# TP 3: Hasting-Metropolis (and Gibbs) samplers

### Exercise 1: Hasting-Metropolis within Gibbs - Stochastic Approximation EM

We observe a group of N (independent) individuals. For the i-th individual, we have  $k_i$  measurements  $y_{i,j} \in \mathbb{R}$ . In studies on the progression of diseases, measurements  $y_{i,j}$  can be measures of weight, volume of brain structures, protein concentration, tumoral score, etc. over time. We assume that each measurement  $y_{i,j}$  are independent and obtained at time  $t_{i,j}$  with  $t_{i,1} < \ldots < t_{i,k_i}$ .

#### 1.A - A population model for longitudinal data

We wish to model an average progression as well as individual-specific progressions of the measurements from the observations  $(y_{i,j})_{i \in [\![1,N]\!], j \in [\![1,k_i]\!]}$ . To do that, we consider a hierarchical model defined as follows.

i. We assume that the average trajectory is the straight line which goes through the point  $p_0$  at time  $t_0$  with velocity  $v_0$ 

$$d(t) := p_0 + v_0(t - t_0)$$

where

$$p_0 \sim \mathcal{N}(\overline{p_0}, \sigma_{p_0}^2)$$
 ;  $t_0 \sim \mathcal{N}(\overline{t_0}, \sigma_{t_0}^2)$  ;  $v_0 \sim \mathcal{N}(\overline{v_0}, \sigma_{v_0}^2)$ 

and  $\sigma_{p_0}$ ,  $\sigma_{t_0}$ ,  $\sigma_{v_0}$  are fixed variance parameters. While we consider straight lines, we can also fix  $p_0$ .

ii. For the *i*-th individual, we assume a trajectory of progression of the form

$$d_i(t) := d(\alpha_i(t - t_0 - \tau_i) + t_0).$$

The trajectory of the *i*-th individual corresponds to an affine reparametrization of the average trajectory. This affine reparametrization, given by  $t \mapsto \alpha_i(t-t_0-\tau_i)+t_0$ , allows to characterize changes in speed and delay in the progression of the *i*-th individual with respect to the average trajectory. Moreover, we assume that for all *i*-th individual measurements

$$\begin{cases} y_{i,j} = d_i(t_{i,j}) + \varepsilon_{i,j} & \text{where} \quad \varepsilon_{i,j} \overset{\text{i.i.d.}}{\sim} \mathcal{N}(0, \sigma^2) \\ \alpha_i = \exp(\xi_i) & \text{where} \quad \xi_i \overset{\text{i.i.d.}}{\sim} \mathcal{N}(0, \sigma_{\xi}^2) \\ \tau_i \overset{\text{i.i.d.}}{\sim} \mathcal{N}(0, \sigma_{\tau}^2) \end{cases} .$$

The parameters of the model are  $\theta = (\overline{t_0}, \overline{v_0}, \sigma_{\xi}, \sigma_{\tau}, \sigma)$ . For all  $i \in [1, N]$ ,  $z_i = (\alpha_i, \tau_i)$  are random variables called random effects and  $z_{pop} = (t_0, v_0)$  are called fixed effects. The fixed effects are used to model the group progression whereas random effects model individual progressions. Likewise, we define  $\theta_i = (\sigma_{\xi}, \sigma_{\tau}, \sigma)$  and  $\theta_{pop} = (\overline{t_0}, \overline{v_0})$ .

We consider a bayesian framework and assume the following a priori on the parameters  $\theta$ :

$$\overline{t_0} \sim \mathcal{N}(\overline{\overline{t_0}}, s_{t_0}^2) \quad ; \quad \overline{v_0} \sim \mathcal{N}(\overline{\overline{v_0}}, s_{v_0}^2)$$

$$\sigma_{\xi}^2 \overset{\text{i.i.d.}}{\sim} \mathcal{W}^{-1}(v_{\xi}, m_{\xi}) \quad ; \quad \sigma_{\tau}^2 \overset{\text{i.i.d.}}{\sim} \mathcal{W}^{-1}(v_{\tau}, m_{\tau}) \quad ; \quad \sigma^2 \overset{\text{i.i.d.}}{\sim} \mathcal{W}^{-1}(v, m) \, .$$

where  $\mathcal{W}^{-1}(v,m)$   $(v>0, m\in\mathbb{N}^*)$  is the inverse-Wishart distribution :

$$f_{\mathcal{W}^{-1}}(\sigma^2) = \frac{1}{\Gamma(\frac{m}{2})} \frac{1}{\sigma^2} \left(\frac{v}{\sigma\sqrt{2}}\right)^m \exp\left(-\frac{v^2}{2\sigma^2}\right).$$

