#### Markov Chain Monte Carlo

#### Dustin Lang Perimeter Institute for Theoretical Physics

Symmetries Graduate School, 2023-01-23

Borrowing heavily from Dan Foreman-Mackey's slides https://speakerdeck.com/dfm/data-analysis-with-mcmc1

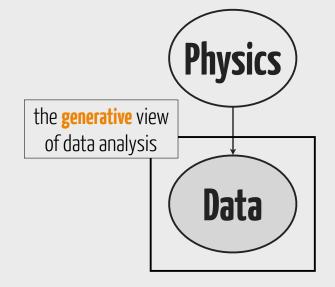
These slides are available at <a href="https://github.com/dstndstn/MCMC-talk">https://github.com/dstndstn/MCMC-talk</a>

data analysis with

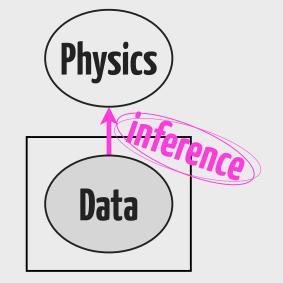
#### Markov chain Monte Carlo

Dan Foreman-Mackey

CCPP@NYU



The graphical model of my research.



The graphical model of my research.

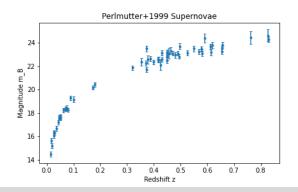
#### p(data | physics)

likelihood function/generative model

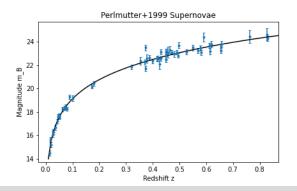


 $p({
m physics}\,|\,{
m data}) \propto p({
m physics})\,p({
m data}\,|\,{
m physics})$  posterior probability

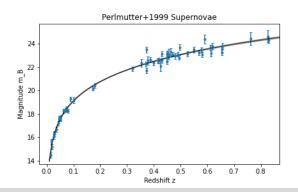
- Perlmutter+1999 (https://arxiv.org/abs/astro-ph/9812133)
- Measured the observed peak brightnesses of a sample of type-1a supernovae (in astronomer "mag" units), and the redshifts ("z") of the supernova host galaxies
- ightharpoonupmag = mag<sub>intrinsic</sub> + luminosity\_distance(z, parameters)



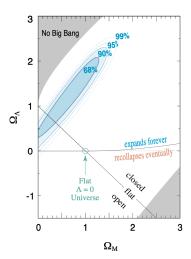
- ► Generative model:  $mag_i = mag_{intrinsic} + luminosity\_distance(z_i, parameters) + \epsilon_i$
- ▶ Probability of data given a model ("likelihood"):  $p(\{\text{mag}_i\} \mid \text{params}) = \prod_i \text{Gaussian}(\text{mag}_i \mid \mu = f(z_i, \text{params}), \sigma_i^2)$
- $p(\text{mag}_i \mid \Omega_M, \Omega_{\Lambda}) = \mathcal{N}(\text{mag}_i \mid \text{mag}_{\text{int}} + D_L(z_i, \Omega_M, \Omega_{\Lambda}), \sigma_i^2)$



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



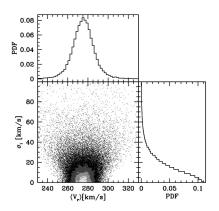
Resulting parameter constraints (blue ellipse):



#### Why we often need MCMC

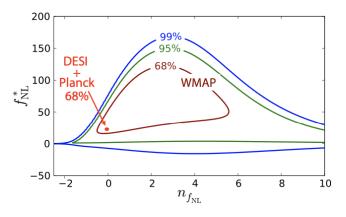
- Real-life models and likelihoods are often complex
- ... so the resulting constraints have complicated distributions (not Gaussians!)
- but we can represent them with samplings
- MCMC is used for drawing samples from probability distributions that we can compute numerically but cannot solve analytically

#### 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

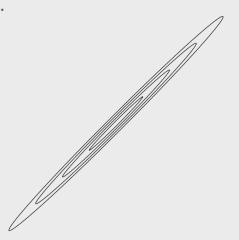
# MCMC

draws samples from a probability function

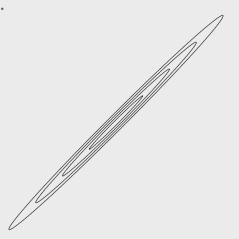
and all you need to be able to do is

## evaluate the function

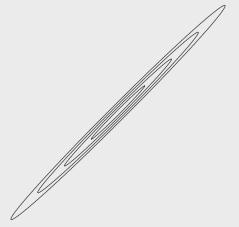
(up to a constant)

