Introduction to Bayesian Inference

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Goals of this lecture

- Understand joint, marginal, and conditional probability distributions
- Understand expectations of functions of a random variable
- Understand how Monte Carlo methods allow us to approximate expectations
- Goal for the subsequent exercise: understand how to implement basic Monte Carlo inference methods



"First I pick a bin, then I pick a single fruit from the bin"

p(red bin) = 2/5p(apple|red) = 2/8



"First I pick a bin, then I pick a single fruit from the bin"

Easy question: what is the probability I pick the red bin?

p(red bin) = 2/5p(apple|red) = 2/8



"First I pick a bin, then I pick a single fruit from the bin"

Easy question: If I first pick the red bin, what is the probability I pick an orange?

p(red bin) = 2/5p(apple|red) = 2/8



"First I pick a bin, then I pick a single fruit from the bin"

Less easy question: What is the overall probability of picking an apple?

p(red bin) = 2/5p(apple|red) = 2/8

"First I pick a bin, then I pick a single fruit from the bin"

Hard question: If I pick an orange, what is the probability that I picked the blue bin?

p(red bin) = 2/5p(apple|red) = 2/8



What is inference?

- The "hard question" requires reasoning backwards in our generative model
- Our generative model specifies these probabilities explicitly:
 - A "marginal" probability *p(bin)*
 - A "conditional" probability p(fruit | bin)
 - A "joint" probability *p(fruit, bin)*
- How can we answer questions about different conditional or marginal probabilities?
 - *p(fruit)*: "what is the overall probability of picking an orange?"
 - p(bin | fruit): "what is the probability I picked the blue bin, given I picked an orange?"

Rules of probability

We just need two basic rules of probability.

• Sum rule:

$$p(\mathbf{y}) = \sum_{\mathbf{x}} p(\mathbf{y}, \mathbf{x}) \quad p(\mathbf{x}) = \sum_{\mathbf{y}} p(\mathbf{y}, \mathbf{x})$$

• Product rule:

$$p(\mathbf{y}, \mathbf{x}) = p(\mathbf{y} | \mathbf{x})p(\mathbf{x}) = p(\mathbf{x} | \mathbf{y})p(\mathbf{y})$$

• These rules define the relationship between *marginal, joint, and conditional* distributions.

Bayes' Rule

Bayes' rule relates two conditional probabilities:



Mini-exercise

 $\sum p(\mathbf{x}|\mathbf{y}) = ???$ X

Use the sum and product rules!

"First I pick a bin, then I pick a single fruit from the bin"

USE THE SUM RULE: What is the overall probability of picking an apple?

p(apple) = p(apple|red)p(red) + p(apple|blue)p(blue) $= 2/8 \times 2/5 + 3/4 \times 3/5$ = 0.55



"First I pick a bin, then I pick a single fruit from the bin"

USE BAYES' RULE: If I pick an orange, what is the probability that I picked the blue bin?

 $p(blue|orange) = \frac{p(orange|blue)p(blue)}{p(orange)}$ $= \frac{1/4 \times 3/5}{6/8 \times 2/5 + 1/4 \times 3/5}$ = 1/3



Continuous probability



A simple continuous example

- Measure the temperature of some water using an inexact thermometer
- The actual water temperature *x* is somewhere near room temperature of 22°; we record an estimate *y*.

 $x \sim \text{Normal}(22, 10)$ $y | x \sim \text{Normal}(x, 1)$

Easy question: what is p(y | x = 25)?

Hard question: what is p(x | y = 25)?

Rules of probability: continuous

• For real-valued *x*, the sum rule becomes an *integral*:

$$p(\mathbf{y}) = \int p(\mathbf{y}, \mathbf{x}) d\mathbf{x}$$

• Bayes' rule:

$$p(\mathbf{x}|\mathbf{y}) = \frac{p(\mathbf{y}|\mathbf{x})p(\mathbf{x})}{p(\mathbf{y})} = \frac{p(\mathbf{y}|\mathbf{x})p(\mathbf{x})}{\int p(\mathbf{y},\mathbf{x})d\mathbf{x}}$$

Integration is harder than addition!

Bayes' rule:
$$p(x|y=25) = \frac{p(x)p(y=25|x)}{p(y=25)}$$

Sum rule, in the denominator:

$$p(y=25) = \int p(x)p(y=25|x)dx$$

In general this integral is intractable, and we can only evaluate up to a normalizing constant

Monte Carlo inference

General problem:



- Our *data* is given by *y*
- Our generative model specifies the prior and likelihood
- We are interested in answering questions about the *posterior* distribution of p(x | y)

General problem:



- Typically we are not trying to compute a probability density function for $p(x \mid y)$ as our end goal
- Instead, we want to compute *expected values* of some function *f(x)* under the posterior distribution

Expectation

• Discrete and continuous:

$$\mathbb{E}[f] = \sum_{x} p(x) f(x)$$
$$\mathbb{E}[f] = \int p(x) f(x) \, \mathrm{d}x.$$

• Conditional on another random variable:

$$\mathbb{E}_x[f|y] = \sum_x p(x|y)f(x)$$

Key Monte Carlo identity

 We can approximate expectations using samples drawn from a distribution p. If we want to compute

$$\mathbb{E}[f] = \int p(x)f(x) \,\mathrm{d}x.$$

we can approximate it with a finite set of points sampled from p(x) using

$$\mathbb{E}[f] \simeq \frac{1}{N} \sum_{n=1}^{N} f(x_n)$$

which becomes exact as N approaches infinity.

How do we draw samples?

