

STA3431 (Monte Carlo Methods) Lecture Notes, Fall 2020

by Jeffrey S. Rosenthal, University of Toronto

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Note: I will update these notes regularly (on the course web page). However, they are just rough, point-form notes, with no guarantee of completeness or accuracy. They should in no way be regarded as a substitute for attending the lectures, doing the homework exercises, studying the course material, or reading the reference books.

INTRODUCTION:

- Introduction to course, handout, references, prerequisites, etc.
 - Course web page: probability.ca/sta3431
 - Lectures: Online on Zoom, synchronous, Mondays 10:10–12:00.
 - If not Stat Dept grad student, must REQUEST enrolment (by e-mail); need advanced undergraduate probability/statistics background, plus basic computer programming (including “R”; see e.g. [this page](#)).
 - Conversely, if you already know lots about MCMC etc., then this course might not be right for you since it’s an INTRODUCTION to these topics.
 - How many of you are graduate students in Statistics? Computer Science? Math? Engineering? Physics? Economics? Management? Other?
- Theme of the course: use (pseudo)randomness on a computer to simulate, and hence estimate, important/interesting quantities.
- Example: Suppose we want to estimate $m := \mathbf{E}[Z^4 \cos(Z)]$, where $Z \sim \text{Normal}(0, 1)$.
 - “Classical” Monte Carlo solution: replicate a large number z_1, \dots, z_n of $\text{Normal}(0,1)$ random variables, and let $x_i = z_i^4 \cos(z_i)$.
 - Their mean $\bar{x} \equiv \frac{1}{n} \sum_{i=1}^n x_i$ is an estimate of $\mathbf{E}[X] \equiv \mathbf{E}[Z^4 \cos(Z)]$.
 - R: `Z = rnorm(100); X = Z^4 * cos(Z); mean(X)` [file “RMC”]
 - Unbiased (good) ... but unstable ... but if replace “100” with “1000000” then \bar{x} is consistently close to -1.213 ... good ...
 - [Aside: In this course we will often use R to automatically sample from simple distributions like Normal, Uniform, Exponential, etc. But how does it work? Discussed later!]
 - Can we quantify the variability?

- Well, can estimate standard deviation of \bar{x} by the estimated “standard error” of \bar{x} , which is:

$$se = \sqrt{\text{Var}(\bar{x})} = \sqrt{\text{Var}\left(\frac{1}{n} \sum_{i=1}^n x_i\right)} = \sqrt{\left(\frac{1}{n}\right)^2 (n \text{ var}(x))}$$

$$= n^{-1/2} \sqrt{\text{var}(x)} \approx n^{-1/2} \sqrt{\frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2} =: n^{-1/2} \text{sd}(x).$$

[file “RMCse”]

- Then what is, say, a 95% confidence interval for m ?
- Well, by the Central Limit Theorem (CLT), for large n , we have $\bar{x} \approx N(m, v) \approx N(m, se^2)$.
 - (Strictly speaking, should use “t” distribution, not normal distribution ... but if n large that doesn’t really matter – ignore it for now.)
 - So $\frac{m-\bar{x}}{se} \approx N(0, 1)$.
 - So, $\mathbf{P}(-1.96 < \frac{m-\bar{x}}{se} < 1.96) \approx 0.95$.
 - So, $\mathbf{P}(\bar{x} - 1.96 se < m < \bar{x} + 1.96 se) \approx 0.95$.
 - i.e., approximate 95% confidence interval is [file “RMCci”]

$$(\bar{x} - 1.96 se, \bar{x} + 1.96 se).$$

- Alternatively, could compute expectation as

$$\int_{-\infty}^{\infty} z^4 \cos(z) \frac{e^{-z^2/2}}{\sqrt{2\pi}} dz.$$

Analytic? Numerical? Better? Worse? [file “RMCcomp”: –1.213]

- [Aside: In fact, by considering it as the real part of $\mathbf{E}(Z^4 e^{iZ})$, with extra work this expectation can be computed exactly, to be $-2/\sqrt{e} \doteq -1.213061$. But not for harder examples.]
- What about higher-dimensional examples? (Can’t do numerical integration!) Can we still sample?
- What if the distribution is too complicated to sample from?
 - (MCMC! Metropolis, Gibbs, etc... Soon!)

HISTORICAL EXAMPLE – BUFFON’S NEEDLE:

- Have series of parallel lines ... line spacing w , needle length $\ell \leq w$ (say $\ell = w$) ... what is prob that needle lands touching line? [<https://mste.illinois.edu/activity/buffon/>]

- Let h be the vertical distance from the bottom end to the nearest line above, and θ be the needle's angle counter-clockwise from horizontal.
- Then $h \sim \text{Uniform}[0, w]$ and $\theta \sim \text{Uniform}[0, \pi]$, independent.
- Touches line iff $\ell \sin(\theta) > h$, i.e. $h < \ell \sin(\theta)$.
- So, the probability the needle touches the line is:

$$\begin{aligned} \frac{1}{\pi} \int_0^\pi \frac{1}{w} \int_0^w \mathbf{1}_{h < \ell \sin(\theta)} dh d\theta &= \frac{1}{\pi} \int_0^\pi \frac{1}{w} \ell \sin(\theta) d\theta \\ &= \frac{1}{\pi} \frac{1}{w} \ell [-\cos(\theta)]_{\theta=0}^{\theta=\pi} = \frac{1}{\pi} \frac{1}{w} \ell [-(-1) + (1)] = \frac{2\ell}{w\pi}. \end{aligned}$$

- Hence, by LLN, if throw needle n times, of which it touches a line m times, then for n large, $m/n \approx 2\ell/w\pi$, so $\pi \approx 2n\ell/mw$.
- (e.g. if $\ell = w$, then $\pi \approx 2n/m$)
- [e.g. recuperating English Captain O.C. Fox, 1864: $\ell = 3$, $w = 4$, $n = 530$, $m = 253$, so $\pi \approx 2n\ell/mw \doteq 3.1423$.]
- But for modern simulations, use computer. How to randomise??

PSEUDORANDOM NUMBERS:

- Goal: generate an i.i.d. sequence $U_1, U_2, U_3, \dots \sim \text{Uniform}[0, 1]$.
- One method: LINEAR CONGRUENTIAL GENERATOR (LCG).
 - Choose (large) positive integers m , a , and b .
 - Start with a “seed” value, x_0 . (e.g., the current time in milliseconds)
 - Then, recursively, $x_n = (ax_{n-1} + b) \bmod m$, i.e. $x_n =$ remainder when $ax_{n-1} + b$ is divided by m .
 - So, $0 \leq x_n \leq m - 1$.
 - Then let $U_n = x_n/m$.
 - Then $\{U_n\}$ will “seem” to be approximately i.i.d. $\sim \text{Uniform}[0, 1]$. (file “Rrng”)
- Choice of m , a , and b ? Many issues ...
 - Need m large (so many possible values);
 - Need a large enough that no obvious “pattern” between U_{n-1} and U_n .
 - Need b to avoid short “cycles” of numbers.
 - Want large “period”, i.e. number of iterations before repeat.
 - Many statistical tests, to try to see which choices provide good randomness, avoid correlations, etc. (e.g. “diehard tests”, “dieharder”: www.phy.duke.edu/~rgb/General/dieharder.php)
 - One common “good” choice: $m = 2^{32}$, $a = 69,069$, $b = 23,606,797$.

- Theorem: the LCG has full period (m) if and only if both:
 - (i) $\gcd(b, m) = 1$, and
 - (ii) every “prime or 4” divisor of m also divides $a - 1$.
 - So, if $m = 2^{32}$, then if b odd and $a - 1$ is a multiple of 4 (like above), then the LCG has full period $m = 2^{32} \doteq 4.3 \times 10^9$; good.
- Many other choices, some good, some bad (ASIDE):
 - e.g. C programming language “glibc” uses $m = 2^{32}$, $a = 1, 103, 515, 245$, $b = 12, 345$. Pretty good.
 - “RANDU” used $m = 2^{31}$, $a = 65539 = 2^{16} + 3$, $b = 0$ for many years, around the 1970s. Seemed okay. But then people noticed:

$$x_{n+2} = ax_{n+1} = a^2x_n = (2^{16} + 3)^2x_n = (2^{32} + 6(2^{16}) + 9)x_n$$

$$\equiv (0 + 6(2^{16} + 3) - 9)x_n \pmod{2^{31}} = 6x_{n+1} - 9x_n.$$
 So, $x_{n+2} = 6x_{n+1} - 9x_n \pmod{m}$. Too much serial correlation! Bad!
 - Microsoft Excel pre-2003: had period $< 10^6$, too small ...
 - Excel 2003 instead used a floating-point “version” of LCG ... which sometimes gave negative numbers! Bad!
- These numbers are not “really” random, just “pseudorandom” ...
 - Can cause problems! Will fail certain statistical tests!
 - Some implementations also use external randomness, e.g. temperature of computer’s CPU / entropy of kernel (e.g. Linux’s “urandom”).
 - Or, the randomness of *quantum mechanics*, e.g. www.fourmilab.ch/hotbits (see “myhotbits”).
 - Or, of atmospheric noise (from lightning etc.), e.g. random.org.
 - But mostly, pseudorandom numbers work pretty well ...
- LCG’s are “good enough”. But other generators include (ASIDE):
 - “Multiply-with-Carry”: $x_n = (ax_{n-r} + b_{n-1}) \pmod{m}$ where $b_n = \lfloor (ax_{n-r} + b_{n-1})/m \rfloor$.
 - ‘Kiss’: $y_n = (x_n + J_n + K_n) \pmod{2^{32}}$, where x_n as above, and J_n and K_n are “shift register generators”, given in bit form by $J_{n+1} = (I + L^{15})(I + R^{17})J_n \pmod{2^{32}}$, and $K_{n+1} = (I + L^{13})(I + R^{18})K_n \pmod{2^{31}}$, where L means “shift left” and R means “shift right”.
 - Mersenne Twister: $x_{n+k} = x_{n+s} \oplus (x_n^{(\text{upper})} | x_{n+1}^{(\text{lower})})A$, where $1 \leq s < k$ where $2^{kw-r} - 1$ is Mersenne prime, and A is $w \times w$ (e.g. 32×32) with $(w - 1) \times (w - 1)$ identity in upper-right, and where the matrix multiplication is done bit-wise mod 2.
 - And many others, too. An entire research area!
 - R’s choice? See “?RNGkind”. Default is Mersenne Twister.

- So, just need computer to do simple arithmetic. No problem, right?

LIMITATIONS OF COMPUTER ARITHMETIC:

- Consider the following computations in R:
 - $> 2 + 1 - 2$
 - $> 2^{10} + 1 - 2^{10}$
 - $> 2^{100} + 1 - 2^{100}$
- Why??
- Question for next class: In R, for what values of n does:
 - $> 2^n + 1 - 2^n$
 - give 0 instead of 1?
- (Similarly in many other computer languages too, e.g. C (powertest.c), Java (powertest.java) ... and Python with floating numbers ... but not Python with *integer* variables (powertest.py), because it then does dynamic memory allocation ...)
- Also, overflow/Inf/underflow problems: 2^{10000} , 2^{-10000} , etc.
- So, numerical computations are approximations, with their own errors.
- We'll usually ignore these issues, but you should BE CAREFUL!
- So how to use pseudorandomness?
 - With LCG etc, we can simulate Uniform[0,1] random variables.
 - What about other random variables?

END WEEK #1

SIMULATING OTHER DISTRIBUTIONS:

- Once we have U_1, U_2, \dots i.i.d. \sim Uniform[0,1] (at least approximately), how do we generate other distributions?
- With transformations, using the “change-of-variable” theorem!
- e.g. to make $X \sim$ Uniform[L, R], set:

$$X = (R - L)U_1 + L.$$
- e.g. to make $X \sim$ Bernoulli(p), set:

$$X = \begin{cases} 1, & U_1 \leq p \\ 0, & U_1 > p \end{cases}$$

- e.g. to make $Y \sim$ Binomial(n, p), either set:

$$Y = X_1 + \dots + X_n \text{ where}$$

$$X_i = \begin{cases} 1, & U_i \leq p \\ 0, & U_i > p \end{cases},$$

or set:

$$Y = \max \left\{ j : \sum_{k=0}^{j-1} \binom{n}{k} p^k (1-p)^{n-k} \leq U_1 \right\}$$

(where by convention $\sum_{k=0}^{-1} (\dots) = 0$). (“Inverse CDF method”; see below)

- More generally, to make $\mathbf{P}(Y = x_i) = p_i$ for some $x_1 < x_2 < x_3 < \dots$, where $p_i \geq 0$ and $\sum_i p_i = 1$, set:

$$Y = \max \left\{ x_j ; \sum_{k=1}^{j-1} p_k \leq U_1 \right\}.$$

(discrete version of “Inverse CDF method”)

- e.g. to make $Z \sim \text{Exponential}(1)$, set:

$$Z = -\log(U_1).$$

– Then for $x > 0$, $\mathbf{P}(Z > x) = \mathbf{P}(-\log(U_1) > x) = \mathbf{P}(\log(U_1) < -x) = \mathbf{P}(U_1 < e^{-x}) = e^{-x}$. Then CDF = $1 - e^{-x}$, and density = e^{-x} .

– Then, to make $W \sim \text{Exponential}(\lambda)$, set:

$$W = Z/\lambda = -\log(U_1)/\lambda. \text{ [So that } W \text{ has density } \lambda e^{-\lambda x} \text{ for } x > 0.]$$

- Suppose we want X to have density $6x^5 \mathbf{1}_{0 < x < 1}$.

– Let $X = U_1^{1/6}$.

– Then for $0 < x < 1$, $\mathbf{P}(X \leq x) = \mathbf{P}(U_1^{1/6} \leq x) = \mathbf{P}(U_1 \leq x^6) = x^6$.

– Hence, $f_X(x) = \frac{d}{dx} [\mathbf{P}(X \leq x)] = \frac{d}{dx} x^6 = 6x^5$ for $0 < x < 1$.

– More generally, for $r > 1$, if $X = U_1^{1/r}$, then $f_X(x) = r x^{r-1}$ for $0 < x < 1$. [CHECK!]

