

STA4502 (Monte Carlo Estimation) Lecture Notes, Jan–Feb 2016

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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 for the test, or reading the reference books.

INTRODUCTION:

- Introduction to course, handout, references, prerequisites, etc.
 - Course web page: www.probability.ca/sta4502
 - Six weeks only; counts for QUARTER-credit only.
 - Sidney Smith Hall room 1080, Wednesdays 11:10–1:00, and Fridays 11:10–12:00.
 - If not Stat Dept grad student, must REQUEST enrolment (by e-mail); need advanced undergraduate probability/statistics background, plus basic computer programming experience.
 - 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 stat grad students? undergrads? math? computer science? physics? economics? management? engineering? other? Auditing??
- Theme of the course: use (pseudo)randomness on a computer to simulate (and hence estimate) important/interesting quantities.
- Example: Suppose want to estimate $m := \mathbf{E}[Z^4 \cos(Z)]$, where $Z \sim \text{Normal}(0, 1)$.
 - 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 (unbiased) 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”]
 - unstable ... but if replace “100” with “1000000” then \bar{x} close to -1.213 ...
 - Variability??
 - Well, can estimate standard deviation of \bar{x} by “standard error” of \bar{x} ,

which is:

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

[file “RMCse”]

- Then what is, say, a 95% confidence interval for m ?
- Well, by central limit theorem (CLT), for large n , 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]

- What about higher-dimensional versions? (Can’t do numerical integration!)
- Note: In this course we will just use R to automatically sample from simple distributions like Normal, Uniform, Exponential, etc.
 - (How does it work? Discussed in e.g. Statistical Computing course.)
- What if distribution too complicated to sample from?
 - (MCMC! ... including Metropolis, Gibbs, tempered, trans-dimensional, ...)

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? [<http://www.metablake.com/pi.swf>]
- Let θ be angle counter-clockwise from line direction, and h distance of top end above nearest line.
- Then $h \sim \text{Uniform}[0, w]$ and $\theta \sim \text{Uniform}[0, \pi]$.

- Touches line iff $h < \ell \sin(\theta)$.
- So, $\text{prob} = \frac{1}{\pi} \int_0^\pi \frac{1}{w} \int_0^w \mathbf{1}_{h < \ell \sin(\theta)} dh d\theta = \frac{1}{\pi} \int_0^\pi \frac{1}{w} \ell \sin(\theta) d\theta = 2\ell/w\pi$.
- 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 $\pi \approx 2n\ell/mw = 2n/m$ (if $\ell = w$).
- [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.
 - many statistical tests, to try to see which choices provide good randomness, avoid correlations, etc. (e.g. “diehard tests”, <http://stat.fsu.edu/pub/diehard/>; “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) $\text{gcd}(b, m) = 1$, 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, then the LCG has full period $m = 2^{32} \doteq 4.3 \times 10^9$; good.
 - Many other choices, e.g. C programming language (glibc) uses $m = 2^{32}$, $a = 1,103,515,245$, $b = 12,345$.
 - One bad choice: $m = 2^{31}$, $a = 65539 = 2^{16} + 3$, $b = 0$ (“RANDU”) ... used for many years (esp. early 1970s) ... but then $x_{n+2} = 6x_{n+1} - 9x_n$

mod m ... too much serial correlation. [Proof: $x_{n+2} = (2^{16} + 3)^2 x_n = (2^{32} + 6(2^{16}) + 9)x_n \equiv (0 + 6(2^{16} + 3) - 9)x_n \pmod{2^{31}} = 6x_{n+1} - 9x_n.$]

- (Microsoft Excel pre-2003: period $< 10^6$, too small ... Excel 2003 used floating-point “version” of LCG, which sometimes gave negative numbers – bad!)
- Not “really” random, just “pseudorandom” ...
 - Can cause problems!
 - Will fail certain statistical tests ...
 - Some implementations also use external randomness, e.g. current 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
 - Or of atmospheric noise, e.g. random.org.
 - But for most purposes, standard pseudorandom numbers are pretty good ...
- We’ll consider LCG’s “good enough for now”, but:
 - Other generators include “Multiply-with-Carry” [$x_n = (ax_{n-r} + b_{n-1}) \pmod{m}$ where $b_n = \lfloor (ax_{n-r} + b_{n-1})/m \rfloor$]; and ‘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 $K_{n+1} = (I + L^{13})(I + R^{18})K_n \pmod{2^{31}}$]; and “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, with matrix mult. done bit-wise mod 2], and many others too.
 - (R implementation: see “?.Random.seed” ... 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:
 - $> 2 + 1 - 2$
[1] 1
 - $> 2^{10} + 1 - 2^{10}$
 - $> 2^{100} + 1 - 2^{100}$

- Why??
- Homework question: for what values of n does:

$> 2\wedge n + 1 - 2\wedge n$
 give 0 instead of 1??

- So, numerical computations are just approximations, with their own errors!
- We'll usually ignore this, but **MUST BE CAREFUL!** Then can simulate ...

