Markov Chain Monte Carlo for Inverse Problems

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1 Theory

1.1 Papers

1.1.1 Stuart et al: Inverse Problems: A Bayesian Perspective [3]

Theoretical Background

1. Notation Central equation:

$$y = \mathcal{G}\left(u\right) + \eta$$

with:

- $y \in \mathbb{R}^q$: data
- $u \in \mathbb{R}^n$: IC ("input to mathematical model")
- $\mathcal{G}(\cdot): \mathbb{R}^n \to \mathbb{R}^q$: observation operator
- η : mean zero RV, observational noise (a.s. $\eta \sim \mathcal{N}(0, \mathcal{C})$)

1.1.2 Cotter et al: MCMC for functions [1]

Implementation, MCMC in infinite dimensions

1.1.3 Schneider et al: Earth System Modeling 2.0 [2]

Example for MCMC on ODE

1.2 Small results

1.2.1 Gaussian in infinite dimensions

Wiki-definition of Gaussian measure: uses Lesbesgue-Measure

However, the Lesbesgue-Measure is not defined in an infinite-dimensional space (wiki).

Can still define a measure to be Gaussian if we demand all push-forward measures via a linear functional onto a finite dimensional space to be Gaussian. (What about the star (E^*, L_*) in the wiki-article? Are they dual-spaces?)

How does this fit with the description in [1]? -> Karhunenen-Loéve

The problem is not actually in $\exp(-1/2x^T\mathcal{C}^{-1}x)$. What about $\exp(-1/2\|\mathcal{C}^{-1/2}x\|)$?

What about the terminology in [1]? Absolutely continuous w.r.t a measure for example?

How is the square root of an operator defined? For matrices, there seems to be a freedom in choosing whether A=BB or $A=BB^T$ for $B=A^{1/2}$. The latter definition seems to be more useful when working with Cholesky factorizations (cf. https://math.stackexchange.com/questions/2767873/why-is-the-square-root-of-cholesky-decomposition-equal-to-the-lower-but for example in the wiki-article about the matrix (operator) square root (https://en.wikipedia.org/wiki/Square_root_of_a_matrix): "The Cholesky factorization provides another particular example of square root, which should not be confused with the unique non-negative square root."

1.2.2 Bayes' Formula & Radon-Nikodym Derivative

Bayes' Formula is stated using the Radon-Nikodym Derivative in both [1] and [3]:

$$\frac{\mathrm{d}\mu}{\mathrm{d}\mu_0} \propto \mathrm{L}(u),$$

where L(u) is the likelihood.

Write the measures as $d\mu = \rho(u)du$ and $d\mu_0 = \rho_0(u)du$ with respect to the standard Lesbesgue measure. Then we have

$$\int f(u)\rho(u)du = \int f(u)d\mu(u) = \int f(u)\frac{d\mu(u)}{d\mu_0(u)}d\mu_0 = \int f(u)\frac{d\mu(u)}{d\mu_0(u)}\rho_0(u)du,$$

provided that $d\mu$, $d\mu_0$ and f are nice enough (which they are since we're working with Gaussians). This holds for all test functions f, so it must hold pointwise:

$$\frac{\mathrm{d}\mu(u)}{\mathrm{d}\mu_0(u)} = \frac{\rho(u)}{\rho_o(u)}.$$

Using this we recover the more familiar formulation of Bayes' formula:

$$\frac{\rho(u)}{\rho_o(u)} \propto L(u).$$

1.2.3 Acceptance Probability for Metropolis-Hastings

A Markov process with transition probabilities t(y|x) has a stationary distribution $\pi(x)$.

• The existence of $\pi(x)$ follows from detailed balance:

$$\pi(x)t(y|x) = \pi(y)t(x|y).$$

Detailed balance is sufficient but not necessary for the existence of a stationary distribution.

- Uniqueness of $\pi(x)$ follows from the Ergodicity of the Markov process. For a Markov process be Ergodic it has to:
 - not return to the same state in a fixed interval
 - reach every state from every other state in finite time

The Metropolis-Hastings algorithm constructs transition probabilities t(y|x) such that the two conditions above are satisfied and that $\pi(x) = P(x)$, where P(x) is the distribution we want to sample from.