- What about normal dist.? Fact: If

$$X = \sqrt{2 \log(1/U_1)} \cos(2\pi U_2),$$

$$Y = \sqrt{2 \log(1/U_1)} \sin(2\pi U_2),$$

then $X, Y \sim N(0, 1)$, and X and Y are independent! [“Box-Muller transformation”: Ann Math Stat 1958, 29, 610-611]

– Proof (Aside): By multidimensional change-of-variable theorem, if

$$(x, y) = h(u_1, u_2) \text{ and } (u_1, u_2) = h^{-1}(x, y), \text{ then}$$

$$f_{X,Y}(x, y) = f_{U_1, U_2}(h^{-1}(x, y)) / |J(h^{-1}(x, y))|.$$

Here $f_{U_1, U_2}(u_1, u_2) = 1$ for $0 < u_1, u_2 < 1$ (otherwise 0), and

$$\begin{aligned} J(u_1, u_2) &= \det \begin{pmatrix} \frac{\partial x}{\partial u_1} & \frac{\partial x}{\partial u_2} \\ \frac{\partial y}{\partial u_1} & \frac{\partial y}{\partial u_2} \end{pmatrix} \\ &= \det \begin{pmatrix} -\cos(2\pi u_2) / u_1 \sqrt{2 \log(1/u_1)} & -2\pi \sin(2\pi u_2) \sqrt{2 \log(1/u_1)} \\ -\sin(2\pi u_2) / u_1 \sqrt{2 \log(1/u_1)} & 2\pi \cos(2\pi u_2) \sqrt{2 \log(1/u_1)} \end{pmatrix} \end{aligned}$$

$$= -2\pi / u_1.$$

But $u_1 = e^{-(x^2+y^2)/2}$, so density of (X, Y) is

$$\begin{aligned} f_{X,Y}(x, y) &= 1/|J(h^{-1}(x, y))| = 1/|-2\pi / e^{-(x^2+y^2)/2}| = e^{-(x^2+y^2)/2}/2\pi \\ &= \left(\frac{1}{\sqrt{2\pi}}e^{-x^2/2}\right)\left(\frac{1}{\sqrt{2\pi}}e^{-y^2/2}\right), \end{aligned}$$

i.e. $X \sim N(0, 1)$ and $Y \sim N(0, 1)$ are independent.

- Most general approach: the general “INVERSE CDF METHOD”:
 - Suppose want $\mathbf{P}(X \leq x) = F(x)$. (“CDF”)
 - For $0 < t < 1$, set $F^{-1}(t) = \min\{x; F(x) \geq t\}$. (“inverse CDF”)
 - Then set $X = F^{-1}(U_1)$.
 - Then $X \leq x$ if and only if $U_1 \leq F(x)$. [Subtle; see e.g. Rosenthal, *A First Look at Rigorous Probability Theory*, 2nd ed., Lemma 7.1.2.]
 - So, $\mathbf{P}(X \leq x) = \mathbf{P}(U_1 \leq F(x)) = F(x)$.
 - Very general, but computing $F^{-1}(t)$ is often very difficult ...
- Overall, generating (pseudo)random numbers for most “standard” one-dimensional distributions is mostly pretty easy or well-known ...
 - So, can get Monte Carlo estimates of expectations involving standard one-dimensional distributions, e.g. $\mathbf{E}[Z^4 \cos(Z)]$ where $Z \sim \text{Normal}(0, 1)$.
- What about other Monte Carlo estimates?

MONTE CARLO INTEGRATION:

- How to compute an integral with Monte Carlo?
 - Re-write it as an expectation!
- EXAMPLE: Want to compute $\int_0^1 \int_0^1 g(x, y) dx dy$.
 - Regard this as $\mathbf{E}[g(X, Y)]$, where X, Y i.i.d. $\sim \text{Uniform}[0, 1]$.
 - Then, similar to before, estimate $\mathbf{E}[g(X, Y)]$ by $\frac{1}{M} \sum_{i=1}^M g(x_i, y_i)$, where $x_i \sim \text{Uniform}[0, 1]$ and $y_i \sim \text{Uniform}[0, 1]$ (all independent).
 - e.g. $g(x, y) = \cos(\sqrt{xy})$. (file “RMCint”) Easy!
 - Get about 0.88 ± 0.003 ... Mathematica gives 0.879544.
- e.g. estimate $I = \int_0^5 \int_0^4 g(x, y) dy dx$, where $g(x, y) = \cos(\sqrt{xy})$.
 - Here

$$\int_0^5 \int_0^4 g(x, y) dy dx = \int_0^5 \int_0^4 5 \cdot 4 \cdot g(x, y) (1/4) dy (1/5) dx = \mathbf{E}[5 \cdot 4 \cdot g(X, Y)],$$
 where $X \sim \text{Uniform}[0, 5]$ and $Y \sim \text{Uniform}[0, 4]$.
 - So, let $X_i \sim \text{Uniform}[0, 5]$, and $Y_i \sim \text{Uniform}[0, 4]$ (all independent).

- Estimate I by $\frac{1}{M} \sum_{i=1}^M (5 \cdot 4 \cdot g(X_i, Y_i))$. (file “RMCint2”)
- Standard error: $se = M^{-1/2} sd(5 \cdot 4 \cdot g(X_1, Y_1), \dots, 5 \cdot 4 \cdot g(X_M, Y_M))$.
- With $M = 10^6$, get about $-4.11 \pm 0.01 \dots$ Mathematica gives -4.11692 .
- e.g. estimate $\int_0^1 \int_0^\infty h(x, y) dy dx$, where $h(x, y) = e^{-y^2} \cos(\sqrt{xy})$.
 - (Can’t use “Uniform” expectations.)
 - Instead, write this as, say, $\int_0^1 \int_0^\infty (e^y h(x, y)) e^{-y} dy dx$.
 - This is the same as $\mathbf{E}[e^Y h(X, Y)]$, where $X \sim \text{Uniform}[0, 1]$ and $Y \sim \text{Exponential}(1)$ are independent.
 - So, estimate it by $\frac{1}{M} \sum_{i=1}^M e^{Y_i} h(X_i, Y_i)$, where $X_i \sim \text{Uniform}[0, 1]$ and $Y_i \sim \text{Exponential}(1)$ (i.i.d.). (file “RMCint3”)
 - With $M = 10^6$ get about $0.767 \pm 0.0004 \dots$ Small error!
 - Mathematica: 0.767211.
- Alternatively, could write this as $\int_0^1 \int_0^\infty (\frac{1}{5} e^{5y} h(x, y)) (5 e^{-5y}) dy dx = \mathbf{E}[\frac{1}{5} e^{5Y} h(X, Y)]$ where $X \sim \text{Uniform}[0, 1]$ and $Y \sim \text{Exponential}(5)$ (indep.).
 - Then, estimate it by $\frac{1}{M} \sum_{i=1}^M \frac{1}{5} e^{5y_i} h(x_i, y_i)$, where $x_i \sim \text{Uniform}[0, 1]$ and $y_i \sim \text{Exponential}(5)$ (i.i.d.).
 - With $M = 10^6$, get about $0.767 \pm 0.0016 \dots$ larger standard error \dots (file “RMCint4”).
 - If replace 5 by 1/5, get about $0.767 \pm 0.0015 \dots$ about the same.
- So which choice is best?
 - Whichever one minimises the standard error!
 - ($\lambda \approx 1.5$, $se \approx 0.00025$?)
- In general, to evaluate $I \equiv \int s(y) dy$, could write it as $I = \int \frac{s(x)}{f(x)} f(x) dx$, where f is easily sampled from, with $f(x) > 0$ whenever $s(x) > 0$.
 - Then $I = \mathbf{E}\left(\frac{s(X)}{f(X)}\right)$, where X has density f .
 - (“Importance Sampling”)
 - So, $I \approx \frac{1}{M} \sum_{i=1}^M \frac{s(x_i)}{f(x_i)}$ where $x_i \sim f$.
 - Can then do classical (iid) Monte Carlo integration, and also get standard errors, confidence intervals, etc.
 - Good if it’s easier to sample from f , and/or if the function $\frac{s(x)}{f(x)}$ is less variable than h itself.
- In general, best to make $\frac{s(x)}{f(x)}$ approximately constant if possible.
 - e.g. extreme case: if $I = \int_0^\infty e^{-3x} dx$, then $I = \int_0^\infty (1/3)(3e^{-3x}) dx = \mathbf{E}[1/3]$ where $X \sim \text{Exponential}(3)$, so $I = 1/3$ (error = 0, no MC needed). [Here $s(x) = e^{-3x}$, and $f(x) = 3e^{-3x}$.]

UNNORMALISED DENSITIES:

- Suppose now that $\pi(y) = c g(y)$, where we know g but don't know c or π . (“Unnormalised density”, e.g. Bayesian posterior.)
 - Obviously, $c = \frac{1}{\int g(y) dy}$, but this might be hard to compute.
 - Still, $I = \int h(x) \pi(x) dx = \int h(x) c g(x) dx = \frac{\int h(x) g(x) dx}{\int g(x) dx}$.
 - Here $\int h(x) g(x) dx = \int (h(x) g(x) / f(x)) f(x) dx = \mathbf{E}[h(X) g(X) / f(X)]$ where $X \sim f$.
 - So, $\int h(x) g(x) dx \approx \frac{1}{M} \sum_{i=1}^M (h(x_i) g(x_i) / f(x_i))$ if $\{x_i\} \sim f$ (i.i.d.).
 - Similarly, $\int g(x) dx \approx \frac{1}{M} \sum_{i=1}^M (g(x_i) / f(x_i))$ if $\{x_i\} \sim f$ (i.i.d.).
 - So, must $I \approx \frac{\sum_{i=1}^M (h(x_i) g(x_i) / f(x_i))}{\sum_{i=1}^M (g(x_i) / f(x_i))}$.
 - (“Importance Sampling”: weighted average)
 - (Because we are taking ratios of (unbiased) estimates, the resulting estimate is not unbiased, and its standard errors are less clear. But it is still consistent as $M \rightarrow \infty$.)

END WEEK #2

- Example: compute $I \equiv \mathbf{E}(Y^2)$ where Y has density $c y^3 \sin(y^4) \cos(y^5) \mathbf{1}_{0 < y < 1}$, where $c > 0$ is unknown (and hard to compute).
 - Here $g(y) = y^3 \sin(y^4) \cos(y^5) \mathbf{1}_{0 < y < 1}$, and $h(y) = y^2$.
 - Let $f(y) = 6 y^5 \mathbf{1}_{0 < y < 1}$.
 - [Recall: if $U \sim \text{Uniform}[0, 1]$, and if $X = U^{1/6}$, then $X \sim f$.]
 - Then $I \approx \frac{\sum_{i=1}^M (h(x_i) g(x_i) / f(x_i))}{\sum_{i=1}^M (g(x_i) / f(x_i))} = \frac{\sum_{i=1}^M (\sin(x_i^4) \cos(x_i^5))}{\sum_{i=1}^M (\sin(x_i^4) \cos(x_i^5) / x_i^2)}$, where $\{x_i\}$ are i.i.d. $\sim f$. (file “Rimp1” ... get about 0.766 ...)
 - Or, let $f(y) = 4 y^3 \mathbf{1}_{0 < y < 1}$. [Then if $U \sim \text{Uniform}[0, 1]$, then $U^{1/4} \sim f$.]
 - Then $I \approx \frac{\sum_{i=1}^M (h(x_i) g(x_i) / f(x_i))}{\sum_{i=1}^M (g(x_i) / f(x_i))} = \frac{\sum_{i=1}^M (\sin(x_i^4) \cos(x_i^5) x_i^2)}{\sum_{i=1}^M (\sin(x_i^4) \cos(x_i^5))}$. (file “Rimp2”)
 - Numerical integration: 0.7661155 (file “Rimp3”).
- With importance sampling, is it important to use the same samples $\{x_i\}$ in both numerator and denominator?
 - What if independent samples are used instead?
 - Let's try it! (file “Rimpind”)
 - Both ways work, but usually(?) the same samples work better.
 - Overall, good to use same sample $\{x_i\}$ for both numerator and denominator: easier computationally, and leads to smaller variance.

- What other methods are available to iid sample from π ?

REJECTION SAMPLER:

- Assume $\pi(x) = c g(x)$, with π and c unknown, g known but hard to sample from.
- Want to sample $X \sim \pi$. (Then if $X_1, X_2, \dots, X_M \sim \pi$ iid, then can estimate $\mathbf{E}_\pi(h)$ by $\frac{1}{M} \sum_{i=1}^M h(X_i)$, etc.)
- Find some other, easily-sampled density f , and known $K > 0$, such that $K f(x) \geq g(x)$ for all x . (i.e., $K f(x) \geq \pi(x) / c$, i.e. $c K f(x) \geq \pi(x)$)
- Sample $X \sim f$, and $U \sim \text{Uniform}[0, 1]$ (indep.).
 - If $U \leq \frac{g(X)}{Kf(X)}$, then accept X (as a draw from π).
 - Otherwise, reject X and start over again.
- Does this algorithm give valid samples?
 - Well, conditional on accepting, we have for any $y \in \mathbf{R}$ that

$$\mathbf{P}(X \leq y \mid U \leq \frac{g(X)}{Kf(X)}) = \frac{\mathbf{P}(X \leq y, U \leq \frac{g(X)}{Kf(X)})}{\mathbf{P}(U \leq \frac{g(X)}{Kf(X)})} = ?$$

- Since $0 \leq \frac{g(x)}{Kf(x)} \leq 1$, therefore $\mathbf{P}(U \leq \frac{g(X)}{Kf(X)} \mid X = x) = \frac{g(x)}{Kf(x)}$.
- Hence, by the double expectation formula, $\mathbf{P}(U \leq \frac{g(X)}{Kf(X)}) = \mathbf{E}[\mathbf{P}(U \leq \frac{g(X)}{Kf(X)} \mid X)] = \mathbf{E}[\frac{g(X)}{Kf(X)}] = \int_{-\infty}^{\infty} \frac{g(x)}{Kf(x)} f(x) dx = \frac{1}{K} \int_{-\infty}^{\infty} g(x) dx$.
- Similarly, for any $y \in \mathbf{R}$, $\mathbf{P}(X \leq y, U \leq \frac{g(X)}{Kf(X)}) = \mathbf{E}[\mathbf{1}_{X \leq y} \mathbf{1}_{U \leq \frac{g(X)}{Kf(X)}}] = \mathbf{E}[\mathbf{1}_{X \leq y} \mathbf{P}(U \leq \frac{g(X)}{Kf(X)} \mid X)] = \mathbf{E}[\mathbf{1}_{X \leq y} \frac{g(X)}{Kf(X)}] = \int_{-\infty}^y \frac{g(x)}{Kf(x)} f(x) dx = \frac{1}{K} \int_{-\infty}^y g(x) dx$.
- So,

$$\mathbf{P}(X \leq y \mid U \leq \frac{g(X)}{Kf(X)}) = \frac{\frac{1}{K} \int_{-\infty}^y g(x) dx}{\frac{1}{K} \int_{-\infty}^{\infty} g(x) dx} = \int_{-\infty}^y \pi(x) dx .$$