- 1. Write the complete log-likelihood of the previous model  $\log q(y, z, \theta)$  and show that the proposed model belongs to the curved exponential family.
- 2. Generate synthetic data from the model by taking some reasonable values for the parameters.

# 1.B - HM-SAEM - Hasting-Metropolis sampler

In order to estimate – by a maximum a posteriori for instance – the parameters of this statistical model, we will use the SAEM – Stochastic Approximation EM algorithm. However, this algorithm requires that we are able to sample from the a posteriori distribution, see algorithm 2.

We will use the Hasting-Metropolis algorithm to that end: actually a direct sampling is not possible in our context. Let q(.|z) be the proposal distribution of the algorithm, *i.e.* the conditional probability of proposing a state  $z^*$  given the current state z, and  $\pi$  be a density defined on an open set  $\mathcal{U}$  of  $\mathbb{R}^n$ . The Hasting-Metropolis algorithm targeting  $\pi$  writes:

### Algorithm 1: Hasting-Metropolis Sampler

- 1 Given  $z^{(0)}$ 2 for k=0 to maxIter do
  3  $\#Proposal: z^* \sim q(.|z^{(k)})$ 4  $\#Acceptance\text{-}Rejection: \alpha(z^{(k)},z^*) = \min\left(1,\frac{q(z^{(k)}|z^*)\pi(z^*)}{q(z^*|z^{(k)})\pi(z^{(k)})}\right)$ 5  $z^{(k+1)} = \begin{cases} z^* \text{ with probability } \alpha(z^{(k)},z^*) \\ z^{(k)} \text{ with probability } 1 \alpha(z^{(k)},z^*) \end{cases}$ 6 end
  - **3.** Propose a Metropolis-Hastings sampler to sample from the *a posteriori* distribution of the latent variable  $z=(z_{pop},z_i)_{i\in \llbracket 1,N\rrbracket}=(t_0,v_0,\xi_i,\tau_i)_{i\in \llbracket 1,N\rrbracket}\in \mathbb{R}^{2N+2}.$

A natural choice for the proposal distribution is to consider a multivariate Gaussian distribution  $\mathcal{N}(z, \sigma_{prop})$ . Thus, the acceptance ratio simply writes  $1 \wedge \frac{\pi(z^*)}{\pi(z^{(k)})}$ . This algorithm is called Symmetric Random Walk Hasting-Metropolis algorithm.

We want to use the Expectation-Maximization algorithm to maximize the likelihood, especially as we have proved at the question 1 that the model belongs to the curved exponential family. Nevertheless, the expectation required by the EM algorithm  $Q_k(\theta^{(k)}) = \mathbb{E}_y\left[q(y,z,\theta^{(k)})|y,\theta^{(k)}\right]$  cannot be calculated here, due to the latent variable z. So, we have to use a stochastic version of the EM algorithm, namely the SAEM algorithm: the Expectation step is split into two steps, the Simulation one and the Stochastic Approximation one, see Algorithm 2.

### 4. Compute the optimal parameters

$$\theta^{(k)} = \operatorname*{argmax}_{\theta \in \Theta} \left\{ -\Phi(\theta) + \langle S_k \mid \Psi(\theta) \rangle \right\}$$

and implement the HM-SAEM in order to find the MAP. In particular, we assume that the MAP exists. Use the question 2 to check your algorithm.

For step-sizes  $\varepsilon_k$  we can choose a parameter  $N_b$  – burn-in parameter – and define

$$\forall k \in \mathbb{N}, \qquad \left\{ \begin{array}{ll} 1 & \text{if } k \in [\![1,N_b]\!] \\ (k-N_b)^{-\alpha} & \text{otherwise} \end{array} \right.$$

where  $\alpha \in [\frac{1}{2}, 1]$  is necessary to ensure the convergence of the MCMC-SAEM. See [AKT10, AK15].

