Coursework 4M24: High-Dimensional MCMC

The performance of MCMC algorithms for high-dimensional problems is analysed in this coursework. The methods seen in the lecture notes: Gaussian random walk Metropolis Hastings (GRW-MH) and the preconditioned Crank-Nicolson (pCN) method) are compared, when sampling latent variables on a 2D domain. Code for the coursework supplied in functions.py contains useful functions for generating matrices, plotting, and skeleton code to be completed. The files simulation.py and spatial.py contain the code to set up the problem and relevant plotting examples.

Part I - Simulation

A Gaussian Process is defined on a 2D domain $\mathbf{x} \in [0,1] \times [0,1]$, and the latent variables are defined as $\mathbf{u} \sim \mathcal{N}(0,K)$, where $K_{ij} = k(\mathbf{x}_i,\mathbf{x}_j)$ and k is a Gaussian covariance function parametrised by a length-scale l.

$$k(\mathbf{x}, \mathbf{x}') = \exp\left(\frac{-\|\mathbf{x} - \mathbf{x}'\|^2}{2l^2}\right)$$

The coordinates of the latent variables $\{\mathbf{x}_i\}_{i=1}^N$ define the GP prior, and are placed on a regular $D \times D$ grid. We generate the data by observing a subsample of the latent variables, with added observation noise. The task will then be to use the noisy data \mathbf{v} to infer the values of all the latent variables \mathbf{u} .

$$\mathbf{v} = \widehat{G}\mathbf{u} + \epsilon$$
 subsample marriz.

Where the matrix G is generated based on the latent variable coordinate locations that are chosen to be observed through random subsampling - the function get_G generates this matrix from the subsampling indexes given by the function subsample. The observation noise is given by $\epsilon \sim \mathcal{N}(0, I)$.

- (a) Sample from the GP prior, and then the observation noise to simulate the data \mathbf{v} set the parameters D=16, l=0.3, subsample_factor = 4. Use the function plot_3D, comment on the effect of varying the length-scale parameter on the GP surface. Visualise the prior sample with the data overlaid using plot_result.
- (b) Complete the functions $\log_{\mathbf{v}}$ and $\log_{\mathbf{v}}$ continuous_likelihood, which should return $\ln p(\mathbf{u})$ and $\ln p(\mathbf{v}|\mathbf{u})$ respectively. Complete the TODO sections in the skeleton functions $\operatorname{\mathsf{grw}}$ and $\operatorname{\mathsf{pcn}}$, which are the GRW-MH and pCN algorithms see Lectures 5 & 11 for details. Sample from $p(\mathbf{u}|\mathbf{v})$ using both these algorithms (use $n=10000, \beta=0.2$), and plot the mean of the inferred \mathbf{u} alongside the data. Visualise the absolute error field between the original \mathbf{u} and the inferred \mathbf{u} , comparing the performance of the two algorithms. Comment on the GRW-MH and pCN acceptance rate for low and high-dimensional latent variable spaces (try D=4 and D=16). Vary the parameter β between 0 and 1 comment on the effect on the acceptance rate. What adverse affect would a very small β have on the samples?

Extension: Demonstrate the robustness of the pCN method to 'mesh refinement' ¹ by increasing the lettent dimension size N for a fixed dataset (i.e. refine the mesh), and comparing the acceptance rate to the GRW.

¹See Fig 1 from Cotter's paper - https://arxiv.org/abs/1202.0709

Now the model is extended to work on a probit classification problem. The data \mathbf{v} is put through a <u>probit transform</u> to give the vector t, where the ith component is given below. This gives the following

(c) Derive the full log-likelihood for the probit classification problem, and use this to complete the

function $\log_{probit_likelihood}$ to return $p(\mathbf{t}|\mathbf{u})$. Generate samples from the posterior $p(\mathbf{u}|\mathbf{t})$ using the pCN method. Predict the true class assignments $\mathbf{t}_{\text{true}} = \text{probit}(\mathbf{u})$, using a MC (Monte-Carlo) estimate of the predictive distribution (Complete the function predict_t). Visualise these simply using the plot_2D function, comparing the true assignments to the predictive probabilities.

Hint: The predictive distribution is given below, use the samples from the posterior to form a MC estimate of this distribution.

ibution.
$$\int_{\mathcal{U}} d\alpha \cdot \frac{1}{|\mathbf{t}|} d\mathbf{u} = \int_{\mathcal{U}} p(t^* = 1|\mathbf{u}|) p(\mathbf{u}|\mathbf{t}) d\mathbf{u}$$

$$p(t_i^* = 1|\mathbf{t}|) = \int_{\mathcal{U}} p(t^* = 1, \mathbf{u}|\mathbf{t}) d\mathbf{u} = \int_{\mathcal{U}} p(t^* = 1|\mathbf{u}|) p(\mathbf{u}|\mathbf{t}) d\mathbf{u}$$

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(d) Now make hard assignments by thresholding the predicted class assignments at $p(t^* = 1|\mathbf{t}) = 0.5$. This can be compared to the true class assignments to give a mean prediction error. Up until now, the true length-scale parameter has been given to the model - you will now attempt to find the parameter through optimisation. Minimise the prediction error with respect to the lengthscale parameter, by performing a simple 1D grid-search between l = 0.01, l = 10 and plotting the error. Compare this to the true value of l used to generate the data - can the length scale be inferred?