- So, conditional on accepting, $X \sim \pi$. Good! iid!
- However, prob. of accepting may be very small.
 - If so, then get very few samples – bad.
- EXAMPLE: $\pi = N(0, 1)$, i.e. $g(x) = \pi(x) = (2\pi)^{-1/2} \exp(-x^2/2)$.
 - Want: $\mathbf{E}_\pi(X^4)$, i.e. $h(x) = x^4$. (Should be 3.)
 - Let f be double-exponential (Laplace) distribution, i.e. $f(x) = \frac{1}{2} e^{-|x|}$.
- If $K = 8$, then:

- For $|x| \leq 2$, $Kf(x) = 8\frac{1}{2} \exp(-|x|) \geq 8\frac{1}{2} \exp(-2) \geq (2\pi)^{-1/2} \geq \pi(x) = g(x)$.
- For $|x| \geq 2$, $Kf(x) = 8\frac{1}{2} \exp(-|x|) \geq 8\frac{1}{2} \exp(-x^2/2) \geq (2\pi)^{-1/2} \exp(-x^2/2) = \pi(x) = g(x)$.
- See graph: file “Rrejgraph”.
- So, can apply rejection sampler with this f and K , to get samples, estimate of $\mathbf{E}[X]$, estimate of $\mathbf{E}[h(X)]$, estimate of $\mathbf{P}[X < -1]$, etc.
 - Try it: file “Rrej”
- For Rejection Sampler, $P(\text{accept}) = \mathbf{E}[P(\text{accept}|X)] = \mathbf{E}\left[\frac{g(X)}{Kf(X)}\right] = \int \frac{g(x)}{Kf(x)} f(x) dx = \frac{1}{K} \int g(x) dx = \frac{1}{cK}$. (Only depends on K , not f .)
 - So, in M attempts, get about M/cK iid samples.
 - (“Rrej” example: $c = 1$, $K = 8$, $M = 10,000$, so get about $M/8 = 1250$ samples.)
 - Since c fixed, try to minimise K .
 - Extreme case: $f(x) = \pi(x)$, so $g(x) = \pi(x)/c = f(x)/c$, and can take $K = 1/c$, whence $P(\text{accept}) = 1$, iid sampling: optimal.
- Note: these algorithms all work in discrete case too.
 - Can let π , f , etc. be “probability functions”, i.e. probability densities with respect to counting measure.
 - Then the algorithms proceed exactly as before.

AUXILIARY VARIABLE APPROACH:

- (related: “slice sampler”)
- Suppose $\pi(x) = c g(x)$, and (X, Y) chosen uniformly under graph of g .
 - i.e., $(X, Y) \sim \text{Uniform}\{(x, y) \in \mathbf{R}^2 : 0 \leq y \leq g(x)\}$.
 - Then $X \sim \pi$, i.e. we have sampled from π .
 - Why? Well, for $a < b$,

$$\mathbf{P}(a < X < b) = \frac{\text{area with } a < x < b}{\text{total area}} = \frac{\int_a^b g(x) dx}{\int_{-\infty}^{\infty} g(x) dx} = \int_a^b \pi(x) dx.$$
 - So, if repeat, get i.i.d. samples from π , can estimate $\mathbf{E}_\pi(h)$ etc.
- Auxiliary Variable rejection sampler:
 - If support of g contained in $[L, R]$, and $|g(x)| \leq K$, then can first sample $(X, Y) \sim \text{Uniform}([L, R] \times [0, K])$, then reject if $Y > g(X)$, otherwise accept as sample with $(X, Y) \sim \text{Uniform}\{(x, y) : 0 \leq y \leq g(x)\}$, hence $X \sim \pi$.
- Example: $g(y) = y^3 \sin(y^4) \cos(y^5) \mathbf{1}_{0 < y < 1}$.

- Then $L = 0, R = 1, K = 1$.
- So, sample $X, Y \sim \text{Uniform}[0, 1]$, then keep X iff $Y \leq g(X)$.
- If $h(y) = y^2$, could compute e.g. $\mathbf{E}_\pi(h)$ as the mean of the squares of the accepted samples. (file “Raux”)

DIGRESSION – QUEUEING THEORY:

- Consider a long line (queue) of customers.
 - Let $Q(t)$ = number of people in queue at time $t \geq 0$.
- Suppose service times $\sim \text{Exponential}(\mu)$ [mean $1/\mu$], and interarrival times $\sim \text{Exponential}(\lambda)$ (“M/M/1 queue”), so $\{Q(t)\}$ Markovian. Then well known [e.g. STA447/2006]:
 - If $\mu \leq \lambda$, then $Q(t) \rightarrow \infty$ as $t \rightarrow \infty$.
 - If $\mu > \lambda$, then $Q(t)$ converges in distribution as $t \rightarrow \infty$:
 - $\mathbf{P}(Q(t) = i) \rightarrow (1 - \frac{\lambda}{\mu})(\frac{\lambda}{\mu})^i$, for $i = 0, 1, 2, \dots$
 - Easy! (e.g. $\mu = 3, \lambda = 2, t = 1000$) [file “Rqueue”]
- Now suppose instead that service times $\sim \text{Uniform}[0, 1]$, and interarrival times have distribution of $|Z|$ where $Z \sim \text{Normal}(0, 1)$. Limits not easily computed. Now what?
 - Simulate it! [file “Rqueue2”]
- Or, to make the means the same as the first example, suppose service times $\sim \text{Uniform}[0, 2/3]$, and interarrival times have distribution of $Z^2/2$ where $Z \sim \text{Normal}(0, 1)$. Now what? [file “Rqueue3”]

END WEEK #3

DIGRESSION – MONTE CARLO IN FINANCE:

- X_t = stock price at time t
- Assume that $X_0 = a > 0$, and $dX_t = bX_t dt + \sigma X_t dB_t$, where $\{B_t\}$ is Brownian motion. (“diffusion”)
 - i.e., for small $h > 0$,

$$(X_{t+h} - X_t | X_t) \approx bX_t(t+h-t) + \sigma X_t(B_{t+h} - B_t) \sim bX_t(t+h-t) + \sigma X_t N(0, h),$$

so

$$(X_{t+h} | X_t) \sim N(X_t + bX_t h, \sigma^2(X_t)^2 h). \quad (*)$$

- A “European call option” is the option to purchase one share of the stock at a fixed time $T > 0$ for a fixed price $q > 0$.

- Question: what is a fair price for this option?
 - At time T , its value is $\max(0, X_T - q)$.
 - So, at time 0, its value is $e^{-rT} \max(0, X_T - q)$, where r is the “risk-free interest rate”.
 - But at time 0, X_T is unknown! So, what is fair price??
- FACT: the fair price is equal to $\mathbf{E}(e^{-rT} \max(0, X_T - q))$, but only after replacing b by r .
 - (Proof: transform to risk-neutral martingale measure ...)
 - Intuition: if b very large, might as well just buy stock itself.
- If σ and r constant, then there’s a formula (“Black-Scholes eqn”) for this price, in terms of $\Phi = \text{cdf of } N(0, 1)$:

$$a \Phi \left(\frac{1}{\sigma\sqrt{T}} \left(\log(a/q) + T(r + \frac{1}{2}\sigma^2) \right) \right) - qe^{-rT} \Phi \left(\frac{1}{\sigma\sqrt{T}} \left(\log(a/q) + T(r - \frac{1}{2}\sigma^2) \right) \right)$$

- But we can also estimate it through (iid) Monte Carlo!
 - Use (*) above (for fixed small $h > 0$, e.g. $h = 0.05$) to generate samples from the diffusion.
 - Any one run is highly variable. (file “RBS”, with $M = 1$)
 - But many runs give good estimate. (file “RBS”, with $M = 1000$)
- An “Asian call option” is similar, but with X_T replaced by $\bar{X}_{k,t} \equiv \frac{1}{k} \sum_{i=1}^k X_{iT/k}$, for some fixed positive integer k (e.g., $k = 8$).
 - Above “FACT” still holds (again with X_T replaced by $\bar{X}_{k,t}$).
 - Now formulas not so simple ... but can still simulate! (file “RAO”)
- So, can iid / importance / rejection / auxiliary sampling solve ALL of our problems? No!
 - Many challenging cases arise, e.g. from Bayesian statistics (later).
 - Some are high-dimensional, and the above algorithms fail.
 - Alternative algorithm: MCMC!

*** MARKOV CHAIN MONTE CARLO (MCMC) ***:

- Suppose have complicated, high-dimensional density $\pi = c g$.
- Want samples $X_1, X_2, \dots \sim \pi$. (Then can do Monte Carlo.) Difficult!
- Define a Markov chain (dependent random process: STA2006) X_0, X_1, X_2, \dots in such a way that for large enough n , $X_n \approx \pi$.

- Then can estimate $\mathbf{E}_\pi(h) \equiv \int h(x) \pi(x) dx$ by:

$$\mathbf{E}_\pi(h) \approx \frac{1}{M - B} \sum_{i=B+1}^M h(X_i),$$

where B (“burn-in”) is chosen large enough so $X_B \approx \pi$, and M is chosen large enough to get good Monte Carlo estimates.

- How to design such a Markov chain? One good way is:
- **METROPOLIS ALGORITHM (1953):**
 - Choose some initial value X_0 (perhaps random).
 - Then, given X_{n-1} , choose a proposal state $Y_n \sim MVN(X_{n-1}, \sigma^2 I)$ for some fixed $\sigma > 0$ (say).
 - Let $A_n = \pi(Y_n) / \pi(X_{n-1}) = g(Y_n) / g(X_{n-1})$, and $U_n \sim \text{Uniform}[0, 1]$.
 - Then, if $U_n < A_n$, set $X_n = Y_n$ (“accept”), otherwise set $X_n = X_{n-1}$ (“reject”).
 - Repeat, for $n = 1, 2, 3, \dots, M$.
 - (Note: only need to compute $\pi(Y_n) / \pi(X_{n-1})$, so the normalising constant c cancels and is not required.)
 - (Why does it work? Markov chain theory – later!)
 - Try it: www.probability.ca/metropolis (Javascript; formerly Java.)
 - Note: This version is called “random walk Metropolis” (RWM). Why? Because the proposals, if we always accepted them, would form a traditional random walk process.
- How large B ? Difficult to say! Some theory (later) ... usually just use trial-and-error / statistical analysis of output, and hope for the best ...
- What initial value X_0 ?
 - Virtually any one will do, but “central” ones best.
 - Can also use an “overdispersed starting distribution”: choose X_0 randomly from some distribution that “covers” the “important” parts of the state space. Good for checking consistency ...
- **EXAMPLE:** $g(y) = y^3 \sin(y^4) \cos(y^5) \mathbf{1}_{0 < y < 1}$.
 - Want to compute (again!) $\mathbf{E}_\pi(h)$ where $h(y) = y^2$.
 - Use Metropolis algorithm with proposal $Y \sim N(X, 1)$. [file “Rmet”]
 - Works pretty well, but lots of variability!
 - Plot: appears to have “good mixing”.
 - acf: has some serial autocorrelation $\mathbf{E}(X_n X_{n+k})$. Important! (Soon.)
- **EXAMPLE:** $\pi(x_1, x_2) = C |\cos(\sqrt{x_1 x_2})| I(0 \leq x_1 \leq 5, 0 \leq x_2 \leq 4)$.

- Want to compute $\mathbf{E}_\pi(h)$, where $h(x_1, x_2) = e^{x_1} + (x_2)^2$.
- Metropolis algorithm (file “Rmet2”) . . . works, but large uncertainty.
- Gets between about 34 and 44 . . . (Mathematica gets 38.7044)
- Individual plots appear to have “good mixing” . . .
- Joint plot shows fewer samples where $x_1 x_2 \approx (\pi/2)^2 \doteq 2.5$. . .

END WEEK #4

- OPTIMAL SCALING:
 - What if we change σ ? How does that affect estimate? plot? acf?
 - Can change proposal distribution to $Y_n \sim MVN(X_{n-1}, \sigma^2 I)$ for any choice of $\sigma > 0$. Which is best?
 - If σ too small, then usually accept, but chain won’t move much.
 - If σ too large, then will usually reject proposals, so chain still won’t move much.
 - Optimal: need σ “just right” to avoid both extremes. (“Goldilocks Principle”)
 - Can experiment (“www.probability.ca/metropolis”, “Rmet”, “Rmet2”).
 - Some theory . . . limited . . . active area of research . . .
 - General principle: the acceptance rate should be far from 0 and far from 1.
 - Surprising Fact: In a certain idealised high-dimensional limit, optimal acceptance rate is 0.234 (!). [Roberts et al., Ann Appl Prob 1997; [Roberts and Rosenthal, Stat Sci 2001](#)] (More later!)

MCMC STANDARD ERROR:

- What about MCMC’s standard error, i.e. uncertainty?
 - It’s usually larger than in the i.i.d. case (due to the positive correlations), and harder to quantify.
- Simplest: re-run the chain many times, with same M and B , with different initial values drawn from some overdispersed starting distribution, and get a fresh estimate each time, and then compute the standard error of the sequence of estimates.
 - Then can analyse the estimates obtained as iid . . .
- But how to estimate standard error from a single run?
- i.e., how to estimate $v \equiv \text{Var}\left(\frac{1}{M-B} \sum_{i=B+1}^M h(X_i)\right)$?
 - For simplicity, let $\bar{h}(x) = h(x) - \mathbf{E}_\pi(h)$, so $\mathbf{E}_\pi(\bar{h}) = 0$.

– And, assume B large enough that $X_i \approx \pi$ for $i > B$.