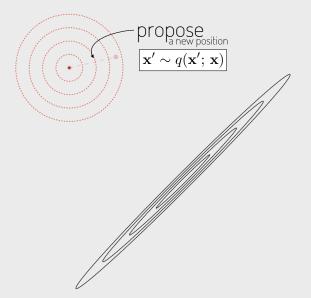


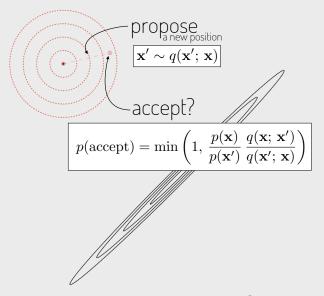
#### **Metropolis-Hastings**

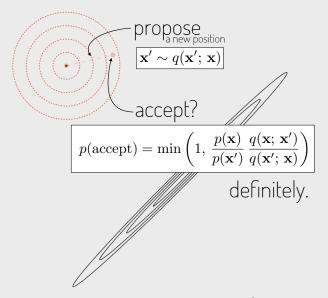


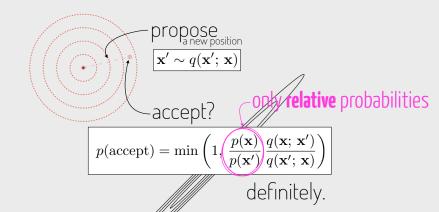


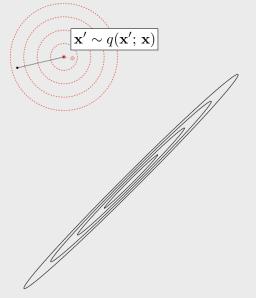


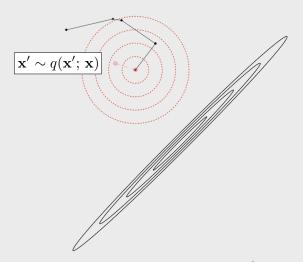


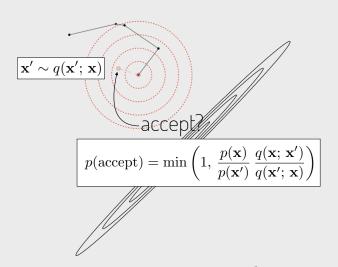


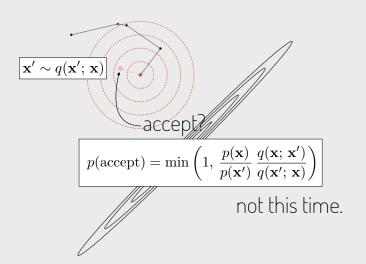


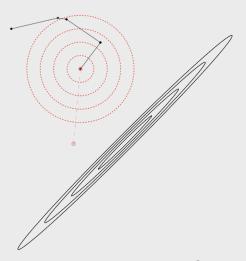


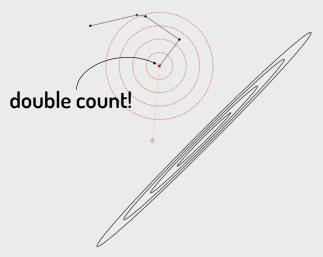


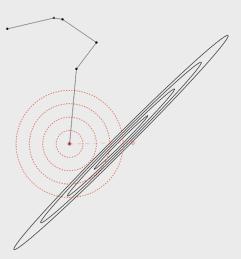














#### About the name

- 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)

```
def mcmc(prob_func, propose_func, initial_pos, nsteps):
     p = initial_pos
     prob = prob_func(p)
     chain = \Pi
     for i in range(nsteps):
         # propose a new position in parameter space
         # . . .
         # compute probability at new position
         # ...
         # decide whether to jump to the new position
         if # ...
             # ...
             # ...
         # save the position
         chain.append(p)
     return chain
```

#### The Algorithm (2)

```
def mcmc(prob_func, propose_func, initial_pos, nsteps):
     p = initial_pos
     prob = prob_func(p)
     chain = \Pi
     for i in range(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
         # save the position
         chain.append(p)
     return chain
```

