcal{B}_{k-1}(\mathbb{R})$ :

$$\mathsf{F}: \begin{cases} \mathcal{B}_{k-1}(\mathbb{R}) & \to \mathcal{S}_{k}^{\mathsf{h}}(\mathbb{R}) \\ \mathsf{V}_{\cdot k} & \mapsto \mathsf{W}_{\cdot k} = \left(\mathsf{V}_{\cdot k}, \sqrt{1 - \|\mathsf{V}_{\cdot k}\|^{2}}\right) \end{cases}$$



General Framework BM Gibbs Constrained Spaces

## Sampling in the Sphere

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 $\mathcal{S}_{2}^{b(\mathbb{R})}$   $W_{2}$  $\mathcal{B}_{1}(\mathbb{R})$   $V_{2}$ 

 $o \mathcal{B}_{k-1}(\mathbb{R})$  to infinite-norm ball  $\mathcal{B}_{k-1}^\infty(\mathbb{R})$  : "LKJ"

$$\mathsf{LKJ}_i(\mathbf{z}) = \begin{cases} z_i & \text{if } i = 1 \\ z_i \prod_{k=1}^{i-1} (1 - z_k^2)^{1/2} & \text{if } 1 < i \le k-1. \end{cases}$$



General Framework BM Gibbs Constrained Spaces

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 Sphere  $\mathcal{S}_k^{\mathsf{h}}(\mathbb{R})$  to euclidean ball  $\mathcal{B}_{k-1}(\mathbb{R})$ :

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$$o \mathcal{B}_{k-1}(\mathbb{R})$$
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 $o \mathcal{B}^\infty_{k-1}(\mathbb{R})$  to  $\mathbb{R}^{k-1}$ : "Fisher Z" transform

$$anh^{-1}: egin{cases} ]-1\,,1[
ightarrow \mathbb{R} \ x\mapsto rac{1}{2}\ln\left(rac{1+x}{1-x}
ight) \end{cases}$$

Variance: LKJ transformation (Lewandowski, Kurowicka, and Joe, 2009)

Decomposition: Use correlation matrix C

$$\mathbf{R} = \begin{pmatrix} \sigma_1 & 0 \\ & \ddots & \\ 0 & & \sigma_p \end{pmatrix} \begin{pmatrix} 1 & C_{kl} \\ & \ddots & \\ C_{kl} & 1 \end{pmatrix} \begin{pmatrix} \sigma_1 & 0 \\ & \ddots & \\ 0 & & \sigma_p \end{pmatrix}$$

- $\sigma$ : Real positive
- C: Correlation matrix

LKJ transformation: Smooth transformation from  $\mathbb{R}^{p(p-1)/2}$  to the space of correlation matrices.

Variance: LKJ transformation (Lewandowski, Kurowicka, and Joe, 2009)

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- $\sigma$ : Real positive
- C: Correlation matrix

LKJ transformation: Smooth transformation from  $\mathbb{R}^{p(p-1)/2}$  to the space of correlation matrices.

Jacobian: Can be computed.

General Framework BM Gibbs Constrained Spaces

## OU: No Gibbs

Likelihood:

 $\mathbf{Y}|\mathbf{A},\mathbf{R},\boldsymbol{\mu}
eq\mathcal{M}\mathcal{N}$ 

General Framework BM Gibbs Constrained Spaces

### OU: No Gibbs

Likelihood:

 $\mathbf{Y}|\mathbf{A}, \mathbf{R}, \boldsymbol{\mu} \not\sim \mathcal{MN}$ 

#### Conjugate Priors: ???

General Framework BM Gibbs Constrained Spaces

## OU: No Gibbs

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Gibbs: Not possible.

General Framework BM Gibbs Constrained Spaces

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Conjugate Priors: ???

Gibbs: Not possible.

Metropolis - Hasting:

Need to sample in constrained spaces (A, R).

back

Likelihood Ancestral State Reconstruction Gradient

### Hamiltonian Monte Carlo

(Betancourt, 2017)

Idea: Introduce the "momentum"  $\boldsymbol{p}$  of the parameters  $\boldsymbol{q}.$ 

Likelihood Ancestral State Reconstruction Gradient

## Hamiltonian Monte Carlo

(Betancourt, 2017)

Idea: Introduce the "momentum"  ${\boldsymbol{p}}$  of the parameters  ${\boldsymbol{q}}.$ 

Hamiltonian  $H(\mathbf{q}, \mathbf{p}) =$  Potential energy + Kinetic energy

Likelihood Ancestral State Reconstruction Gradient

## Hamiltonian Monte Carlo

(Betancourt, 2017)

Idea: Introduce the "momentum"  $\boldsymbol{p}$  of the parameters  $\boldsymbol{q}.$ 

Hamiltonian  $H(\mathbf{q}, \mathbf{p}) = \text{Potential energy} + \text{Kinetic energy}$ =  $U(\mathbf{q}) + K(\mathbf{p})$ =  $-\log \underbrace{p(\mathbf{Y}|\mathbf{q})p(\mathbf{q})}_{\text{posterior}} - \log \underbrace{\phi(\mathbf{p})}_{\text{Gaussian}}$
Bayesian for Continuous Traits Pruning Algorithms Phylogeography Likelihood Ancestral State Reconstruction Gradient

## Hamiltonian Monte Carlo

(Betancourt, 2017)

Idea: Introduce the "momentum"  $\boldsymbol{p}$  of the parameters  $\boldsymbol{q}.$ 

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H total energy invariant by Hamiltonian dynamic:

$$\begin{cases} \frac{d\mathbf{p}}{dt} = \nabla_{\mathbf{q}} U(\mathbf{q}) \\ \frac{d\mathbf{q}}{dt} = -\nabla_{\mathbf{p}} K(\mathbf{p}) \end{cases}$$

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Hamiltonian  $H(\mathbf{q}, \mathbf{p}) =$  Potential energy + Kinetic energy  $= U(\mathbf{q}) + K(\mathbf{p})$  $= -\log \underbrace{p(\mathbf{Y}|\mathbf{q})p(\mathbf{q})}_{i} - \log \underbrace{\phi(\mathbf{p})}_{i}$ *H* total energy invariant by Hamiltonian dynamic:  $\begin{cases} \frac{d\mathbf{p}}{dt} = \nabla_{\mathbf{q}} U(\mathbf{q}) \\ \frac{d\mathbf{q}}{dt} = -\nabla_{\mathbf{p}} \mathcal{K}(\mathbf{p}) \end{cases}$  Draw random moments p. Propose a new q from the dynamic.