– Then, for large $M - B$,

$$\begin{aligned}
 v &\approx \mathbf{E}_\pi \left[\left(\left[\frac{1}{M-B} \sum_{i=B+1}^M h(X_i) \right] - \mathbf{E}_\pi(h) \right)^2 \right] = \mathbf{E}_\pi \left[\left(\frac{1}{M-B} \sum_{i=B+1}^M \bar{h}(X_i) \right)^2 \right] \\
 &= \frac{1}{(M-B)^2} \left[(M-B) \mathbf{E}_\pi[\bar{h}(X_i)^2] + 2(M-B-1) \mathbf{E}_\pi[\bar{h}(X_i)\bar{h}(X_{i+1})] \right. \\
 &\quad \left. + 2(M-B-2) \mathbf{E}_\pi[\bar{h}(X_i)\bar{h}(X_{i+2})] + \dots \right] \\
 &\approx \frac{1}{M-B} \left(\mathbf{E}_\pi[\bar{h}(X_i)^2] + 2 \mathbf{E}_\pi[\bar{h}(X_i)\bar{h}(X_{i+1})] + 2 \mathbf{E}_\pi[\bar{h}(X_i)\bar{h}(X_{i+2})] + \dots \right) \\
 &= \frac{1}{M-B} \left(\text{Var}_\pi(h) + 2 \text{Cov}_\pi(h(X_i), h(X_{i+1})) + 2 \text{Cov}_\pi(h(X_i), h(X_{i+2})) + \dots \right) \\
 &= \frac{1}{M-B} \text{Var}_\pi(h) \left(1 + 2 \text{Corr}_\pi(h(X_i), h(X_{i+1})) + 2 \text{Corr}_\pi(h(X_i), h(X_{i+2})) + \dots \right) \\
 &\equiv \frac{1}{M-B} \text{Var}_\pi(h) (\text{varfact}) = (\text{iid variance}) (\text{varfact}),
 \end{aligned}$$

where

$$\begin{aligned}
 \text{“varfact”} &= 1 + 2 \sum_{k=1}^{\infty} \text{Corr}_\pi(h(X_0), h(X_k)) \equiv 1 + 2 \sum_{k=1}^{\infty} \rho_k \\
 &= 2 \left(\sum_{k=0}^{\infty} \rho_k \right) - 1 = \sum_{k=-\infty}^{\infty} \rho_k
 \end{aligned}$$

since $\rho_0 = 1$ and $\rho_{-k} = \rho_k$.

- Also called “integrated auto-correlation time” or “ACT”.
- Then can estimate both iid variance, and varfact, from the sample run.
- Note: to compute varfact, don’t sum over all k , just e.g. until, say, $|\rho_k| < 0.05$ or $\rho_k < 0$ or ...
 - (Can use R’s built-in “acf” function, hopefully with a good choice of the “lag.max” parameter. Or can write your own – better.)
- Then standard error = $se = \sqrt{v} = (\text{iid-se}) \sqrt{\text{varfact}}$.
- e.g. the files Rmet and Rmet2. (Recall: true answers are about 0.766 and 38.7, respectively.)
 - Usually varfact $\gg 1$; try to get “better” chains so varfact smaller.
 - Sometimes even try to design chain to get varfact < 1 (“antithetic”).
 - Work in parallel? (Antithetically??) [Some work](#), but limited. (Project?)

CONFIDENCE INTERVALS:

- Suppose we estimate $u \equiv \mathbf{E}_\pi(h)$ by the quantity $e = \frac{1}{M-B} \sum_{i=B+1}^M h(X_i)$, and obtain an estimate e and an approximate variance (as above) v .
- Then what is, say, a 95% confidence interval for u ?
- Well, if have central limit theorem (CLT), then for large $M - B$, $e \approx N(u, v)$.
 - So $(e - u) v^{-1/2} \approx N(0, 1)$.
 - So, $\mathbf{P}(-1.96 < (e - u) v^{-1/2} < 1.96) \approx 0.95$.
 - So, $\mathbf{P}(-1.96 \sqrt{v} < e - u < 1.96 \sqrt{v}) \approx 0.95$.
 - i.e., with probability 95%, the interval $(e - 1.96 \sqrt{v}, e + 1.96 \sqrt{v})$ will contain u .
 - (Again, strictly speaking, should use something like a “t” distribution, not the normal distribution . . . but if $M - B$ large that doesn’t really matter – ignore it for now.)
- e.g. the files Rmet and Rmet2. (Recall: true answers are about 0.766 and 38.7, respectively.)
- But does a CLT even hold??
 - Does not follow from classical i.i.d. CLT. Does not always hold. But often does.
 - For example, CLT holds if chain is “geometrically ergodic” (later!) and $\mathbf{E}_\pi(|h|^{2+\delta}) < \infty$ for some $\delta > 0$.
 - (If chain also reversible then don’t need δ : [Roberts and Rosenthal, “Geometric ergodicity and hybrid Markov chains”, ECP 1997.](#))
 - Can get alternative (slightly larger) confidence intervals even without a CLT, if have consistent variance estimator: [Rosenthal, “Simple confidence intervals for MCMC without CLTs”, EJS 2017.](#))
- So MCMC is more complicated than standard Monte Carlo.
 - Why should we bother? Some problems are too challenging for other methods! (e.g. Bayesian – later)

METROPOLIS-HASTINGS ALGORITHM:

- The Metropolis algorithm doesn’t always work well.
 - Sometimes other MCMC algorithms can help too.
 - With above theory, can derive other valid algorithms!
- Note: key facts about $q(x, y)$ are symmetry, and irreducibility.
 - So, could replace $Y_n \sim N(X_{n-1}, 1)$ by e.g. $Y_n \sim \text{Uniform}[X_{n-1} - 1, X_{n-1} + 1]$, or (on discrete space) $Y_n = X_{n-1} \pm 1$ prob. $\frac{1}{2}$ each, etc.

- Metropolis algorithm still works provided proposal distribution is symmetric, i.e. $q(x, y) = q(y, x)$.
- But what if q is not symmetric?
- Hastings, Biometrika 1970 [Canadian! see www.probability.ca/hastings]:
 - Claim: If we replace “ $A_n = \pi(Y_n) / \pi(X_{n-1})$ ” by $A_n = \frac{\pi(Y_n)q(Y_n, X_{n-1})}{\pi(X_{n-1})q(X_{n-1}, Y_n)}$, then the algorithm is still valid even if q is not symmetric.
 - That is, we still accept if $U_n < A_n$, otherwise reject.
 - (Intuition: if $q(x, y) \gg q(y, x)$, then Metropolis chain would spend too much time at y and not enough at x , so need to accept fewer moves $x \rightarrow y$.)
 - Do require that $q(x, y) > 0$ iff $q(y, x) > 0$.
 - Why is it valid? Later!
- EXAMPLE: again $\pi(x_1, x_2) = C |\cos(\sqrt{x_1 x_2})| I(0 \leq x_1 \leq 5, 0 \leq x_2 \leq 4)$, and $h(x_1, x_2) = e^{x_1} + (x_2)^2$. (Mathematica gives $\mathbf{E}_\pi(h) \doteq 38.7044$.)
 - Proposal distribution: $Y_n \sim MVN(X_{n-1}, \sigma^2(1 + |X_{n-1}|)^2 I)$.
 - (Intuition: larger proposal variance if farther from center.)
 - So, $q(x, y) = C(1 + |x|^2)^{-2} \exp(-|y - x|^2 / 2\sigma^2(1 + |x|^2)^2)$.
 - Then, can run Metropolis-Hastings algorithm. (file “RMH”)
 - Usually get between 34 and 43, with claimed standard error ≈ 2 .

INDEPENDENCE SAMPLER:

- Propose $\{Y_n\} \sim q(\cdot)$, i.e. the $\{Y_n\}$ are i.i.d. from some fixed density q , independent of X_{n-1} . (e.g. $Y_n \sim MVN(0, I_d)$)
 - Then accept if $U_n < A_n$ where $U_n \sim \text{Uniform}[0, 1]$ and $A_n = \frac{\pi(Y_n)q(X_{n-1})}{\pi(X_{n-1})q(Y_n)}$.
 - Special case of the Metropolis-Hastings algorithm, where $Y_n \sim q(X_{n-1}, \cdot)$, and $A_n = \frac{\pi(Y_n)q(Y_n, X_{n-1})}{\pi(X_{n-1})q(X_{n-1}, Y_n)}$.
 - Very special case: if $q(y) \equiv \pi(y)$, i.e. propose exactly from target density π , then $A_n \equiv 1$, i.e. make great proposals, and always accept them (iid).
- e.g. independence sampler with $\pi(x) = e^{-x}$ and $q(y) = ke^{-ky}$ for $x > 0$.
 - Then if $X_{n-1} = x$ and $Y_n = y$, then $A_n = \frac{e^{-y}ke^{-ky}}{e^{-x}ke^{-ky}} = e^{(k-1)(y-x)}$. (file “Rind”)
 - $k = 1$: iid sampling (great).
 - $k = 0.01$: proposals way too large (so-so).
 - $k = 5$: proposals somewhat too small (terrible).
 - And with $k = 5$, confidence intervals often miss 1. (file “Rind2”)

- Why is large k so much worse than small k ? (Later!)

LANGEVIN ALGORITHM:

- Special case of Metropolis-Hastings algorithm.
- $Y_n \sim MVN(X_{n-1} + \frac{1}{2} \sigma^2 \nabla \log \pi(X_{n-1}), \sigma^2 I)$.
- Intuition: tries to move in direction where π increasing.
- Based on discrete approximation to “Langevin diffusion”.
- Usually more efficient, but requires knowledge and computation of $\nabla \log \pi$.
- For theory, see e.g. Roberts & Tweedie, Bernoulli **2(4)**, 341–363, 1996; Roberts & Rosenthal, JRSSB **60**, 255–268, 1998.

COMPONENTWISE (VARIABLE-AT-A-TIME) MCMC:

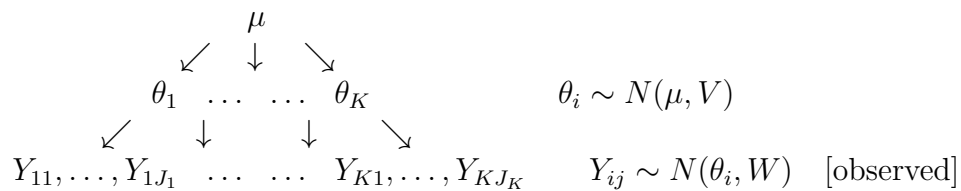
- Propose to move just one coordinate at a time, leaving all the other coordinates fixed (since changing all coordinates at once may be difficult).
 - e.g. proposal Y_n has $Y_{n,i} \sim N(X_{n-1,i}, \sigma^2)$, with $Y_{n,j} = X_{n-1,j}$ for $j \neq i$.
 - (Here $Y_{n,i}$ is the i^{th} coordinate of Y_n .)
- Then accept/reject with usual Metropolis rule (symmetric proposals: “Componentwise Metropolis”, or “Variable-at-a-time Metropolis”, or “Metropolis-within-Gibbs”) or Metropolis-Hastings rule (non-symmetric proposals: “Componentwise Metropolis-Hastings”, or “Variable-at-a-time Metropolis-Hastings”, or “Metropolis-Hastings-within-Gibbs”).
- Need to choose which coordinate to update each time ...
 - Could choose coordinates in sequence $1, 2, \dots, d, 1, 2, \dots$ (“systematic-scan”).
 - Or, choose coordinate $\sim \text{Uniform}\{1, 2, \dots, d\}$ each time (“random-scan”).
 - Note: one systematic-scan iteration corresponds to d random-scan ones ...
- EXAMPLE: again $\pi(x_1, x_2) = C |\cos(\sqrt{x_1 x_2})| I(0 \leq x_1 \leq 5, 0 \leq x_2 \leq 4)$, and $h(x_1, x_2) = e^{x_1} + (x_2)^2$. (Recall: Mathematica gives $\mathbf{E}_\pi(h) \doteq 38.7044$.)
 - Works with systematic-scan (file “Rcompwise”) or random-scan (file “Rcompwise2”).
- So, lots of MCMC algorithms to choose from.
 - Why do we need them all?
 - To compute with complicated models! For example ...

BAYESIAN STATISTICS:

- Have unknown parameter(s) θ , and a statistical model (likelihood function) for how the distribution of the data Y depends on θ : $\mathcal{L}(Y | \theta)$.
- Have a prior distribution, representing our “initial” (subjective?) probabilities for θ : $\mathcal{L}(\theta)$.
- Combining these gives a full joint distribution for θ and Y , i.e. $\mathcal{L}(\theta, Y)$.
- Then posterior distribution of θ , $\pi(\theta)$, is then the conditional distribution of θ , conditioned on the observed data y , i.e. $\pi(\theta) = \mathcal{L}(\theta | Y = y)$.
 - In terms of densities, if have prior density $f_\theta(\theta)$, and likelihood $f_{Y|\theta}(y, \theta)$, then joint density is $f_{\theta, Y}(\theta, y) = f_\theta(\theta) f_{Y|\theta}(y, \theta)$, and posterior density is

$$\pi(\theta) = \frac{f_{\theta, Y}(\theta, y)}{f_Y(y)} = C f_{\theta, Y}(\theta, y) = C f_\theta(\theta) f_{Y|\theta}(y, \theta).$$

- Bayesian Statistics Example: VARIANCE COMPONENTS MODEL (a.k.a. “random effects model”):



- Suppose some population has overall mean μ (unknown).
- Population consists of K groups.
- Observe Y_{i1}, \dots, Y_{iJ_i} from group i , for $1 \leq i \leq K$.
- Assume $Y_{ij} \sim N(\theta_i, W)$ (cond. ind.), where θ_i and W unknown.
- Assume the different θ_i are “linked” by $\theta_i \sim N(\mu, V)$ (cond. ind.), with μ and V also unknown.
- Want to estimate some or all of $V, W, \mu, \theta_1, \dots, \theta_K$.
- Bayesian approach: use prior distributions, e.g. (“conjugate”):

$$V \sim IG(a_1, b_1); \quad W \sim IG(a_2, b_2); \quad \mu \sim N(a_3, b_3)$$

(indep), where a_i, b_i known constants, and $IG(a, b)$ is the “inverse gamma” distribution, with density $\frac{b^a}{\Gamma(a)} e^{-b/x} x^{-a-1}$ for $x > 0$.

