Markov Chain Monte Carlo

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PSI Numerical Methods 2024

Borrowing heavily from Dan Foreman-Mackey's slides https://speakerdeck.com/dfm/data-analysis-with-mcmc

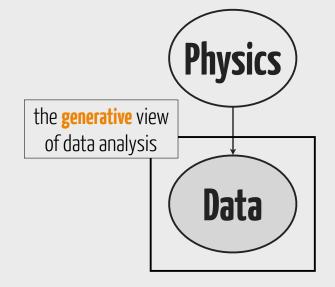
These slides are available at https://github.com/dstndstn/MCMC-talk

data analysis with

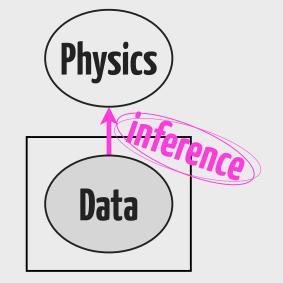
Markov chain Monte Carlo

Dan Foreman-Mackey

CCPP@NYU



The graphical model of my research.



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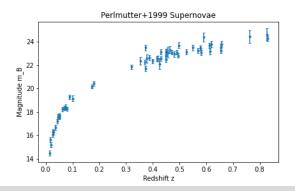
p(data | physics)

likelihood function/generative model

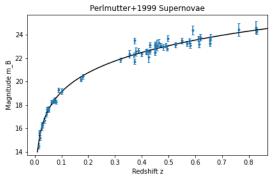


 $p({
m physics}\,|\,{
m data}) \propto p({
m physics})\,p({
m data}\,|\,{
m physics})$ posterior probability

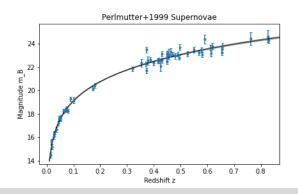
- Perlmutter+1999 (https://arxiv.org/abs/astro-ph/9812133)
- Measured the observed peak brightnesses of a sample of type-1a supernovae (in astronomer "mag" units), and the redshifts ("z") of the supernova host galaxies
- ightharpoonupmag = mag_{intrinsic} + luminosity_distance(z, parameters)



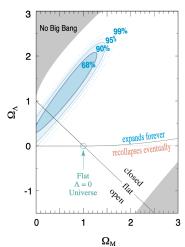
- ► Generative model: $mag_i = mag_{intrinsic} + luminosity_distance(z_i, parameters) + \epsilon_i$
- Probability of a data point given a model ("likelihood"): $p(\text{mag}_i | \text{params}) = \text{Gaussian}(\text{mag}_i | \mu = f(z_i, \text{params}), \sigma_i^2)$
- Probability of whole data set given a model: $p(\{ \max_i \} \mid \Omega_M, \Omega_\Lambda) = \prod_i \mathcal{N}(\max_i \mid \max_{i=1}^n + D_L(z_i, \Omega_M, \Omega_\Lambda), \sigma_i^2)$



- ▶ Use Bayes' theorem to convert data likelihoods into contraints on the model parameters $\theta = \{\Omega_M, \Omega_\Lambda\}$
- $ightharpoonup p(\theta \mid \text{data}) \propto p(\theta) \, p(\text{data} \mid \theta)$
- $p(\Omega_M, \Omega_\Lambda | \{ \text{mag}_i \}) \propto$ $p(\Omega_M, \Omega_\Lambda) \prod_i \mathcal{N}(\text{mag}_i | \text{mag}_{\text{int}} + D_L(z_i, \Omega_M, \Omega_\Lambda), \sigma_i^2)$



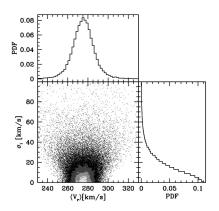
- ► Then they ran MCMC...
- Resulting parameter constraints (blue ellipse):