Rewrite detailed balance as

$$\frac{t(y|x)}{t(x|y)} = \frac{P(y)}{P(x)}.$$

Split up the transition probability into proposal g(y|x) and acceptance a(y,x). Then detailed balance requires

$$\frac{a(y,x)}{a(x,y)} = \frac{P(y)g(x|y)}{P(x)g(y|x)}.$$

Choose

$$a(y,x) = \min\left\{1, \frac{P(y)g(x|y)}{P(x)g(y|x)}\right\}$$

to ensure that detailed balance is always satisfied. Choose g(y|x) such that ergodicity is fulfilled.

If the proposal is symmetric (g(y|x) = g(x|y)), then the acceptance takes the simpler form

$$a(y,x) = \min\left\{1, \frac{P(y)}{P(x)}\right\}. \tag{1}$$

Since the target distribution P(x) only appears as a ratio, normalizing factors can be ignored.

1.2.4 Potential for Bayes'-MCMC when sampling from analytic distributions

How can we use formulations of Metropolis-Hastings-MCMC algorithms designed to sample from posteriors when want to sample from probability distribution with an easy analytical expression?

Algorithms for sampling from a posterior sample from

$$\rho(u) \propto \rho_0(u) \exp(-\Phi(u)),$$

where ρ_0 is the prior and $\exp(-\Phi(u))$ is the likelihood. Normally, we have an efficient way to compute the likelihood.

When we have an efficient way to compute the posterior ρ and we want to sample from it, the potential to do that is:

$$\Phi(u) = \ln(\rho_0(u)) - \ln(\rho(u)),$$

where an additive constant from the normalization was omitted since only potential differences are relevant.

When working with a Gaussian prior $\mathcal{N}(0,\mathcal{C})$, the potential takes the form

$$\Phi(u) = -\ln \rho(u) - \frac{1}{2} \|\mathcal{C}^{-1/2}u\|^2.$$

When inserting this into the acceptance probability for the standard random walk MCMC given in formula (1.2) in [1], the two Gaussian-expressions cancel, as do the logarithm and the exponentiation, leaving the simple acceptance described in 1.

This cancellation does not happen when using the pCN-Acceptance probability. This could explain the poorer performance of pCN when directly sampling a probability distribution.

1.2.5 Acceptance Probabilities for different MCMC Proposers

Start from Bayes' formula and rewrite the likelyhood L(u) as $\exp(-\Phi(u))$ for a positive scalar function Φ called the potential:

$$\frac{\rho(u)}{\rho_o(u)} \propto \exp(\Phi(u)).$$

Assuming our prior to be a Gaussian $(\mu_0 \sim \mathcal{N}(0, \mathcal{C}))$.

Then

$$\rho(u) \propto \exp\left(-\Phi(u) + \frac{1}{2} \left\|C^{-1/2}u\right\|^2\right),$$

since $u^T C^{-1} u = (C^{-1/2} u)^T (C^{-1/2} u) = \langle C^{-1/2} u, C^{-1/2} u \rangle = \|C^{-1/2} u\|^2$, where in the first equality we used C being symmetric.

This is formula (1.2) in [1] and is used in the acceptance probability for the standard random walk (see also Acceptance Probability for Metropolis-Hastings)

 $C^{-1/2}u$ makes problems in infinite dimensions.

Todo: Why exactly is the second term (from the prior) cancelled when doing pCN?

1.2.6 Different formulations of multivariate Gaussians

Is an RV $\xi \sim \mathcal{N}\left(0,C\right)$ distributed the same as $C^{1/2}\xi_0$, with $\xi_0 \sim \mathcal{N}\left(0,\mathcal{I}\right)$? From wikipedia: Affine transformation Y=c+BX for $X\sim \mathcal{N}\left(\mu,\Sigma\right)$ is also a Gaussian $Y\sim \mathcal{N}\left(c+B\mu,B\Sigma B^T\right)$. In our case $X\sim \mathcal{N}\left(0,\mathcal{I}\right)$, so $Y\sim \mathcal{N}\left(0,C^{1/2}\mathcal{I}C^{1/2}^T\right)=\mathcal{N}\left(0,C\right)$, since the covariance matrix is positive definite, which means it's square root is also positive definite and thus symmetric.

On second thought, it also follows straight from the definition:

$$\mathbf{X} \sim \mathcal{N}(\mu, \Sigma) \Leftrightarrow \exists \mu \in \mathbb{R}^k, A \in \mathbb{R}^{k \times l} \text{ s.t. } \mathbf{X} = \mu + A\mathbf{Z} \text{ with } \mathbf{Z}_n \sim \mathcal{N}(0, 1) \text{ i.i.d}$$

where $\Sigma = AA^T$.