- Combining the above dependencies, we see that the joint density is (for $V, W > 0$):

$$\begin{aligned} & f(V, W, \mu, \theta_1, \dots, \theta_K, Y_{11}, Y_{12}, \dots, Y_{KJ_K}) \\ &= \left(\frac{b_1^{a_1}}{\Gamma(a_1)} e^{-b_1/V} V^{-a_1-1} \right) \left(\frac{b_2^{a_2}}{\Gamma(a_2)} e^{-b_2/W} W^{-a_2-1} \right) \left(\frac{1}{\sqrt{2\pi b_3}} e^{-(\mu-a_3)^2/2b_3} \right) \times \end{aligned}$$

$$\begin{aligned}
& \times \left(\prod_{i=1}^K \frac{1}{\sqrt{2\pi V}} e^{-(\theta_i - \mu)^2/2V} \right) \left(\prod_{i=1}^K \prod_{j=1}^{J_i} \frac{1}{\sqrt{2\pi W}} e^{-(Y_{ij} - \theta_i)^2/2W} \right) \\
& = C_2 e^{-b_1/V} V^{-a_1-1} e^{-b_2/W} W^{-a_2-1} e^{-(\mu - a_3)^2/2b_3} V^{-K/2} W^{-\frac{1}{2} \sum_{i=1}^K J_i} \times \\
& \quad \times \exp \left[-\sum_{i=1}^K (\theta_i - \mu)^2/2V \right] \exp \left[-\sum_{i=1}^K \sum_{j=1}^{J_i} (Y_{ij} - \theta_i)^2/2W \right].
\end{aligned}$$

- Then

$$\begin{aligned}
& \pi(V, W, \mu, \theta_1, \dots, \theta_K) \\
& = f(V, W, \mu, \theta_1, \dots, \theta_K, Y_{11}, Y_{12}, \dots, Y_{KJ_K}) / f_Y(Y_{11}, Y_{12}, \dots, Y_{KJ_K}) \\
& \quad \propto f(V, W, \mu, \theta_1, \dots, \theta_K, Y_{11}, Y_{12}, \dots, Y_{KJ_K}) \\
& = C_3 e^{-b_1/V} V^{-a_1-1} e^{-b_2/W} W^{-a_2-1} e^{-(\mu - a_3)^2/2b_3} V^{-K/2} W^{-\frac{1}{2} \sum_{i=1}^K J_i} \times \\
& \quad \times \exp \left[-\sum_{i=1}^K (\theta_i - \mu)^2/2V \right] \exp \left[-\sum_{i=1}^K \sum_{j=1}^{J_i} (Y_{ij} - \theta_i)^2/2W \right].
\end{aligned}$$

- NOTE: Many applications of variance components model, e.g.:
 - Predicting success at law school (D. Rubin, JASA 1980), $K = 82$ schools.
 - Melanoma (skin cancer) recurrence (http://www.mssanz.org.au/MODSIM07/papers/52_s24/Analysing_Clinicals24_Bartolucci_.pdf), with $K = 19$ different patient categories.
 - Comparing baseball home-run hitters (J. Albert, The American Statistician 1992), $K = 12$ players.
 - Analysing fabric dyes (Davies 1967; Box/Tiao 1973; Gelfand/Smith JASA 1990), $K = 6$ batches of dyestuff, $J_i \equiv 5$. (data in file “[Rdye](#)”)
- Here, the dimension is $d = K + 3$, e.g. $K = 19$, $d = 22$. High!
- How to compute/estimate, say, $\mathbf{E}_\pi(W/V)$, or the effect of changing b_1 ?
 - Numerical integration? No, too high-dimensional!
 - Importance sampling? Perhaps, but what “ f ”? Too inefficient!
 - Rejection sampling? What “ f ”? What “ K ”? Virtually no samples!
 - Perhaps MCMC can work!
 - But need clever, useful MCMC algorithms!
 - Perhaps Metropolis, or ...
- ASIDE: For big complicated π , often better to work with logarithms, e.g. accept iff $\log(U_n) < \log(A_n) = \log(\pi(Y_n)) - \log(\pi(X_{n-1}))$.
 - Then only need to compute $\log(\pi(x))$; helps avoid overflow problems.
 - So, better to program on log scale: $\log \pi(V, W, \mu, \theta_1, \dots, \theta_K) = \dots$

- Also sometimes simpler, e.g. if $\pi(x) = \exp\left(\sum_{i<j} |x_j - x_i|\right)$, then $\log(\pi(x)) = \sum_{i<j} |x_j - x_i|$. (Best to type in the log formula directly.)

GIBBS SAMPLER:

- (Special case of Componentwise Metropolis-Hastings.)
- Proposal distribution for i^{th} coordinate is equal to the full conditional distribution of that coordinate (according to π), conditional on the current values of all the other coordinates.
 - Can use either systematic or random scan, just like above.
 - Then, always accept. Why? Later!
 - (Intuition: if start in stationary distribution, then update one coordinate from its conditional stationary distribution (and always accept), then the distribution remains the same, i.e. stationarity is preserved.)

END WEEK #5

- EXAMPLE: Variance Components Model:
 - Update of μ (say) should be from conditional density of μ , conditional on current values of all the other coordinates: $\mathcal{L}(\mu | V, W, \theta_1, \dots, \theta_K, Y_{11}, \dots, Y_{JK})$.
 - This conditional density is proportional to the full joint density, but with all variables besides μ treated as constant.
 - Recall: full joint density is:

$$= C_3 e^{-b_1/V} V^{-a_1-1} e^{-b_2/W} W^{-a_2-1} e^{-(\mu-a_3)^2/2b_3} V^{-K/2} W^{-\frac{1}{2} \sum_{i=1}^K J_i} \times \\ \times \exp\left[-\sum_{i=1}^K (\theta_i - \mu)^2/2V\right] \exp\left[-\sum_{i=1}^K \sum_{j=1}^{J_i} (Y_{ij} - \theta_i)^2/2W\right].$$

- So, combining “constants” (w.r.t. μ), the conditional density of μ is

$$C_4 e^{-(\mu-a_3)^2/2b_3} \exp\left[-\sum_{i=1}^K (\theta_i - \mu)^2/2V\right].$$

- This equals (verify this! HW!)

$$C_5 \exp\left(-\mu^2\left(\frac{1}{2b_3} + \frac{K}{2V}\right) + \mu\left(\frac{a_3}{b_3} + \frac{1}{V} \sum_{i=1}^K \theta_i\right)\right).$$

- Side calculation: if $\mu \sim N(m, v)$, then density $\propto e^{-(\mu-m)^2/2v} \propto e^{-\mu^2(1/2v) + \mu(m/v)}$.
- Hence, here $\mu \sim N(m, v)$, where $1/2v = \frac{1}{2b_3} + \frac{K}{2V}$ and $m/v = \frac{a_3}{b_3} + \frac{1}{V} \sum_{i=1}^K \theta_i$.

- Solve: $v = b_3 V / (V + K b_3)$, and $m = (a_3 V + b_3 \sum_{i=1}^K \theta_i) / (V + K b_3)$.
- So, in Gibbs Sampler, each time μ is updated, we sample it from $N(m, v)$ for this m and v (and always accept).
- Similarly (HW!), conditional distribution for V is:

$$C_6 e^{-b_1/V} V^{-a_1-1} V^{-K/2} \exp \left[- \sum_{i=1}^K (\theta_i - \mu)^2 / 2V \right], \quad V > 0.$$

- Recall that “ $IG(r, s)$ ” has density $\frac{s^r}{\Gamma(r)} e^{-s/x} x^{-r-1}$ for $x > 0$.
- So, conditional distribution for V equals $IG(a_1 + K/2, b_1 + \frac{1}{2} \sum_{i=1}^K (\theta_i - \mu)^2)$.
- Can similar compute conditional distributions for W and θ_i (HW!).
- The systematic-scan Gibbs sampler then proceeds (HW!) by:
 - Update V from its conditional distribution $IG(\dots, \dots)$.
 - Update W from its conditional distribution $IG(\dots, \dots)$.
 - Update μ from its conditional distribution $N(\dots, \dots)$.
 - Update θ_i from its conditional distribution $N(\dots, \dots)$, for $i = 1, 2, \dots, K$.
 - Repeat all of the above M times.
- Or, the random-scan Gibbs sampler proceeds by choosing one of $V, W, \mu, \theta_1, \dots, \theta_K$ uniformly at random, and then updating that coordinate from its corresponding conditional distribution.
 - Then repeat this step M times [or $M(K + 3)$ times?].
 - How well does it work? HW!

SUBSAMPLING (THINNING):

- The autocorrelations (acf) of an MCMC run usually start near 1, and then decrease until they become negligible after some lag L . (file “Rmetnorm”)
 - This means that every L^{th} iteration of the chain is approximately independent.
 - But also, after the burn-in B , the chain is approximately in stationarity, i.e. has distribution approximately π .
 - So, the states $X_B, X_{B+L}, X_{B+2L}, X_{B+3L}, \dots$ are approximately i.i.d. samples from π . Good?
- When running MCMC, some people use “subsampling” (or, “thinning”) by just using every S^{th} sample, i.e. by using the Markov chain $\{X_{B+S_n}\}$.
 - If $S \geq L$, then the $\{X_{B+S_n}\}$ are approximately independent.
 - This could be useful: i.i.d. samples, classical standard error and confidence intervals, good tests for accuracy, etc.

- But does it improve the actual estimator? e.g. its variance?
 - The number of samples is reduced from $M - B$ to $(M - B)/S$, which increases the estimator variance by a factor of S . Bad.
 - But the varfact is reduced from $1 + 2 \sum_{k=1}^{\infty} \rho_k$ to $1 + 2 \sum_{k=1}^{\infty} \rho_{Sk}$ (good).
 - So, which is larger, $1 + 2 \sum_{k=1}^{\infty} \rho_k$, or $S (1 + 2 \sum_{k=1}^{\infty} \rho_{Sk})$?
 - Well, we can re-write $S (1 + 2 \sum_{k=1}^{\infty} \rho_{Sk})$ as

$$(1 + \dots + 1) + (2\rho_S + \dots + 2\rho_S) + (2\rho_{2S} + \dots + 2\rho_{2S}) + \dots$$

- We want to compare this to

$$(1 + 2\rho_1 + \dots + 2\rho_{S-1}) + (2\rho_S + 2\rho_{S+1} + \dots + 2\rho_{2S-1}) + (2\rho_{2S} + \dots) + \dots$$
- Assuming the ρ_i are decreasing, and ignoring some of the “2” factors, the first (subsampled) variance is larger.
- This calculation seems to suggest that subsampling is not advantageous.
 - In fact, for “reversible” Markov chains (defined later), consecutive sums $\rho_{2i} + \rho_{2i+1}$ are decreasing, and the first (subsampled) variance is always larger: [Geyer 1992](#)
 - More generally, even if not reversible, subsampling always increases the estimator variance: [Maceachern & Berliner, 1994](#)
- Conclusion: Subsampling makes the estimator variance larger, i.e. subsampling does not improve the estimator.
 - However, if lots of extra computation is required to compute the functional values $h(X_i)$, then subsampling might help: [Owen 2017](#)

JUSTIFICATION: WHY DOES METROPOLIS ALG WORK?:

- (Uses Markov chain theory ... e.g. STA447/2006 ... already know?)
- Basic fact: if a Markov chain is “irreducible” and “aperiodic”, with “stationarity distribution” π , then $\mathcal{L}(X_n) \rightarrow \pi$ as $n \rightarrow \infty$. More precisely:
- THEOREM: If Markov chain is irreducible, with stationarity probability density π , then for π -a.e. initial value $X_0 = x$,
 - (a) if $\mathbf{E}_\pi(|h|) < \infty$, then $\lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i=1}^n h(X_i) = \mathbf{E}_\pi(h) \equiv \int h(x) \pi(x) dx$;
 - and (b) if chain aperiodic, then also $\lim_{n \rightarrow \infty} \mathbf{P}(X_n \in S) = \int_S \pi(x) dx$ for all $S \subseteq \mathcal{X}$.
- Let’s figure out what this all means ...
- Notation: $P(i, j) = \mathbf{P}(X_{n+1} = j | X_n = i)$ (discrete case), or $P(x, A) = \mathbf{P}(X_{n+1} \in A | X_n = x)$ (general case). Also $\Pi(A) = \int_A \pi(x) dx$.

- Well, irreducible means that you have positive probability of eventually getting from anywhere to anywhere else.
 - Discrete case: for all $i, j \in \mathcal{X}$ (the state space), there is $n \in \mathbf{N}$ such that $P(X_n = j | X_0 = i) > 0$.
 - Actually, we only need to require this for states j such that $\pi(j) > 0$.
 - General case: for all $x \in \mathcal{X}$, and for all $A \subseteq \mathcal{X}$ with $\Pi(A) > 0$, there is $n \in \mathbf{N}$ such that $P(X_n \in A | X_0 = x) > 0$. (“ π -irreducible”)
 - (Since usually $P(X_n = y | X_0 = x) = 0$ for all y .)
 - Irreducibility is usually satisfied for MCMC.
- And, aperiodic means there are no forced cycles, i.e. there do not exist disjoint non-empty subsets $\mathcal{X}_1, \mathcal{X}_2, \dots, \mathcal{X}_d$ for $d \geq 2$, such that $P(x, \mathcal{X}_{i+1}) = 1$ for all $x \in \mathcal{X}_i$ ($1 \leq i \leq d-1$), and $P(x, \mathcal{X}_1) = 1$ for all $x \in \mathcal{X}_d$. [Diagram.]
 - (Discrete case: Equivalent that $\gcd\{n : p^n(i, i) > 0\} = 1 \forall i$.)
 - This is true for virtually any Metropolis algorithm, e.g. it’s true if $P(x, \{x\}) > 0$ for any one state $x \in \mathcal{X}$, e.g. if positive prob of rejection.
 - Also true if $P(x, \cdot)$ has positive density throughout S , for all $x \in S$, for some $S \subseteq \mathcal{X}$ with $\Pi(S) > 0$. (e.g. Normal proposals)
 - Not quite guaranteed, e.g. $\mathcal{X} = \{0, 1, 2, 3\}$, and π uniform on \mathcal{X} , and $Y_n = X_{n-1} \pm 1 \pmod{4}$. [Diagram.] But almost always holds.
- What about Π being a stationary distribution?
 - This means that if we start with the probabilities Π , and then run the Markov chain for one step, that we will still have the probabilities Π .
 - Will this be true for the Metropolis algorithm?