#### The Algorithm (3)

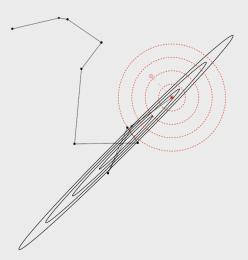
```
def mcmc(logprob_func, propose_func, initial_pos, nsteps):
     p = initial_pos
     logprob = logprob_func(p)
     chain = \Pi
     for i in range(nsteps):
         # propose a new position in parameter space
         p_new = propose_func(p)
         # compute probability at new position
         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
         # save the position
         chain.append(p)
     return chain
```

#### The Algorithm (4)

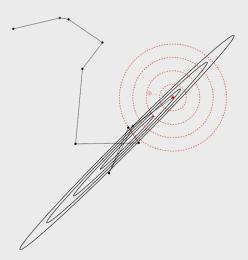
```
def mcmc(logprob_func, propose_func, initial_pos, nsteps):
     p = initial_pos
     logprob = logprob_func(p)
     chain = []
     naccept = 0
     for i in range(nsteps):
         # propose a new position in parameter space
         p_new = propose_func(p)
         # compute probability at new 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
         # save the position
         chain.append(p)
     return chain, naccept/nsteps
```

#### **Practicalities**

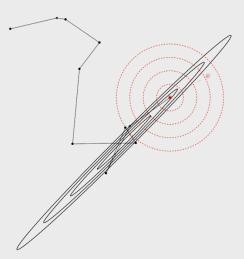
- How do I choose a proposal distribution?
- How many steps do I have to take?

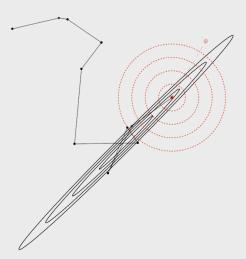


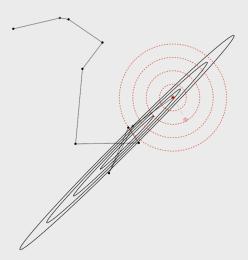
## Metropolis-Hastings in the real world



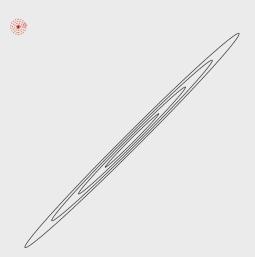
## Metropolis-Hastings in the real world

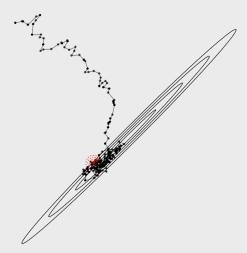




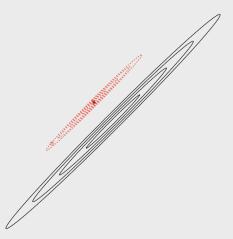


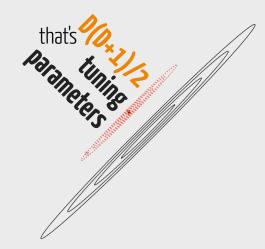


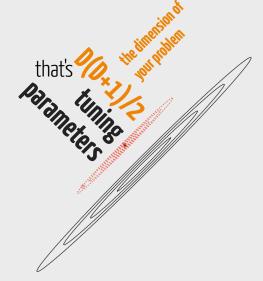


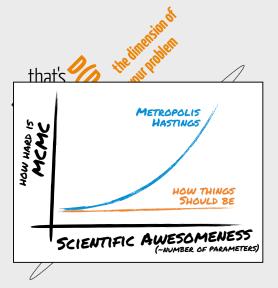






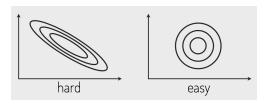






#### A connection to symmetries

- 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?
    - Can measure the autocorrelation time τ
    - ▶ 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?"

#### 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?
    - $\blacktriangleright$  Can measure the *autocorrelation time*  $\tau$
    - 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?"

#### 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?
    - $\blacktriangleright$  Can measure the *autocorrelation time*  $\tau$
    - 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?"

#### as you learned in middle school

$$\int f(\mathbf{x}) p(\mathbf{x}) d\mathbf{x} \approx \frac{1}{N} \sum_{n=1}^{N} f(\mathbf{x}_n)$$
where:  $\mathbf{x}_n \sim p(\mathbf{x})$ 

error: 
$$\delta \propto \frac{1}{\sqrt{N'}}$$
 number of independent camples

#### 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  $\to \infty$

#### This afternoon's tutorial/lab session

- ▶ Bob Room, 3:15–4:30
- Time to play with MCMC yourself!
- We'll use Google CoLab no need to install anything on your computer
- In the Python language