2 Implementation

2.1 Framework/Package Structure

The framework is designed to support an easy use case:

```
proposer = StandardRWProposer(beta=0.25, dims=1)
accepter = AnalyticAccepter(my_distribution)
rng = np.random.default_rng(42)
sampler = MCMCSampler(rw_proposer, accepter, rng)
samples = sampler.run(x_0=0, n_samples=1000)
```

There is only one source of randomness, shared among all classes and supplied by the user. This facilitates reproducability.

Tests are done with pytest.

2.1.1 Distributions

A class for implementing probability distributions.

```
class DistributionBase(ABC):
    @abstractmethod
    def sample(self, rng):
        """Return a point sampled from this distribution"""
```

The most important realisation is the GaussianDistribution, used in the proposers.

The design of this class is based on the implementation in muq2. The precision / sqrt_precision is implemented through a Cholesky decomposition, computed in the constructor. This makes applying them pretty fast $(\mathcal{O}(n^2))$.

At the moment the there is one class for both scalar and multivariate Gaussians. This introduces some overhead as it has to work with both float and np.array. Maybe two seperate classes would be better.

2.1.2 Proposers

Propose a new state v based on the current one u.

```
class ProposerBase(ABC):
    @abstractmethod
    def __call__(self, u, rng):
         ...
```

 $1. \ Standard RWP roposer$

Propose a new state as $v = u + \sqrt{2\delta}\xi,$ with either $\xi \sim \mathcal{N}\left(0, \mathcal{I}\right)$ or $\xi \sim \mathcal{N}\left(0, \mathcal{C}\right)$ (see section 4.2 in [1]).

This leads to a well-defined algorithm in finite dimensions. This is not the case when working on functions (as described in section 6.3 in [1])

2. pCNProposer

Propose a new state as

$$v = \sqrt{1 - \beta^2}u + \beta\xi,$$

with
$$\xi \sim \mathcal{N}(0, \mathcal{C})$$
 and $\beta = \frac{8\delta}{(2+\delta)^2} \in [0, 1]$ (see formula (4.8) in [1]).

This approach leads to an improved algorithm (quicker decorrelation in finite dimensions, nicer properties for infinite dimensions) (see sections 6.2 + 6.3 in [1]).

The wikipedia-article on the Cholesky-factorization mentions the usecase of obtaining a correlated sample from an uncorrelated one by the Cholesky-factor. This is not implemented here.

2.1.3 Accepters

Given a current state u and a proposed state v, decide if the new state is accepted or rejected.

For sampling from a distribution P(x), the acceptance probability for a symmetric proposal is $a = \min\{1, \frac{P(v)}{P(u)}\}$ (see 1.2.3)

```
class ProbabilisticAccepter(AccepterBase):
    def __call__(self, u, v, rng):
        """Return True if v is accepted"""
        a = self.accept_probability(u, v)
        return a > rng.random()

    @abstractmethod
    def accept_probability(self, u, v):
        ...
```

1. AnalyticAccepter

Used when there is an analytic expression of the desired distribution.

```
class AnalyticAccepter(ProbabilisticAccepter):
    def accept_probability(self, u, v):
        return self.rho(v) / self.rho(u)
```

2. StandardRWAccepter

Based on formula (1.2) in [1]:

$$a = \min\{1, \exp(I(u) - I(v))\},\$$

with

$$I(u) = \Phi(u) + \frac{1}{2} \left\| \mathcal{C}^{-1/2} u \right\|^2$$

.

See also 1.2.5.

3. pCNAccepter

Works together with the pCNProposer to achieve the simpler expression for the acceptance

$$a = \min\{1, \exp(\Phi(u) - \Phi(v))\}.$$

4. CountedAccepter

Stores and forwards calls to an "actual" accepter. Counts calls and accepts and is used for calculating the acceptance ratio.

2.1.4 Sampler

class MCMCSampler:

The structure of the sampler is quite simple, since it can rely heavily on the functionality provided by the Proposers and Accepters.

```
def __init__(self, proposal, acceptance, rng):
    ...

def run(self, u_0, n_samples, burn_in=1000, sample_interval=200):
    ...

def _step(self, u, rng):
```

2.2 Results

2.2.1 Analytic sampling from a bimodal Gaussian

1. Setup

Attempting to recreate the "Computational Illustration" from [1]. They use, among other algorithms, pCN to sample from a 1-D bimodal Gaussian

$$\rho \propto (\mathcal{N}(3,1) + \mathcal{N}(-3,1)) \mathbb{1}_{[-10,10]}.$$

Since the density estimation framework for a known distribution is not quite clear to me from the paper, I don't expect to perfectly replicate their results.