END WEEK #6

- Begin with DISCRETE CASE (e.g. www.probability.ca/metropolis).
 - State space is \mathcal{X} , e.g. $\mathcal{X} \equiv \{1, 2, 3, 4, 5, 6\}$.
- Here stationary means that if $X_0 \sim \pi$, i.e. $\mathbf{P}(X_0 = i) = \pi(i)$ for all i , then also $X_1 \sim \pi$, i.e. $\mathbf{P}(X_1 = j) = \pi(j)$ for all j .
 - But $\mathbf{P}(X_1 = j) = \sum_{i \in S} \mathbf{P}(X_0 = i, X_1 = j) = \sum_{i \in S} \mathbf{P}(X_0 = i) P(i, j)$.
 - So, π is stationary if $\sum_{i \in S} \pi(i) P(i, j) = \pi(j)$ for all j .
- Let $q(x, y) = \mathbf{P}(Y_n = y | X_{n-1} = x)$ be the proposal distribution, e.g. perhaps $q(x, x+1) = q(x, x-1) = 1/2$.
 - Assume that q is symmetric, i.e. $q(x, y) = q(y, x)$ for all $x, y \in \mathcal{X}$.
 - Then if $\alpha(x, y)$ is the probability of accepting a proposed move from

x to y , then

$$\begin{aligned}\alpha(x, y) &= \mathbf{P}(U_n < A_n \mid X_{n-1} = x, Y_n = y) \\ &= \mathbf{P}\left(U_n < \frac{\pi(y)}{\pi(x)}\right) = \min\left[1, \frac{\pi(y)}{\pi(x)}\right].\end{aligned}$$

– (Assume for simplicity that $\pi(x) > 0$.)

- Then we compute that for $i, j \in \mathcal{X}$ with $i \neq j$,

$$P(i, j) = q(i, j) \alpha(i, j) = q(i, j) \min\left(1, \frac{\pi(j)}{\pi(i)}\right).$$

– Hence, using the symmetry of q ,

$$\begin{aligned}\pi(i) P(i, j) &= q(i, j) \min(\pi(i), \pi(j)) \\ &= q(j, i) \min(\pi(i), \pi(j)) = \pi(j) P(j, i).\end{aligned}$$

– This (obviously) still holds if $i = j$.

– It follows that chain is “(time) reversible”, i.e.

$$\pi(i) P(i, j) = \pi(j) P(j, i) \quad \forall i, j \in \mathcal{X}.$$

– (Intuition: if $X_0 \sim \pi$, i.e. $\mathbf{P}(X_0 = i) = \pi(i)$ for all $i \in \mathcal{X}$, then $\mathbf{P}(X_0 = i, X_1 = j) = \pi(i) P(i, j) = \mathbf{P}(X_0 = j, X_1 = i) \dots$)

- Suppose now that $X_0 \sim \pi$, i.e. that $\mathbf{P}(X_0 = i) = \pi(i)$ for all $i \in \mathcal{X}$.

– Then using reversibility, we have

$$\begin{aligned}\mathbf{P}(X_1 = j) &= \sum_{i \in \mathcal{X}} \mathbf{P}(X_0 = i) P(i, j) = \sum_{i \in \mathcal{X}} \pi(i) P(i, j) \\ &= \sum_{i \in \mathcal{X}} \pi(j) P(j, i) = \pi(j) \sum_{i \in \mathcal{X}} P(j, i) = \pi(j),\end{aligned}$$

i.e. $X_1 \sim \pi$ too!

– So, the Markov chain “preserves” π , i.e. π is a stationary distribution.

– This is true for any Metropolis algorithm!

- It then follows from the Theorem (i.e., “Basic Fact”) that as $n \rightarrow \infty$, $\mathcal{L}(X_n) \rightarrow \pi$, i.e. $\lim_{n \rightarrow \infty} P(X_n = i) = \pi(i)$ for all $i \in \mathcal{X}$.

– Also follows that if $\mathbf{E}_\pi(|h|) < \infty$, then $\lim_{n \rightarrow \infty} \frac{1}{n} \sum_{i=1}^n h(X_i) = \mathbf{E}_\pi(h) \equiv \int h(x) \pi(x) dx$. (“LLN”)

JUSTIFICATION: GENERAL CONTINUOUS CASE:

- Some notation:

– Let \mathcal{X} be the state space of all possible values. (Usually $\mathcal{X} \subseteq \mathbf{R}^d$, e.g. $\mathcal{X} = \mathbf{R}^2$, or $\mathcal{X} = (0, \infty) \times (0, 1) \times \mathbf{R}^5$, or \dots)

- Let $\pi(x)$ be the target density function.
- Let $q(x, y)$ be the proposal density for y given x . (e.g. $q(x, y) = (2\pi\sigma)^{-d/2} \exp(-\sum_{i=1}^d (y_i - x_i)^2 / 2\sigma^2)$.)
- Assume again that q is symmetric: $q(x, y) = q(y, x)$.
- Let $\alpha(x, y) = \min[1, \frac{\pi(y)}{\pi(x)}]$ be probability of accepting a proposed move from x to y .
- Let $P(x, S) = \mathbf{P}(X_1 \in S \mid X_0 = x)$ be the transition probabilities.
- (Don't use $P(x, y)$ since that is usually 0.)
- Then if $x \notin S$, then

$$\begin{aligned} P(x, S) &= \mathbf{P}(Y_1 \in S, U_1 < A_1 \mid X_0 = x) \\ &= \int_S q(x, y) \alpha(x, y) dy = \int_S q(x, y) \min[1, \pi(y)/\pi(x)] dy. \end{aligned}$$

- Shorthand: write “ $P(x, dy)$ ” for the transition measure, i.e. a quantity whose integral over any subset S is equal to the transition probability to S , meaning that $P(x, S) = \int_{y \in S} P(x, dy)$.
- Then, for $x \neq y$, $P(x, dy) = q(x, y) \min[1, \pi(y)/\pi(x)] dy$.
- Hence, for $x \neq y$,

$$\begin{aligned} \pi(x) dx P(x, dy) &= \pi(x) dx q(x, y) \min[1, \pi(y)/\pi(x)] dy \\ &= q(x, y) \min[\pi(x), \pi(y)] dy dx. \end{aligned}$$

- This is symmetric in x and y , i.e. $\pi(x) dx P(x, dy) = \pi(y) dy P(y, dx)$ for all $x, y \in \mathcal{X}$.
- Shorthand: $\Pi(dx) P(x, dy) = \Pi(dy) P(y, dx)$. (“reversible”)
- How does “reversible” help? Just like for discrete chains!
- Indeed, suppose $X_0 \sim \Pi$, i.e. we “start in stationarity”. Then

$$\begin{aligned} \mathbf{P}(X_1 \in S) &= \int_{x \in \mathcal{X}} \pi(x) dx \int_{y \in S} P(x, dy) = \int_{x \in \mathcal{X}} \int_{y \in S} \Pi(dx) P(x, dy) \\ &= \int_{x \in \mathcal{X}} \int_{y \in S} \Pi(dy) P(y, dx) = \int_{y \in S} \Pi(dy) \int_{x \in \mathcal{X}} P(y, dx) \\ &= \int_{y \in S} \Pi(dy) = \int_{y \in S} \pi(y) dy \equiv \Pi(S), \end{aligned}$$

so also $X_1 \sim \Pi$. So, chain “preserves” Π , i.e. Π is stationary distribution.

- And, again, almost always irreducible and aperiodic.
- So, again, the Theorem applies.

EXAMPLES RE WHY DOES MCMC WORK:

- EXAMPLE #1: Metropolis algorithm where $\mathcal{X} = \mathbf{Z}$, $\pi(x) = 2^{-|x|}/3$, and $q(x, y) = \frac{1}{2}$ if $|x - y| = 1$, otherwise 0.
 - Reversible? Yes, it's a Metropolis algorithm!
 - π stationary? Yes, follows from reversibility!
 - Aperiodic? Yes, since $P(x, \{x\}) > 0$!
 - Irreducible? Yes: $\pi(x) > 0$ for all $x \in \mathcal{X}$, so can get from x to y in $|x - y|$ steps.
 - So, by theorem, probabilities and expectations converge to those of π
 - good.
- EXAMPLE #2: Same as #1, except now $\pi(x) = 2^{-|x|-1}$ for $x \neq 0$, with $\pi(0) = 0$.
 - Still reversible, π stationary, aperiodic, same as before.
 - Irreducible? No – can't go from positive to negative!
- EXAMPLE #3: Same as #2, except now $q(x, y) = \frac{1}{4}$ if $1 \leq |x - y| \leq 2$, otherwise 0.
 - Still reversible, π stationary, aperiodic, same as before.
 - Irreducible? Yes – can “jump over 0” to get from positive to negative, and back!
- EXAMPLE #4: Metropolis algorithm with $\mathcal{X} = \mathbf{R}$, and $\pi(x) = C e^{-x^6}$, and proposals $Y_n \sim \text{Uniform}[X_{n-1} - 1, X_{n-1} + 1]$.
 - Reversible? Yes since it's Metropolis, and $q(x, y)$ still symmetric.
 - π stationary? Yes since reversible!
 - Irreducible? Yes, since the n -step transitions $P^n(x, dy)$ have positive density whenever $|y - x| < n$.
 - Aperiodic? Yes since if periodic, then if e.g. $\mathcal{X}_1 \cap [0, 1]$ has positive measure, then possible to go from \mathcal{X}_1 directly to \mathcal{X}_1 , i.e. if $x \in \mathcal{X}_1 \cap [0, 1]$, then $P(x, \mathcal{X}_1) > 0$. (Or, even simpler: since $P(x, \{x\}) > 0$ for all $x \in \mathcal{X}$.)
 - So, by theorem, probabilities and expectations converge to those of π
 - good.
- EXAMPLE #5: Same as #4, except now $\pi(x) = C_1 e^{-x^6} (\mathbf{1}_{x < 2} + \mathbf{1}_{x > 4})$.
 - Still reversible and stationary and aperiodic, same as before.
 - But no longer irreducible: cannot jump from $[4, \infty)$ to $(-\infty, 2]$ or back.
 - So, does not converge.

- EXAMPLE #6: Same as #5, except now proposals are $Y_n \sim \text{Uniform}[X_{n-1} - 5, X_{n-1} + 5]$.
 - Still reversible and stationary and aperiodic, same as before.
 - And now irreducible, too: now can jump from $[4, \infty)$ to $(-\infty, 2]$ or back.
- EXAMPLE #7: Same as #6, except now $Y_n \sim \text{Uniform}[X_{n-1} - 5, X_{n-1} + 10]$.
 - Makes no sense – proposals not symmetric, so not a Metropolis algorithm! (Not even symmetrically zero, for the Metropolis-Hastings algorithm below, e.g. have positive density $3 \rightarrow 9$ but not $9 \rightarrow 3$.)

END WEEK #7

JUSTIFICATION OF METROPOLIS-HASTINGS:

- Can we modify the above proof to work for Metropolis-Hastings, too?
- For Metropolis, key was that the Markov chain is reversible, i.e. $\pi(x) P(x, y) = \pi(y) P(y, x)$, i.e. $q(x, y) \alpha(x, y) \pi(x)$ is symmetric in x and y .
 - If instead $A_n = \frac{\pi(Y_n) q(Y_n, X_{n-1})}{\pi(X_{n-1}) q(X_{n-1}, Y_n)}$, i.e. acceptance prob. $\equiv \alpha(x, y) = \min \left[1, \frac{\pi(y) q(y, x)}{\pi(x) q(x, y)} \right]$, then:

$$q(x, y) \alpha(x, y) \pi(x) = q(x, y) \min \left[1, \frac{\pi(y) q(y, x)}{\pi(x) q(x, y)} \right] \pi(x)$$

$$= \min \left[\pi(x) q(x, y), \pi(y) q(y, x) \right].$$
 - So, $\pi(x) P(x, y)$ is still symmetric, even if q wasn't.
 - So, still reversible. So, still have stationary distribution Π .
 - So, if irreducible and aperiodic (nearly always true), then can again apply usual Theorem, and again conclude that it converges to Π .
- Conclusion: For the Metropolis-Hastings algorithm, if we replace “ $A_n = \pi(Y_n) / \pi(X_{n-1})$ ” by $A_n = \frac{\pi(Y_n) q(Y_n, X_{n-1})}{\pi(X_{n-1}) q(X_{n-1}, Y_n)}$, and keep everything else the same (i.e., still accept if $U_n < A_n$, otherwise reject), then it is still valid, and still converges to the correct distribution Π .

JUSTIFICATION FOR VARIABLE-AT-A-TIME:

- The exact same justification works just like for the “regular” (full-dimensional) Metropolis and Metropolis-Hastings algorithms:
 - If we update the variables one-at-a-time (e.g. Metropolis-within-Gibbs, Metropolis-Hastings-within-Gibbs, etc.), then each individual step is still reversible (for the same reason), so π is still stationary.

- So, like any irreducible, aperiodic Markov chain with stationary distribution π , it will eventually converge to π .

JUSTIFICATION OF GIBBS SAMPLER:

- Special case of Metropolis-Hastings-within-Gibbs.
- Proposal distribution for i^{th} coordinate is equal to the conditional distribution of that coordinate (according to π), conditional on the current values of all the other coordinates.
- That is, $q_i(x, y) = C[x^{(-i)}] \pi(y)$ whenever $x^{(-i)} = y^{(-i)}$, where $x^{(-i)}$ means all coordinates except the i^{th} one.
- (And $q_i(x, y) = 0$ if $x^{(-i)} \neq y^{(-i)}$.)
- Here $C[x^{(-i)}]$ is the appropriate normalising constant (which depends on $x^{(-i)}$). So, will always have $C[x^{(-i)}] = C[y^{(-i)}]$.
- Then $A_n = \frac{\pi(Y_n) q_i(Y_n, X_{n-1})}{\pi(X_{n-1}) q_i(X_{n-1}, Y_n)} = \frac{\pi(Y_n) C[Y_n^{(-i)}] \pi(X_{n-1})}{\pi(X_{n-1}) C[X_{n-1}^{(-i)}] \pi(Y_n)} = 1$.
- So, always accept (i.e., can ignore the accept-reject step).