Why we often need MCMC

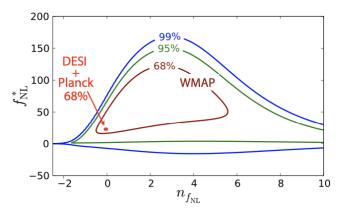
- Real-life models and likelihoods are often complex
- ... so the resulting constraints have complicated distributions (not Gaussians!)
- ... but we can represent them with samplings
- MCMC is used for drawing samples from probability distributions that we can compute numerically but cannot solve analytically

Samplings to represent constraints - examples



- From https://arxiv.org/abs/1910.04899
- With a sampling: Marginalize over a parameter by projecting it out

Samplings to represent constraints - examples



From https://arxiv.org/abs/1611.00036

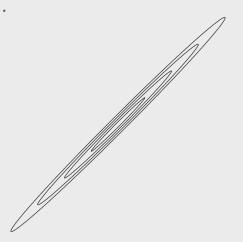
MCMC

draws samples from a probability function

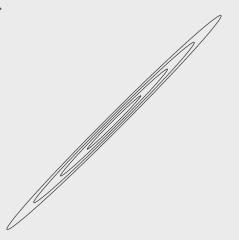
and all you need to be able to do is

evaluate the function

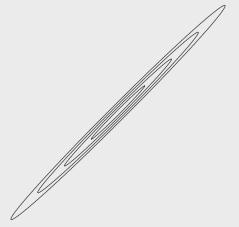
(up to a constant)