They use a formulation of the prior based on the Karhunen-Loéve Expansion that doesn't make sense to me in the 1-D setting (how do I sum infinite eigenfunctions of a scalar?).

The potential for density estimation described in section is also not clear to me (maybe for a similar reason? What is u in the density estimate case?).

I ended up using a normal $\mathcal{N}(0,1)$ as a prior and the potential described before, and compared the following samplers:

- ullet (1) StandardRWProposer ($\delta=0.25$) + AnalyticAccepter
- ullet (2) StandardRWProposer ($\delta=0.25$) + StandardRWAccepter
- (3) pCNProposer ($\beta = 0.25$) + pCNAccepter

The code is in analytic.py.

2. Result

All three samplers are able to reproduce the target density 1 2 2.

The autocorrelation decays for all samplers: 4, 5. However, the pCN doens't do nearly as well as expected. This could be the consequence of the awkward formulation of the potential or a bad prior.

A peculiar thing about the decorrelation of the pCN sampling process is that it somehow is tied to the number of samples, compare 6 and 7. Is this a bug or a misunderstanding of the autocorrelation function?

2.2.2 Bayesian inverse problem for $\mathcal{G}(u) = \langle g, u \rangle$

For $\mathcal{G}(u) = \langle g, u \rangle$ the resulting posterior under a Gaussian prior is again a Gaussian. The model equation is

$$y = \mathcal{G}\left(u\right) + \eta$$

with:

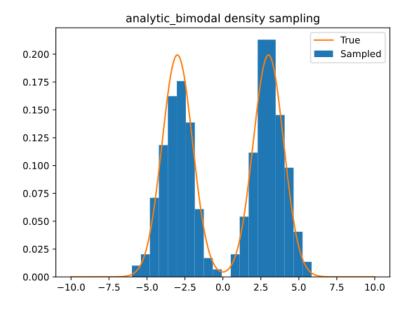


Figure 1: analytic

- $y \in \mathbb{R}$
- $u \in \mathbb{R}^n$
- $\eta \sim \mathcal{N}\left(0, \gamma^2\right)$ for $\gamma \in \mathbb{R}$

A concrete realization with scalar u:

- \bullet u=2
- g = 3
- $\gamma = 0.5$
- y = 6.172
- prior $\mathcal{N}(0, \Sigma_0 = 1)$

leads to a posterior with mean $\mu = \frac{(\Sigma_0 g)y}{\gamma^2 + \langle g, \Sigma_0 g \rangle} \approx 2$, which is what we see when we plot the result 8. The pCN-Sampler with $\beta = 0.25$ had an acceptance rate of 0.567.

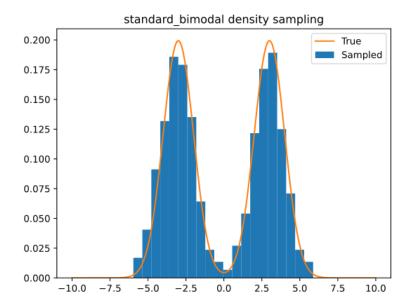


Figure 2: standard rw

For n > 2, the resulting posterior can not be plotted anymore. However, it is still Gaussian with given mean & covariance. Can just compare the analytical values to the sample values. Verify that the error decays like $\frac{1}{\sqrt{N}}$.

2.2.3 Bayesian inverse problem for $G(u) = g(u + \beta u^3)$

Since the observation operator is not linear anymore, the resulting posterior is not Gaussian in general. However, since the dimension of the input u is 1, it can still be plotted.

The concrete realization with:

- g = [3, 1]
- u = 0.5
- $\beta = 0$
- y = [1.672, 0.91]
- $\gamma = 0.5$

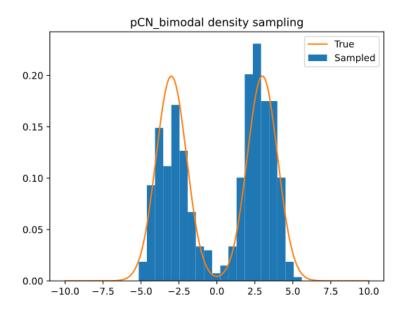


Figure 3: pCN

- $\eta \sim \mathcal{N}\left(0, \gamma^2 I\right)$
- prior $\mathcal{N}(0, \Sigma_0 = 1)$

however leads to a Gaussian thanks to $\beta=0$. The mean is $\mu=\frac{\langle g,y\rangle}{\gamma^2+|g|^2}\approx 0.58$. Plot: 9

The pCN-Sampler with $\beta=0.25$ (different beta) had an acceptance rate of 0.576.