INITIAL DISTRIBUTION CONDITION:

- Why does the above Theorem say “ π -a.e.” $X_0 = x$?
- Example: $\mathcal{X} = \{1, 2, 3, 4, \dots\}$, and $P(1, \{1\}) = 1$, and for $x \geq 2$, $P(x, \{x+1\}) = 1 - (1/x^2)$ and $P(x, \{1\}) = 1/x^2$.
 - Stationary distribution?
 $\Pi(\cdot) = \delta_1(\cdot)$, i.e. $\Pi(S) = \mathbf{1}_{1 \in S}$ for $S \subseteq \mathcal{X}$.
 - Irreducible?
 Yes, since if $\Pi(S) > 0$ then $1 \in S$ so $P(x, S) \geq P(x, \{1\}) > 0$ for all $x \in \mathcal{X}$.
 - Aperiodic?
 Yes, since $P(1, \{1\}) > 0$.
 - Converges?
 Yes, by Theorem, for π -a.e. X_0 , have $\lim_{n \rightarrow \infty} \mathbf{P}(X_n \in S) = \Pi(S)$, i.e. $\lim_{n \rightarrow \infty} \mathbf{P}(X_n = 1) = 1$.
 - From everywhere?
 No! If $X_0 = x \geq 2$, then $\mathbf{P}[X_n = x+n \text{ for all } n] = \prod_{j=x}^{\infty} (1 - (1/j^2)) > 0$ (since $\sum_{j=x}^{\infty} (1/j^2) < \infty$), so $\lim_{n \rightarrow \infty} \mathbf{P}(X_n = 1) \neq 1$.
 - Convergence holds if $X_0 = 1$, which is π -a.e. since $\Pi(1) = 1$, but not from $X_0 = x \geq 2$.
- So, convergence subtle. But usually holds from any $x \in \mathcal{X}$. (“Harris recurrent”, see e.g. <http://probability.ca/jeff/ftpdire/harris.pdf>)

TEMPERED MCMC:

- Suppose $\Pi(\cdot)$ is multi-modal, i.e. has distinct “parts” (e.g., $\Pi = \frac{1}{2} N(0, 1) + \frac{1}{2} N(20, 1)$)
- Usual RWM with $Y_n \sim N(X_{n-1}, 1)$ (say) can explore well within each mode, but how to get from one mode to the other?
- Idea: if $\Pi(\cdot)$ were flatter, e.g. $\frac{1}{2} N(0, 10^2) + \frac{1}{2} N(20, 10^2)$, then much easier to get between modes.
- So: define a sequence $\Pi_1, \Pi_2, \dots, \Pi_m$ where $\Pi_1 = \Pi$ (“cold”), and Π_τ is flatter for larger τ (“hot”). (e.g. $\Pi_\tau = \frac{1}{2} N(0, \tau^2) + \frac{1}{2} N(20, \tau^2)$; file “Rtempered”)
- In the end, only “count” those samples where $\tau = 1$.
- Proceed by defining a joint Markov chain (x, τ) on $\mathcal{X} \times \{1, 2, \dots, m\}$, with stationary distribution $\bar{\Pi}$ defined by $\bar{\Pi}(S \times \{\tau\}) = \frac{1}{m} \Pi_\tau(S)$.
 - (Can also use other weights besides $\frac{1}{m}$.)
- The Markov chain should have both spatial moves (change x) and temperature moves (change τ).
 - e.g. perhaps chain alternates between:
 - (a) propose $x' \sim N(x, 1)$, accept with prob $\min\left(1, \frac{\bar{\pi}(x', \tau)}{\bar{\pi}(x, \tau)}\right) = \min\left(1, \frac{\pi_\tau(x')}{\pi_\tau(x)}\right)$.
 - (b) propose $\tau' = \tau \pm 1$ (prob $\frac{1}{2}$ each), accept with prob $\min\left(1, \frac{\bar{\pi}(x, \tau')}{\bar{\pi}(x, \tau)}\right) = \min\left(1, \frac{\pi_{\tau'}(x)}{\pi_\tau(x)}\right)$.
- Chain should converge to $\bar{\Pi}$.
- Then, as above, only “count” those samples where $\tau = 1$. (red)
- EXAMPLE: $\Pi = \frac{1}{2} N(0, 1) + \frac{1}{2} N(20, 1)$
 - Assume proposals are $Y_n \sim N(X_{n-1}, 1)$.
 - Mixing for Π : terrible! (file “Rtempered” with dotempering=FALSE and temp=1; note the small claimed standard error!)
 - Define $\Pi_\tau = \frac{1}{2} N(0, \tau^2) + \frac{1}{2} N(20, \tau^2)$, for $\tau = 1, 2, \dots, 10$.
 - Mixing better for larger τ ! (file “Rtempered” with dotempering=FALSE and temp=1,2,3,4,...,10)
 - (Compare graphs of π_1 and π_8 : plot commands at bottom of “Rtempered” ...)
 - So, use above “(a)–(b)” algorithm; converges fairly well to $\bar{\Pi}$. (file “Rtempered”, with dotempering=TRUE)
 - So, conditional on $\tau = 1$, converges to Π . (“points” command at end of file “Rtempered”)

- So, average of those $h(x)$ with $\tau = 1$ gives good estimate of $\mathbf{E}_\pi(h)$.

FINDING THE TEMPERED DENSITIES:

- Usually won't "know" about e.g. $\Pi_\tau = \frac{1}{2} N(0, \tau^2) + \frac{1}{2} N(20, \tau^2)$.
- Instead, can e.g. let $\pi_\tau(x) = c_\tau (\pi(x))^{1/\tau}$. (Sometimes write $\beta = 1/\tau$)
 - Then $\Pi_1 = \Pi$, and π_τ flatter for larger τ – good.
 - (e.g. if $\pi(x)$ density of $N(\mu, \sigma^2)$, then $c_\tau (\pi(x))^{1/\tau}$ density of $N(\mu, \tau\sigma^2)$.)
 - Then temperature acceptance probability is:

$$\min\left(1, \frac{\pi_{\tau'}(x)}{\pi_\tau(x)}\right) = \min\left(1, \frac{c_{\tau'}}{c_\tau} (\pi(x))^{(1/\tau') - (1/\tau)}\right).$$

- But this depends on the c_τ , which are usually unknown – bad.
- e.g. in above example, could try $\pi_\tau(x) = \left(\frac{1}{2}N(0, 1; x) + \frac{1}{2}N(20, 1; x)\right)^{1/\tau}$, but don't know normalising constants c_τ so not valid! (“Rtempered2”)
- What to do?

PARALLEL TEMPERING:

- (a.k.a. replica exchange: Swendsen and Wang, 1986)
- (a.k.a. Metropolis-Coupled MCMC, or MCMCMC: Geyer, 1991)
- Alternative to tempered MCMC.
- Again have a sequence $\Pi_1, \Pi_2, \dots, \Pi_m$ where $\Pi_1 = \Pi$ (“cold”), and Π_τ is flatter for larger τ (“hot”).
 - e.g. $\pi_\tau(x) = c_\tau (\pi(x))^{1/\tau}$, where τ ranges over $\tau_1 = 1, \tau_2, \tau_3, \dots, \tau_m$.
- Use state space \mathcal{X}^m , with m chains, i.e. one chain for each temperature.
 - So, state at time n is $X_n = (X_{n1}, X_{n2}, \dots, X_{nm})$, where $X_{n\tau}$ is “at temperature τ ”.
- Stationary distribution is now $\bar{\Pi} = \Pi_1 \times \Pi_2 \times \dots \times \Pi_m$, i.e. $\bar{\Pi}(X_1 \in S_1, X_2 \in S_2, \dots, X_m \in S_m) = \Pi_1(S_1) \Pi_2(S_2) \dots \Pi_m(S_m)$.
- Then, can update the chain $X_{n-1, \tau}$ at temperature τ (for each $1 \leq \tau \leq m$), by proposing e.g. $Y_{n, \tau} \sim N(X_{n-1, \tau}, 1)$, and accepting with probability $\min\left(1, \frac{\pi_\tau(Y_{n, \tau})}{\pi_\tau(X_{n-1, \tau})}\right)$.
- Or, can also choose temperatures τ and τ' (e.g., at random), and propose to “swap” the values $X_{n, \tau}$ and $X_{n, \tau'}$, and accept this with probability $\min\left(1, \frac{\pi_\tau(X_{n, \tau'}) \pi_{\tau'}(X_{n, \tau})}{\pi_\tau(X_{n, \tau}) \pi_{\tau'}(X_{n, \tau'})}\right)$.
 - Now, normalising constants cancel, e.g. if $\pi_\tau(x) = c_\tau (\pi(x))^{1/\tau}$, then

acceptance probability is:

$$\min \left(1, \frac{c_\tau \pi(X_{n,\tau'})^{1/\tau} c_{\tau'} \pi(X_{n,\tau})^{1/\tau'}}{c_\tau \pi(X_{n,\tau})^{1/\tau} c_{\tau'} \pi(X_{n,\tau'})^{1/\tau'}} \right) = \min \left(1, \frac{\pi(X_{n,\tau'})^{1/\tau} \pi(X_{n,\tau})^{1/\tau'}}{\pi(X_{n,\tau})^{1/\tau} \pi(X_{n,\tau'})^{1/\tau'}} \right),$$

so c_τ and $c_{\tau'}$ are not required.

- Hence, can set $g_\tau(x) = \pi(x)^{1/\tau}$, no problem.
- EXAMPLE: again $\Pi = \frac{1}{2} N(0, 1) + \frac{1}{2} N(20, 1)$.
 - Now can set $g_\tau(x) = \pi(x)^{1/\tau}$, and ignore c_τ .
 - Then run parallel tempering ... works pretty well. (file “Rpara”)

END WEEK #8

MONTE CARLO OPTIMISATION – Simulated Annealing:

- General method to find highest mode of π .
- Idea: mode of π is same as mode of a flatter or a more peaked version π_τ , for any $\tau > 0$.
 - e.g. $\pi_\tau \equiv \pi^{1/\tau}$. Flatter if $\tau > 1$, more peaked if $\tau < 1$. (“tempered”)
 - For large τ , MCMC explores a lot; good at beginning of search.
 - For small τ , MCMC narrows in on local mode; good at end of search.
- So, use tempered MCMC, but where $\tau = \tau_n \searrow 0$, so π_{τ_n} becomes more and more concentrated at mode as $n \rightarrow \infty$.
- Need to choose $\{\tau_n\}$, the “cooling schedule”.
 - e.g. geometric ($\tau_n = \tau_0 r^n$ for some $r < 1$).
 - or linear ($\tau_n = \tau_0 - dn$ for some $d > 0$, chosen so $\tau_M = \tau_0 - dM \geq 0$).
 - or logarithmic ($\tau_n = \tau_0 / \log(1 + n)$).
 - or ...
 - Theorem:: if $c \geq \sup \pi$, then simulated annealing with $\tau_n = c / \log(1 + n)$ will converge to the global maximum as $n \rightarrow \infty$. (But very slow.)
- EXAMPLE: $\Pi_\tau = 0.3 N(0, \tau^2) + 0.7 N(20, \tau^2)$. (file “Rsimann”)
 - Highest mode is at 20 (for any τ).
 - If run usual Metropolis algorithm, it will either jump forever between modes (if τ large), or get stuck in one mode or the other with equal probability (if τ small) – bad.
 - But if $\tau_n \searrow 0$ slowly, then can usually find the highest mode (20) – good.
 - Try both geometric and linear (better?) cooling ... (file “Rsimann”)

- EXAMPLE with real density powers:
 - Set $\pi_\tau(x) = (0.3 N(0, 1) + 0.7 N(20, 1))^{1/\tau}$.
 - Need longer run, and smaller final τ .
 - Then it works pretty well. (file “Rsimann2”)

DIGRESSION – CODE BREAKING:

- Try it out: “decipherdemo”. [uses file “decipher.c”]
- Data is the coded message text: $s_1 s_2 s_3 \dots s_N$, where $s_i \in \mathcal{A} = \{A, B, C, \dots, Z, \text{space}\}$.
- State space \mathcal{X} is set of all bijections (for now) of \mathcal{A} , i.e. one-to-one onto mappings $f : \mathcal{A} \rightarrow \mathcal{A}$, subject to $f(\text{space}) = \text{space}$.
 - [“substitution cipher”]
- Use a reference text (e.g. “War and Peace”) to get matrix $M(x, y) = 1 + \text{number of times } y \text{ follows } x$, for $x, y \in \mathcal{A}$.
- Then for $f \in \mathcal{X}$, let $\pi(f) = \prod_{i=1}^{N-1} M(f(s_i), f(s_{i+1}))$.
 - (Or raise this all to a power, e.g. 0.25.)
- Idea: if $\pi(f)$ is larger, then f leads to pair frequencies which more closely match the reference text, so f is a “better” choice.
- Would like to find the choice of f which maximises $\pi(f)$.
- To do this, run a “Metropolis algorithm” for π :
 - Choose $a, b \in \mathcal{A} \setminus \{\text{space}\}$, uniformly at random.
 - Propose to replace f by g , where $g(a) = f(b)$, $g(b) = f(a)$, and $g(x) = f(x)$ for all $x \neq a, b$.
 - Accept with probability $\min\left(1, \frac{\pi(g)}{\pi(f)}\right)$.
- Easily seen to be an irreducible, aperiodic, reversible Markov chain.
- So, converges (quickly!) to correct answer, breaking the code.
- References: S. Conner (2003), “Simulation and solving substitution codes”. P. Diaconis (2008), “The Markov Chain Monte Carlo Revolution”.
- We later extended this, to transposition ciphers and more: J. Chen and J.S. Rosenthal (2010), “[Decrypting Classical Cipher Text Using Markov Chain Monte Carlo](#)” (*Statistics and Computing* **22**(2), 397–413, 2011).

DIGRESSION – PATTERN DETECTION:

- Data is an image, given in terms of a grid of pixels (each “on” or “off”).
- Want to “find” the face in the image.