**Remark**: Contrary to Bayesian inference, where burn-in traditionally refers to a certain amount of samples which are discarded, here the term burn-in refers to memoryless approximation steps. In other words, during the burn-in phase, the information contained in  $z^{(k)}$  is not used in the approximation of the sufficient statistics. In practice, the burn-in period is chosen to be half of the maximum number of iterations.

### Algorithm 2: MCMC-SAEM (for curved exponential family)

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1 Given data y and intial guess \theta^{(0)}

2 #Initialization : z^{(0)} = 0, S^{(0)} = 0 and step-sizes (\varepsilon_k)_{k\geqslant 0}.

3 for k=0 to maxIter do

4 | #Simulation z^{(k+1)} \sim q(.|y,\theta^{(k)}) (sampler intialized at z^{(k)})

5 | #Stochastic Approximation : S^{(k+1)} = S^{(k)} + \varepsilon_k \left( S(y,z^{(k+1)}) - S^{(k)} \right),

6 | #Maximization : \theta^{(k+1)} = \underset{\theta \in \Theta}{\operatorname{argmax}} \left\{ -\Phi(\theta^{(k)}) + \left\langle S^{(k+1)} \mid \Psi(\theta^{(k)}) \right\rangle \right\}

7 end
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#### 1.C - HMwG-SAEM - Hasting-Metropolis within Gibbs sampler

However, the dimension of the latent variable z may become high if we consider a large cohort and so the a posteriori distribution of the latent variable difficult to sample. In that case, we can use a Gibbs sampler which consists in generating an instance from the distribution of each (sub)-variable in turn, conditional on the current values of the other (sub)-variables. Gibbs sampling is more generally applicable when the joint distribution is not known explicitly or is difficult to sample from directly,

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but the conditional distribution of each variable is known and is easy (or at least, easier) to sample from.

If we consider  $\pi$ , a density defined on an open set  $\mathcal{U}$  of  $\mathbb{R}^n$   $(n \ge 2)$  and if we denote, for  $\ell \in [1, n]$ ,  $\pi_{\ell}$  the  $\ell^{\text{th}}$  full conditional of  $\pi$  given by

$$\pi_{\ell}(z_{\ell} \mid z_{-\ell}) \propto \pi(z)$$

where  $z_{-\ell} = \{z_1, \dots, z_{\ell-1}, z_{\ell+1}, \dots, z_n\}$ , we recall that the classical Gibbs sampler writes as follows:

## Algorithm 3: Gibbs Sampler

1 Given  $z^{(k)} = (z_1^{(k)}, \dots, z_n^{(k)})$ 2 for  $\ell = 1$  to n do
3  $| z_{\ell}^{(k+1)} \sim \pi_{\ell}(z_{\ell} | z_1^{(k+1)}, \dots, z_{\ell-1}^{(k+1)}, z_{\ell+1}^{(k)}, \dots, z_n^{(k)})$ 

When direct sampling from the full conditionals is not possible, the step  $(\star)$  is often replaced with a Metropolis-Hastings step. The resulting MCMC algorithm is called *hybrid Gibbs sampler* or *Metropolis-Hastings within Gibbs sampler*.

- **5.** Propose a Metropolis-Hastings within Gibbs sampler to sample from the *a posteriori* distribution of  $z_i = (\xi_i, \tau_i)$ .
- **6.** Likewise, propose a HMwG sampler for the a posteriori distribution of  $z_{pop} = (t_0, v_0)$ .
- 7. Implement the HMwG-SAEM in order to find the MAP.

We can improve the sampling step for big dataset by considering a Block HMwG sampler instead of a "one-at-a-time" as described above HMwG sampler. In the Block version, each Metropolis-Hastings step of the algorithm consists in a multivariate symmetric random walk. Then, the Block MHwG sampler updates simultaneously block (or sets) of latent variables given the others.

- **8.** Explain what is the advantages of a Block Gibbs sampler over a "one-at-a-time" Gibbs sampler for our model.
- **9.** Implement a Block HMwG sampler by choosing a block for the fixed effects and a block by individuals.

The model studied in this exercise is a very simplified version of the model proposed by Jean-Baptiste Schiratti in his Phd-Thesis. For more details, you can refer to [SACD15, Sch16, COA17].