For $\beta \neq 0$, the resulting posterior is not a Gaussian. Still n=1, so it can be plotted. Just numerically normalize the analytical expression of the posterior?

2.2.4 Geophysics example

References

[1] S. L. Cotter, G. O. Roberts, A. M. Stuart, and D. White. MCMC Methods for Functions: Modifying Old Algorithms to Make Them Faster. Statistical Science, 28(3):424–446, August 2013. Publisher: Institute of Mathematical Statistics.

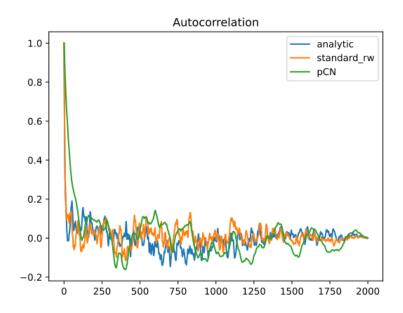


Figure 4: AC of standard normal. All samplers decorrelate quickly

- [2] Tapio Schneider, Shiwei Lan, Andrew Stuart, and João Teixeira. Earth System Modeling 2.0: A Blueprint for Models That Learn From Observations and Targeted High-Resolution Simulations. *Geophysical Research Letters*, 44(24):12,396–12,417, 2017. _eprint: https://agupubs.onlinelibrary.wiley.com/doi/pdf/10.1002/2017GL076101.
- [3] A. M. Stuart. Inverse problems: A Bayesian perspective. *Acta Numerica*, 19:451–559, May 2010.

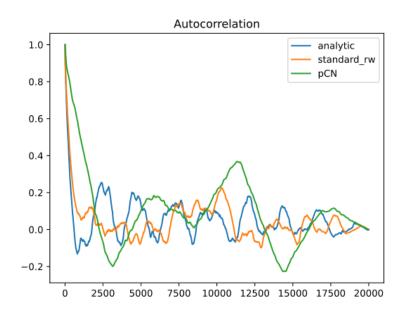


Figure 5: AC of bimodal distribution. pCN takes forever to decorrelate

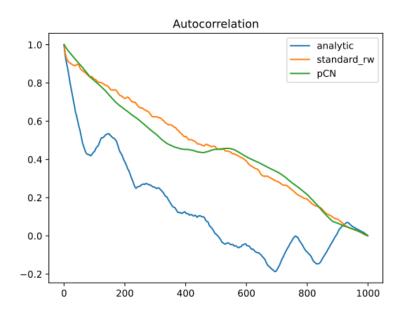


Figure 6: AC of bimodal distribution.

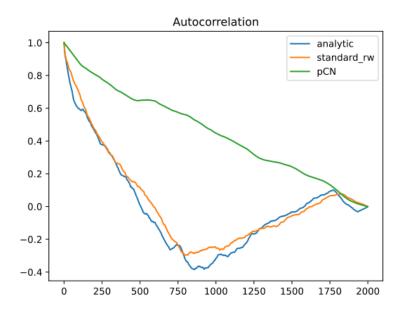


Figure 7: AC of bimodal distribution.

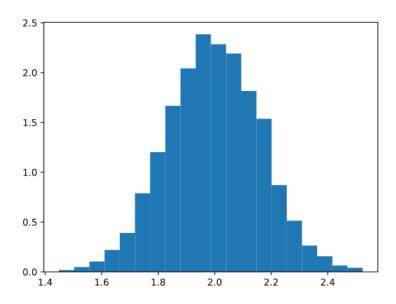


Figure 8: $N = 5000, \mu \approx 2$

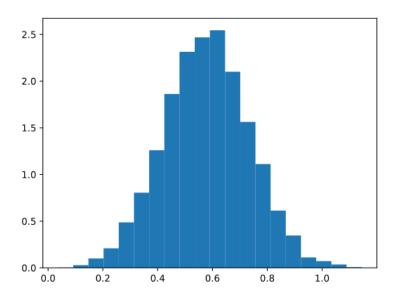


Figure 9: $N = 5000, \mu \approx 0.58$