- (Harder for computers than for humans!)
- Java applet: [faces.html](#) (See [before](#) and [after](#) images.)
- Define the face location by a vector θ of various parameters (face center, eye width, nose height, etc.).
- Then define a score function $S(\theta)$ indicating how well the image agrees with having a face in the location corresponding to the parameters θ .
- Then run a “mixed” Monte Carlo search (sometimes updating by small RWM moves, sometimes starting fresh from a random vector) over the entire parameter space, searching for $\operatorname{argmax}_{\theta} S(\theta)$, i.e. for the parameter values which maximise the score function.
 - Keep track of the best θ so far – this allows for greater flexibility in trying different search moves without needing to preserve a stationary distribution.
 - Works pretty well, and fast! (“[faces.html](#)” Java applet)
 - For details, see Java applet source code file “[faces.java](#)”, or the paper [J.S. Rosenthal, Optimising Monte Carlo Search Strategies for Automated Pattern Detection. F. E. J. Math. Sci. 2009.](#)
- Here, again, we want to maximise (i.e., optimise) π , not sample from π .

MCMC CONVERGENCE RATES THEORY:

- $\{X_n\}$: Markov chain on \mathcal{X} , with stationary distribution $\Pi(\cdot)$.
- Let $P^n(x, S) = \mathbf{P}[X_n \in S \mid X_0 = x]$ be the probabilities for the Markov chain after n steps, when started at x .
 - Hope that for large n , $P^n(x, S) \approx \Pi(S)$.
- Let $D(x, n) = \|P^n(x, \cdot) - \Pi(\cdot)\| \equiv \sup_{S \subseteq \mathcal{X}} |P^n(x, S) - \Pi(S)|$.
- DEFN: chain is ergodic if $\lim_{n \rightarrow \infty} D(x, n) = 0$, for Π -a.e. $x \in \mathcal{X}$.
- Theorem (mentioned earlier): if chain is irreducible and aperiodic and $\Pi(\cdot)$ stationary, then chain is ergodic, i.e. converges asymptotically to Π .
- DEFN: chain is geometrically ergodic if there is $\rho < 1$, and $M : \mathcal{X} \rightarrow [0, \infty]$ which is Π -a.e. finite, such that $D(x, n) \leq M(x) \rho^n$ for all $x \in \mathcal{X}$ and $n \in \mathbf{N}$.
- Fact (mentioned earlier): CLT holds for $\frac{1}{n} \sum_{i=1}^n h(X_i)$ if chain is geometrically ergodic and $\mathbf{E}_{\pi}(|h|^{2+\delta}) < \infty$ for some $\delta > 0$.
 - (If chain also reversible then don’t need δ : [Roberts and Rosenthal, “Geometric ergodicity and hybrid Markov chains”, ECP 1997.](#))
 - If CLT holds, then (as before) have 95% confidence interval $(e - 1.96 \sqrt{v}, e + 1.96 \sqrt{v})$, where $v \approx \frac{1}{M-B} \operatorname{Var}_{\pi}(h)(\text{varfact})$.

- DEFN: chain is uniformly ergodic if there is $\rho < 1$, and $M < \infty$ such that $D(x, n) \leq M \rho^n$ for all $x \in \mathcal{X}$ and $n \in \mathbf{N}$.
- DEFN: a quantitative bound on convergence is an actual number n^* such that $D(x, n^*) < 0.01$ (say). [Then sometimes say chain “converges in n^* iterations”.]
- Quantitative bounds often difficult (though I’ve worked on them a lot, see e.g. [Rosenthal, “Quantitative convergence rates of Markov chains: A simple account”, Elec Comm Prob 2002](#) and the references therein), but “geometric ergodicity” is often easier to verify.
- But what about convergence rates, i.e. bounds on $D(x, n)$?
- Useful Theorem: If $P(x, dy) \geq \delta \pi(dy)$ for all $x, y \in \mathcal{X}$, then $D(x, n) \leq (1 - \delta)^n$. (“minorisation condition” ... proof by “coupling” ...)
 - If true, shows uniform ergodicity and can give a quantitative bound.
- Special Case: INDEPENDENCE SAMPLER (mentioned earlier):
 - Proposals $\{Y_n\}$ i.i.d. from some fixed distribution (say, $Y_n \sim MVN(0, I)$).
 - Another special case of Metropolis-Hastings algorithm, where $q(x, y) = q(y)$ depends only on y . So, Π is a stationary distribution.
 - By above Theorem, independence sampler is ergodic provided $q(x) > 0$ whenever $\pi(x) > 0$ (since then it must be irreducible and aperiodic).
 - But does that guarantee that it will work well?
 - No, e.g. previous “Rind” example with $k = 5$: ergodic (of course), but performs terribly.
 - FACT: independence sampler is geometrically ergodic IF AND ONLY IF there is $\delta > 0$ such that $q(x) \geq \delta \pi(x)$ for π -a.e. $x \in \mathcal{X}$.
 - If so, then we compute that

$$\begin{aligned}
 p(x, y) &\geq q(y) \min\left[1, \frac{\pi(y) q(x)}{\pi(x) q(y)}\right] = \min[q(y), \pi(y) (q(x) / \pi(x))] \\
 &\geq \min[\delta \pi(y), \delta \pi(y)] = \delta \pi(y).
 \end{aligned}$$

Hence, from the above, $D(x, n) \leq (1 - \delta)^n$ for π -a.e. $x \in \mathcal{X}$.

- PREVIOUS EXAMPLE: Independence sampler with $\pi(x) = e^{-x}$ and $q(x) = k e^{-kx}$ for $x > 0$. (file “Rind”)
 - If $0 < k \leq 1$, then setting $\delta = k$, we have that $q(x) = k e^{-kx} \geq k e^{-x} = k \pi(x) = \delta \pi(x)$ for all $x > 0$, so it’s geometrically ergodic, and furthermore $D(x, n) \leq (1 - k)^n$.
 - e.g. if $k = 0.01$, then $D(x, 459) \leq (0.99)^{459} \doteq 0.0099 < 0.01$ for all $x > 0$, i.e. “converges after 459 iterations”.

- But if $k > 1$, then cannot find any $\delta > 0$ such that $q(x) \geq \delta\pi(x)$ for all x , so it is not geometrically ergodic.
- If $k > 2$, then no CLT (Roberts, J. Appl. Prob. **36**, 1210–1217, 1999).
- So, if $k = 5$ (as in “Rind”), then not geometrically ergodic, and CLT does not hold. Indeed, confidence intervals often miss 1. (file “Rind2”)
- Fact: if $k = 5$, then $D(0, n) > 0.01$ for all $n \leq 4,000,000$, while $D(0, n) < 0.01$ for all $n \geq 14,000,000$, i.e. “convergence” takes between 4 million and 14 million iterations. Slow! [Roberts and Rosenthal, “Quantitative Non-Geometric Convergence Bounds for Independence Samplers”, MCAP 2011.]
- What about other MCMC algorithms (besides independence sampler)?
- FACT: if state space is finite, and chain is irreducible and aperiodic, then always ergodic (of course) and also geometrically ergodic. (See e.g. J.S. Rosenthal, SIAM Review 37:387-405, 1995.)
- What about for the “random-walk Metropolis algorithm” (RWM), i.e. where $\{Y_n - X_{n-1}\} \sim q$ (i.i.d.) for some fixed symmetric density q ?
 - e.g. $Y_n \sim N(X_{n-1}, \sigma^2 I)$, or $Y_n \sim \text{Uniform}[X_{n-1} - \delta, X_{n-1} + \delta]$.
- FACT: RWM is geometrically ergodic essentially if and only if π has exponentially light tails, i.e. there are $a, b, c > 0$ such that $\pi(x) \leq ae^{-b|x|}$ whenever $|x| > c$. (Requires a few technical conditions: π and q continuous and positive; q has finite first moment; and π non-increasing in the tails, with (in higher dims) bounded Gaussian curvature.) [Mengersen and Tweedie, Ann Stat 1996; Roberts and Tweedie, Biometrika 1996]

END WEEK #9

- [11 STUDENT PRESENTATIONS!]
- EXAMPLES: RWM on \mathbf{R} with usual proposals: $Y_n \sim N(X_{n-1}, \sigma^2)$:
- CASE #1: $\Pi = N(5, 4^2)$, and functional $h(y) = y^2$, so $\mathbf{E}_\pi(h) = 5^2 + 4^2 = 41$. (file “Rnorm” ... $\sigma = 1$ v. $\sigma = 4$ v. $\sigma = 16$)
 - Geometrically ergodic? Yes! (By above.)
 - Does CLT hold? Yes! (geometrically ergodic, and $\mathbf{E}_\pi(|h|^p) < \infty$ for all p .)
 - Indeed, confidence intervals “usually” contain 41. (file “Rnorm2”)
- CASE #2: $\pi(y) = c \frac{1}{(1+y^4)}$, and functional $h(y) = y^2$, so

$$\mathbf{E}_\pi(h) = \frac{\int_{-\infty}^{\infty} y^2 \frac{1}{(1+y^4)} dy}{\int_{-\infty}^{\infty} \frac{1}{(1+y^4)} dy} = \frac{\pi/\sqrt{2}}{\pi/\sqrt{2}} = 1.$$

- Not exponentially light tails, so not geometrically ergodic; estimates less stable, confidence intervals often miss 1. (file “Rheavy”)
- CASE #3: $\pi(y) = \frac{1}{\pi(1+y^2)}$ (Cauchy), and functional $h(y) = \mathbf{1}_{-10 < y < 10}$.
 - Recall that for Cauchy, $\Pi(0 < X < y) = \arctan(y)/\pi$.
 - So, $\mathbf{E}_\pi(h) = \Pi(|X| < 10) = 2 \arctan(10)/\pi = 0.93655$.
 - Again, not exponentially light tails, so not geometrically ergodic.
 - Confidence intervals often miss 0.93655. (file “Rcauchy”)
- CASE #4: $\pi(y) = \frac{1}{\pi(1+y^2)}$ (Cauchy), and functional $h(y) = \min(y^2, 100)$.
 - [Numerical integration: $\mathbf{E}_\pi(h) \doteq 11.77$]
 - Once again, not exponentially light tails, so not geometrically ergodic.
 - And, 95% CI often miss 11.77, though iid MC does better. (“Rcauchy2”)
- **NOTE:** Even when CLT holds, it can be rather unstable, e.g. it requires that chain has converged to Π , so it might underestimate v .
 - Estimate of v is very important! And “varfact” is not always reliable!
 - Repeated runs?
 - Another approach is “batch means”, whereby chain is broken into m large “batches”, which are assumed to be approximately i.i.d.

OPTIMAL RWM PROPOSAL SHAPE:

- Consider RWM on $\mathcal{X} = \mathbf{R}^d$, where $Y_n \sim MVN(X_{n-1}, \Sigma)$ for some $d \times d$ proposal covariance matrix Σ .
- What is best choice of Σ ?
 - Usually we take $\Sigma = \sigma^2 I_d$ for some $\sigma > 0$, and then choose σ so acceptance rate not too small, not too large (e.g. 0.234).
 - But can we do better?
- Suppose for now that $\Pi = MVN(\mu_0, \Sigma_0)$ for some fixed μ_0 and Σ_0 , in $\text{dim}=5$. Try RWM with various proposal distributions (file “Ropt”):
 - first version: $Y_n \sim MVN(X_{n-1}, I_d)$. ($\text{acc} \approx 0.06$; $\text{varfact} \approx 220$)
 - second version: $Y_n \sim MVN(X_{n-1}, 0.1 I_d)$. ($\text{acc} \approx 0.234$; $\text{varfact} \approx 300$)
 - third version: $Y_n \sim MVN(X_{n-1}, \Sigma_0)$. ($\text{acc} \approx 0.31$; $\text{varfact} \approx 15$)
 - fourth version: $Y_n \sim MVN(X_{n-1}, 1.4 \Sigma_0)$. ($\text{acc} \approx 0.234$; $\text{varfact} \approx 7$)
- Or in $\text{dim}=20$ (file “Ropt2”, with file “Rtarg20”):
 - $Y_n \sim MVN(X_{n-1}, 0.025 I_d)$. ($\text{acc} \approx 0.234$; $\text{varfact} \approx 400$ or more)
 - $Y_n \sim MVN(X_{n-1}, 0.283 \Sigma_0)$. ($\text{acc} \approx 0.234$; $\text{varfact} \approx 50$)

- Conclusion: acceptance rates near 0.234 are better.
- But also, proposals shaped like the target are better.
 - Indeed, best is when proposal covariance = $((2.38)^2/d)\Sigma_0$.
 - This has been proved for targets which are orthogonal transformations of independent components (Roberts et al., Ann Appl Prob 1997; [Roberts and Rosenthal, Stat Sci 2001](#) ; [Bédard, Ann Appl Prob 2007](#)).
 - And it’s “approximately” true for most unimodal targets . . .
- Problem: Σ_0 would usually be unknown; then what?
 - Can perhaps “adapt“!

ADAPTIVE MCMC:

- Recall: RWM optimal proposal covariance is $((2.38)^2/d)\Sigma_0$.
- What if target covariance Σ_0 is unknown??
- Can estimate Σ_0 based on run so far, to get empirical covariance Σ_n .
- Then update proposal covariance “on the fly”.
- “Learn as you go”: see e.g. the [Javascript](#) simulation.
- For Adaptive MCMC, could use proposal $Y_n \sim MVN(X_{n-1}, ((2.38)^2/d)\Sigma_n)$.
 - Hope that for large n , $\Sigma_n \approx \Sigma_0$, so proposals “nearly” optimal.
 - (Usually also add ϵI_d to proposal covariance, to improve stability, e.g. $\epsilon = 0.05$.)
- Try R version, for the same MVN example as in Ropt (file “Radapt”):
 - Need much longer burn-in, e.g. $B = 20,000$, for adaption to work.
 - Get varfact of last 4000 iterations of about 18 . . . “competitive” with Ropt optimal . . .
 - The longer the run, the more benefit from adaptation.
 - Can also compute “slow-down factor”, $s_n \equiv d \left(\sum_{i=1}^d \lambda_{in}^{-2} / (\sum_{i=1}^d \lambda_{in}^{-1})^2 \right)$, where $\{\lambda_{in}\}$ eigenvals of $\Sigma_n^{1/2} \Sigma_0^{-1/2}$. Starts large, should converge to 1. [Motivation: if $\Sigma_n = \Sigma_0$, then $\lambda_{in} \equiv 1$, so $s_n = d(d/d^2) \equiv 1$.] See [Roberts and Rosenthal, Examples of Adaptive MCMC, JCGS 2009](#).
- Higher dimensions: figure “[RplotAMx200.png](#)” (dim=200). (beautiful!)
 - Works well, but it takes many iterations before the adaption is helpful.