- Simple, well-known distributions: samplers exist (for the moment take as given)
- We will look at:
 - 1. Build samplers for complicated distributions out of samplers for simple distributions compositionally
 - 2. Rejection sampling
 - 3. Likelihood weighting
 - 4. Markov chain Monte Carlo

Ancestral sampling from a model

 In our example with estimating the water temperature, suppose we already know how to sample from a normal distribution.

 $x \sim \text{Normal}(22, 10)$

 $y|x \sim Normal(x, 1)$

We can sample *y* by literally simulating from the generative process: we first sample a "true" temperature *x*, and then we sample the observed *y*.

• This draws a sample from the **joint** distribution p(x, y).

Samples from the joint distribution



Conditioning via rejection

- What if we want to sample from a conditional distribution? The simplest form is via rejection.
- Use the ancestral sampling procedure to simulate from the generative process, draw a sample of x and a sample of y. These are drawn together from the joint distribution p(x, y).
- To estimate the posterior $p(x \mid y = 25)$, we say that x is a sample from the posterior if its corresponding value y = 25.
- **Question:** is this a good idea?

Conditioning via rejection



Black bar shows measurement at y = 25. How many of these samples from the joint have y = 25?

- One option is to sidestep sampling from the posterior $p(x \mid y = 3)$ entirely, and draw from some proposal distribution q(x) instead.
- Instead of computing an expectation with respect to p(x|y), we compute an expectation with respect to q(x): $\mathbb{E}_{p(x|y)}[f(x)] = \int f(x)p(x|y)dx$

$$[f(x)] = \int f(x)p(x|y)dx$$
$$= \int f(x)p(x|y)\frac{q(x)}{q(x)}dx$$
$$= \mathbb{E}_{q(x)}\left[f(x)\frac{p(x|y)}{q(x)}\right]$$

- Define an "importance weight" $W(x) = \frac{p(x|y)}{q(x)}$
- Then, with $x_i \sim q(x)$

$$\mathbb{E}_{p(x|y)}[f(x)] = \mathbb{E}_{q(x)}\left[f(x)W(x)\right] \approx \frac{1}{N} \sum_{i=1}^{N} f(x_i)W(x_i)$$

 Expectations now computed using *weighted* samples from *q(x)*, instead of unweighted samples from *p(x|y)*

 Typically, can only evaluate W(x) up to a constant (but this is not a problem):

$$W(x_i) = \frac{p(x_i|y)}{q(x_i)} \qquad \qquad w(x_i) = \frac{p(x_i, y)}{q(x_i)}$$

• Approximation:

$$W(x_i) \approx \frac{w(x_i)}{\sum_{j=1}^N w(x_j)}$$

$$\mathbb{E}_{p(x|y)}[f(x)] \approx \sum_{i=1}^{N} \frac{w(x_i)}{\sum_{j=1}^{N} w(x_j)} f(x_i)$$

- We already have very simple proposal distribution we know how to sample from: the prior p(x).
- The algorithm then resembles the rejection sampling algorithm, except instead of sampling both the latent variables and the observed variables, we only sample the latent variables
- Then, instead of a "hard" rejection step, we use the values of the latent variables and the data to assign "soft" weights to the sampled values.



Draw a sample of x from the prior



What does p(y|x) look like for this sampled x?



What does p(y|x) look like for this sampled x?



What does p(y|x) look like for this sampled x?



Compute p(y|x) for all of our x drawn from the prior



Assign weights (vertical bars) to samples for a representation of the posterior

Conditioning via MCMC

- Problem: Likelihood weighting degrades poorly as the dimension of the latent variables increases, unless we have a very well-chosen proposal distribution q(x).
- **An alternative**: Markov chain Monte Carlo (MCMC) methods draw samples from a target distribution by performing a biased random walk over the space of the latent variables *x*.
- Idea: create a Markov chain such that the sequence of states x_0 , x_1 , x_2 , ... are samples from $p(x \mid y)$



Conditioning via MCMC

- MCMC also uses a proposal distribution, but this proposal distribution makes **local** changes to the latent variables *x*. The proposal *q(x' | x)* defines a conditional distribution over *x'* given a current value *x*.
 - Typical choice: add small amount of Gaussian noise
- We use the proposal and the joint density to define an "acceptance ratio"

$$A(x \to x') = \min\left(1, \frac{p(x', y)q(x|x')}{p(x, y)q(x'|x)}\right)$$

 With probability A we "move" state with the new value x', otherwise we stay at x.



MCMC initialization



Initialize arbitrarily (e.g. with a sample from the prior)

First MCMC step



Propose a local move on x from a transition distribution

1 MCMC iteration



10 MCMC iterations



Continue: propose a local move, and accept or reject. At first, this will look like a stochastic search algorithm!

100 MCMC iterations



Once in a high-density region, it will explore the space

200 MCMC iterations



Once in a high-density region, it will explore the space



Helpful diagnostic: a "trace plot" of the path of the sampled values, as the number of MCMC iterations increases



Histogram of trace plot, overlaid on prior probability density

Now: exercises

- **Part one:** a model much like the model we just looked at, Gaussian data with a latent Gaussian distributed mean
 - A. implement likelihood weighting for this model
 - **B.** this is one of the *very few* continuous models where exact inference is possible. Do the math, and check if your sampler is correct!
- Part two: seven scientists are performing an experiment to estimate the value of a particular physical constant. Most of them find similar results, but a few differ by surprisingly much. Do I trust all these scientists equally? What is the "real" value? Write an MCMC sampler to find out!