Part II - Spatial Data

This section looks at predicting the number of bike thefts in Lewisham borough (Figure 1a) ² over 2015. The dataset given in data.csv contains a lists of x, y locations of $400m^2$ cells and the corresponding number of bike thefts in that location in 2015. The task is to subsample this count data, then infer the field **u** on the original coordinates given this data, and evaluate performance based on predictions of the original data.

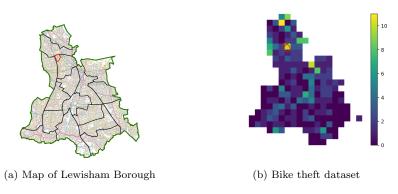


Figure 1

²https://www.lgbce.org.uk/_flysystem/s3/2018-07/lewisham_0.jpg

The likelihood needed here is the Poisson likelihood. The field \mathbf{u} is mapped to \mathbb{R}^+ using the exponential function for positivity, giving $\theta_i = \exp([G\mathbf{u}]_i)$, which is used as the Poisson rate for each data point location. We denote the subsampled count data by the vector \mathbf{c} , and the likelihood of \mathbf{c} given the rate parameters at the subsampled locations θ is shown below:

sampled locations
$$\boldsymbol{\theta}$$
 is shown below:
$$p(\mathbf{c}|\boldsymbol{\theta}) = \prod_{i=1}^{M} f(c_i|\theta_i) = \prod_{i=1}^{M} \frac{e^{-\theta_i}\theta_i^{c_i}}{c_i!}$$

(e) Derive the full Poisson log-likelihood, and use this to complete the function $\log_{poisson_likelihood}$ to return $p(\mathbf{c}|\mathbf{u})$ Generate samples from the posterior $p(\mathbf{u}|\mathbf{c})$ using the pCN method.

The inferred expected counts are given by:

$$\mathbb{E}_{p(c^*|\mathbf{c})}[c^*] = \sum_{k=0}^{\infty} kp(c^* = k|\mathbf{c})$$

We have samples from $p(\mathbf{u}|\mathbf{c})$, so using the predictive distribution of bike theft counts at a test location:

$$p(c^* = k|\mathbf{c}) = \int p(c^* = k, \mathbf{u}|\mathbf{c}) d\mathbf{u} = \int p(c^* = k|\mathbf{u}) p(\mathbf{u}|\mathbf{c}) d\mathbf{u} = \int \frac{1}{|\mathbf{c}|} \int \left(\mathcal{C}_i^* = k |\underline{\mathcal{U}}_i^{(\dagger)} \right) d\mathbf{u} = \int \frac{1}{|\mathbf{c}|} \int \frac{1}{|\mathbf{c}|} \int \frac{1}{|\mathbf{c}|} \int \frac{1}{|\mathbf{c}|} d\mathbf{u} d\mathbf{u} d\mathbf{u} = \int \frac{1}{|\mathbf{c}|} \int \frac{1}{|\mathbf{c}|} \int \frac{1}{|\mathbf{c}|} d\mathbf{u} d\mathbf{u}$$

 $\mathbb{E}_{f(c^*|\theta)}[c^*] = \theta^*$, where c^* is the count of a test location \mathbf{x}^* , and θ^* the respective transformed field value u^* :

$$\mathbb{E}_{f(c^{\star}|\theta)}[c^{\star}] = \theta^{\star}, \text{ where } c^{\star} \text{ is the count of a test location } \mathbf{x}^{\star}, \text{ and } \theta^{\star} \text{ the respective transformed}$$

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$$\mathbb{E}_{p(c^{\star}|\mathbf{c})}[c^{\star}|\mathcal{L}^{\star}] = \mathbb{E}_{p(c^{\star}|\mathbf{c})}[c^{\star}|\mathcal{L}^{\star}] = \mathbb{E}_{p(c^{\star}|\mathbf{c})}[c^{\star}] = \mathbb{E}_{p(c^{\star}|\mathbf{c})}[c^{$$

Use this to infer the expected counts and compare these to the bike theft counts from the briginal dataset. Visualise four plots - the original count data, the subsampled data, the inferred counts, and the error field. Run the model using extreme length-scale parameter values (low and high), and comment on the predicted counts. Suggest a reasonable length-scale value.