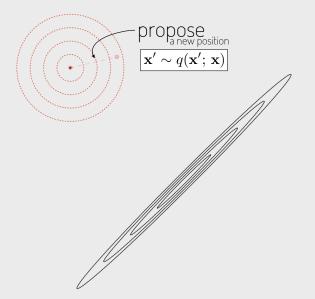


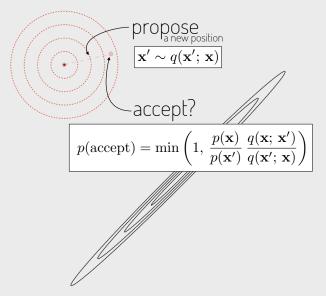
Metropolis-Hastings

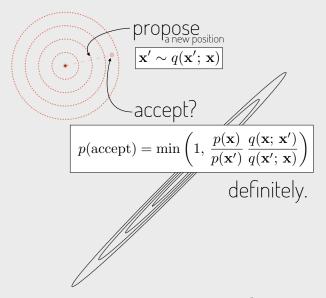


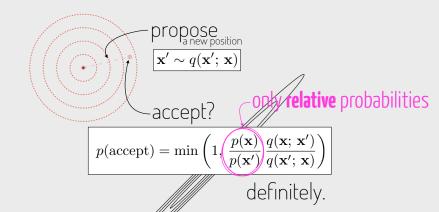


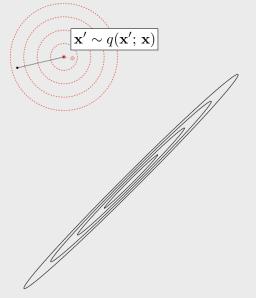


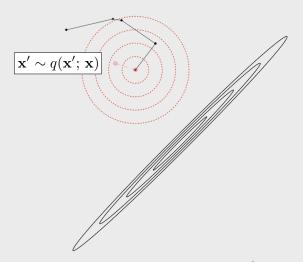


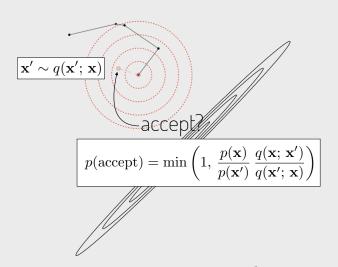


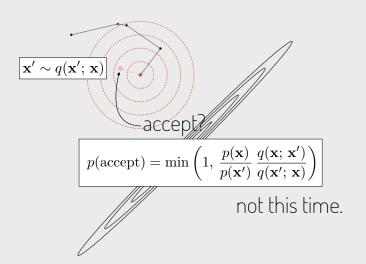


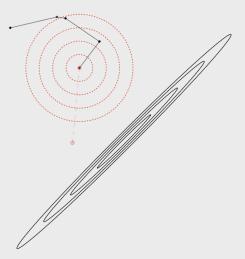


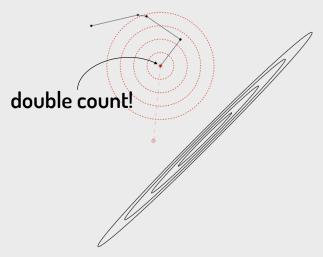


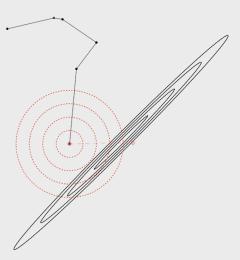














Note

In the previous slides, there is an unfortunate typo! In all cases,

$$p(\text{accept}) = \min(1, \frac{p(x)}{p(x')} \frac{q(x; x')}{q(x'; x)})$$

should actually be

$$p(\text{accept}) = \min(1, \frac{p(x')}{p(x)} \frac{q(x'; x)}{q(x; x')})$$

That is, how good is the new place divided by how good was the old place?

About the name

- Monte Carlo: a reference to the famous Monte Carlo Casino in Monaco, alluding to the randomness used in the algorithm
- Markov Chain: a list of samples, where each one is generated by a process that only looks at the previous one.
- Markov: a 19th-centure Russian mathematician and impressive-moustache-haver with an extensive list of things named after him
- Metropolis—Hastings: lead authors of 1953 and 1970 papers (resp.) giving the algorithm with symmetric and general proposal distributions (resp.)

The Algorithm (1)

```
function mcmc(prob_func, propose_func, initial_pos, nsteps)
    p = initial_pos
    prob = prob_func(p)
    chain = []
    for i in 1:nsteps
        # propose a new position in parameter space
        # ...
        # compute probability at new position
        # ...
        # decide whether to jump to the new position
        if # ...
            # ...
            # . . .
        end
        # save the position
        append! (chain, p)
    end
    return chain
end
```

The Algorithm (2)

```
function mcmc(prob_func, propose_func, initial_pos, nsteps)
    p = initial_pos
    prob = prob_func(p)
    chain = []
    for i in 1:nsteps
        # propose a new position in parameter space
        p_new = propose_func(p)
        # compute probability at new position
        prob_new = prob_func(p_new)
        # decide whether to jump to the new position
        if prob_new / prob > uniform_random()
            p = p_new
            prob = prob_new
        end
        # save the position
        append! (chain, p)
    end
    return chain
end
```

The Algorithm (3)

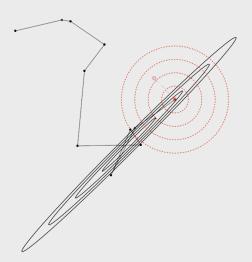
```
function mcmc(logprob_func, propose_func, initial_pos, nsteps)
    p = initial_pos
    logprob = logprob_func(p)
    chain = []
    for i in 1:nsteps
        # propose a new position in parameter space
        p_new = propose_func(p)
        # compute probability at new position
        logprob_new = logprob_func(p_new)
        # decide whether to jump to the new position
        if exp(logprob_new - logprob) > uniform_random()
            p = p_new
            logprob = logprob_new
        end
        # save the position
        append! (chain, p)
    end
    return chain
end
```

The Algorithm (4)

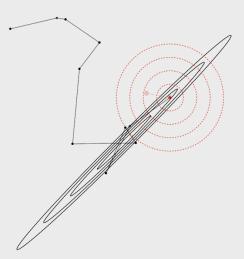
```
function mcmc(logprob_func, propose_func, initial_pos, nsteps)
    p = initial_pos
    logprob = logprob_func(p)
    chain = []
    naccept = 0
    for i in 1:nsteps
        # propose a new position in parameter space
        p_new = propose_func(p)
        # compute probability at new position
        logprob_new = logprob_func(p_new)
        # decide whether to jump to the new position
        if exp(logprob_new - logprob) > uniform_random():
            p = p_new
            logprob = logprob_new
            naccept += 1
        end
        # save the position
        append! (chain, p)
    end
    return chain, naccept/nsteps
```