#### **Exercise 2: Multiplicative Hasting-Metropolis**

Let f be a density function on ]-1,1[. We consider the multiplicative Hasting-Metropolis algorithm defined as follows.

Let X be the current state of the Markov chain.

- (i) We sample  $\varepsilon$  from the probability density function f and a random variable  $\mathcal B$  from the Bernoulli distribution with parameter  $\frac{1}{2}$ .
- (ii) If  $\mathcal{B} = 1$ , we set  $Y = \varepsilon X$ . Otherwise, we set  $Y = \frac{X}{\varepsilon}$ . Then, we accept the candidate Y with a probability given by  $\alpha(X,Y)$ , the usual Hasting-Metropolis acceptation ratio.
- **1.** Determine the density of the jumping distribution  $Y \sim q(X,Y)$ .
- **2.** Compute the acceptation ratio  $\alpha$  so that the chain has a given distribution  $\pi$  as invariant distribution.
- 3. Implement this sampler for two different target distributions: the first one being a distribution from which we can sample using the inverse transform method and the second one is of your choice.
- **4.** Evaluate, in each case, the match of your samples with the true distribution.

### **Exercise 3: Data augmentation**

Let  $f:(x,y)\in\mathbb{R}^p\times\mathbb{R}^q\mapsto f(x,y)\in\mathbb{R}^+$  be a density with respect to the Lebesgue measure on  $\mathbb{R}^{p+q}$ . Let us define

$$f_X(x) := \int f(x, y) dy$$
;  $f_Y(y) = \int f(x, y) dx$ ;

and

$$\forall y \in \mathbf{Y} := \{ y \in \mathbb{R}^q \mid f_Y(y) > 0 \}, \qquad f_{X|Y}(x,y) := \frac{f(x,y)}{f_Y(y)} ;$$

$$\forall x \in \mathbf{X} := \{ x \in \mathbb{R}^p \mid f_X(x) > 0 \}, \qquad f_{X|Y}(x,y) := \frac{f(x,y)}{f_Y(y)}.$$

We define a bivariate chain  $\{(X_n, Y_n), n \ge 0\}$  as in the following algorithm.

### Algorithm 4: Data augmentation

- 1 Given  $(X_0, Y_0) \in \mathbb{R}^p \times \mathbb{R}^q$  and  $N \in \mathbb{N}$
- $\mathbf{2} \ \mathbf{for} \ n = 1 \ to \ N \ \mathbf{do}$
- $\mathbf{3} \mid X_n \sim f_{X|Y}(\cdot, Y_{n-1})$
- $4 \mid Y_n \sim f_{Y|X}(X_n, \cdot)$
- 5 end
- 6 return  $\{(X_n, Y_n), 0 \leqslant n \leqslant N\}$ 
  - 1. Show that the bivariate process  $\{(X_n, Y_n), n \ge 0\}$  is a Markov chain. Give the expression of its transition kernel as a function of the quantities defined above.

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**2.** Show that  $\{Y_n, n \ge 0\}$  is a Markov chain: give the expression of its transition kernel and prove that  $f_Y(y) dy$  is invariant for this kernel.

Hereafter, we consider the case when

$$f(x,y) = \frac{4}{\sqrt{2\pi}} y^{\frac{3}{2}} \exp\left[-y\left(\frac{x^2}{2} + 2\right)\right] \mathbbm{1}_{\mathbb{R}^+}(y)$$

**3.** Describe a Gibbs algorithm to approximate the distribution on  $\mathbb{R} \times \mathbb{R}$  with density f.

We can use a gamma distribution sampler: numpy.random.gamma in python or gamrnd from the Statistics and Machine Learning Toolbox in Matlab. We can also find a Matlab toolbox-free Gamma Generator in the Handbook of Monte Carlo Methods [KTB13]:

https://people.smp.uq.edu.au/DirkKroese/montecarlohandbook/probdist/.

**4.** Let H be a bounded function on  $\mathbb{R}$ . Explain how to approximate

$$\int_{\mathbb{R}} \frac{H(x)}{(4+x^2)^{\frac{5}{2}}} \, \mathrm{d}x$$

from the output  $\{(X_n, Y_n), 0 \le n \le N\}$  of this Gibbs sampler.

### References

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