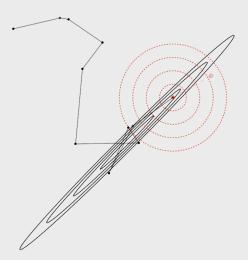
Practicalities

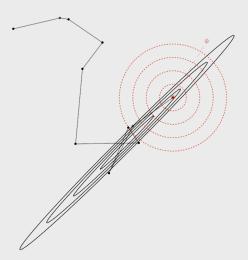
- How do I choose a proposal distribution?
- ► How many steps do I have to take?

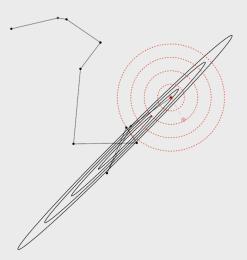


Metropolis-Hastings in the real world

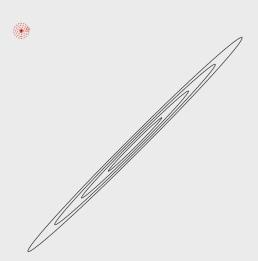


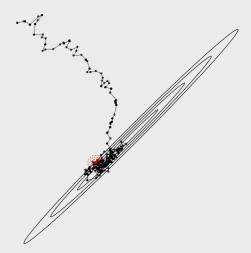




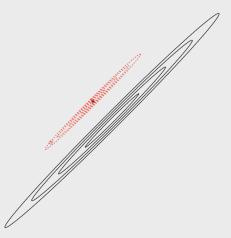


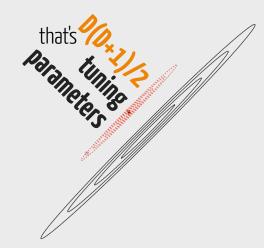


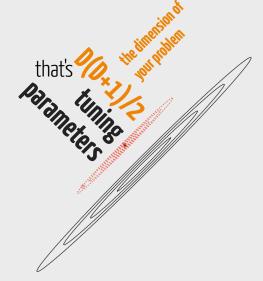


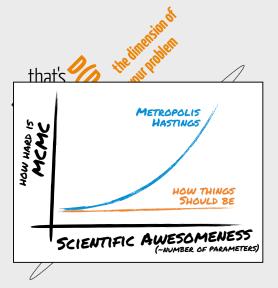






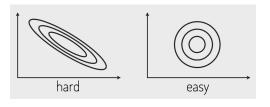






A connection to symmetries

- In Metropolis—Hastings MCMC, the proposal distribution needs tuning parameters, especially as dimensionality increases
- Can be seen as a lack of symmetry in the algorithm—the algorithm is sensitive to the parameterization of the problem
- For example, it's not invariant to an affine transformation
- Next lecture, I'll show you an alternative algorithm that does have affine invariance



How many samples do I need?

- ▶ Burn-in skip the first *N* samples
- Has my chain converged?
- MCMC produces correlated samples, so
 - How correlated are my samples?
 - ightharpoonup Can measure the *autocorrelation time* au
 - ightharpoonup Keep $1/\tau$ of the MCMC samples
 - eg https://github.com/dfm/acor
 - How many uncorrelated samples do I need?
 - ► No easy general answer to this question!
 - "How many can you afford?"

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Conclusions

- MCMC remains an essential tool for probabilistic inference
- For science: lets us contrain model parameters based on data (Bayesian inference)
- Beguilingly simple algorithm, but difficult practicalities
- ▶ MCMC has beautiful theoretical guarantees... as compute time $\to \infty$