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 “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 $Y = X_1 + \dots + X_n$ 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”)

- 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 $\sum_i p_i = 1$, simply set

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

- e.g. to make $Z \sim$ Exponential(1), set $Z = -\log(U_1)$.
 - Then $\mathbf{P}(Z > x) = \mathbf{P}(-\log(U_1) > x) = \mathbf{P}(\log(U_1) < -x) = \mathbf{P}(U_1 < e^{-x}) = e^{-x}$.
 - Then, to make $W \sim$ Exponential(λ), set $W = Z/\lambda = -\log(U_1)/\lambda$.
- What if 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, $f_X(x) = \frac{d}{dx}x^6 = 6x^5$ for $0 < x < 1$.
- More generally, for $r > 1$, if $X = U_1^{1/r}$, then $f_X(x) = r x^{r-1}$ for $0 < x < 1$.
- 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 $X, Y \sim N(0, 1)$ (independent!). [“Box-Muller transformation”: Ann Math Stat 1958, 29, 610-611]

- Proof: By multidimensional change-of-variable theorem, if $(x, y) = h(u_1, u_2)$ and $(u_1, u_2) = h^{-1}(x, y)$, 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

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

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.

- Another approach: “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 $U_1 \leq F(x)$.
 - So, $\mathbf{P}(X \leq x) = \mathbf{P}(U_1 \leq F(x)) = F(x)$.
 - Very general, but computing $F^{-1}(t)$ could be difficult ...
- So, generating (pseudo)random numbers for most “standard” one-dimensional distributions is pretty easy ...
 - So, can get Monte Carlo estimates of expectations involving standard one-dimensional distributions, e.g. $\mathbf{E}[Z^4 \cos(Z)]$ where $Z \sim \text{Normal}(0, 1)$.
- But what if distribution is complicated, multidimensional, etc.? Simulate!

SIMULATION EXAMPLE: QUEUEING THEORY:

- $Q(t)$ = number of people in queue at time $t \geq 0$.
- Suppose service times \sim Exponential(μ) [mean $1/\mu$], and interarrival times \sim Exponential(λ) (“M/M/1 queue”), so $\{Q(t)\}$ Markovian. Then well known:
 - 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 Uniform[0, 1], and interarrival times have distribution of $|Z|$ where $Z \sim$ 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 Uniform[0, 2/3], and interarrival times have distribution of $Z^2/2$ where $Z \sim$ Normal(0, 1). Now what? [file “Rqueue3”]

END WEDNESDAY #1

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 Uniform[0, 1].
 - e.g. $g(x, y) = \cos(\sqrt{xy})$. (file “RMCint”) Easy!
 - Get about $0.88 \pm 0.003 \dots$ 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 $X \sim$ Uniform[0, 5] and $Y \sim$ Uniform[0, 4].
 - So, let $X_i \sim$ Uniform[0, 5], and $Y_i \sim$ 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 $\int_0^1 \int_0^\infty (e^y h(x, y)) e^{-y} dy dx$.
 - 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 ... (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 $I \equiv \mathbf{E}[h(Y)] = \int h(y) \pi(y) dy$, where Y has density π , could instead re-write this as $I = \int h(x) \frac{\pi(x)}{f(x)} f(x) dx$, where f is easily sampled from, with $f(x) > 0$ whenever $\pi(x) > 0$.
 - Then $I = \mathbf{E}\left(h(X) \frac{\pi(X)}{f(X)}\right)$, where X has density f . ("Importance Sampling")
 - Can then do classical (iid) Monte Carlo integration, get standard errors etc.
 - Good if easier to sample from f than π , and/or if the function $h(x) \frac{\pi(x)}{f(x)}$ is less variable than h itself.
- In general, best to make $h(x) \frac{\pi(x)}{f(x)}$ approximately constant.
 - 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).

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, $I = \int h(x) \pi(x) dx = \int h(x) c g(x) dx = \frac{\int h(x) g(x) dx}{\int g(x) dx}$.

- If sample $\{x_i\} \sim f$ (i.i.d.), then $\int h(x)g(x)dx = \int (h(x)g(x)/f(x))f(x)dx = \mathbf{E}[h(X)g(X)/f(X)]$ where $X \sim f$.
- So, $\int h(x)g(x)dx \approx \frac{1}{M} \sum_{i=1}^M (h(x_i)g(x_i)/f(x_i))$.
- Similarly, $\int g(x)dx \approx \frac{1}{M} \sum_{i=1}^M (g(x_i)/f(x_i))$.
- So, $I \approx \frac{\sum_{i=1}^M (h(x_i)g(x_i)/f(x_i))}{\sum_{i=1}^M (g(x_i)/f(x_i))}$. (“Importance Sampling”: weighted average)
- (Not unbiased, standard errors less clear, but still consistent.)
- (Good to use same sample $\{x_i\}$ for both numerator and denominator: easier computationally, and smaller variance.)
- 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 $c > 0$ 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 $f(y) = 6 y^5 \mathbf{1}_{0 < y < 1}$. [Fact (check): if $U \sim \text{Uniform}[0, 1]$, then $X \equiv U^{1/6} \sim f$.]
 - Then $I \approx \frac{\sum_{i=1}^M (h(x_i)g(x_i)/f(x_i))}{\sum_{i=1}^M (g(x_i)/f(x_i))} = \frac{\sum_{i=1}^M (\sin(x_i^4) \cos(x_i^5))}{\sum_{i=1}^M (\sin(x_i^4) \cos(x_i^5) / x_i^2)}$. (file “Rimp” ... 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 “Rimp”)
- 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.
- 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 $U \leq g(X)/Kf(X)$, then accept X (as a draw from π).
- Otherwise, reject X and start over again.
- Now, $\mathbf{P}(U \leq g(X)/Kf(X) \mid X = x) = g(x)/Kf(x)$, so conditional on accepting, we have that

$$\begin{aligned} \mathbf{P}\left(X \leq y \mid U \leq \frac{g(X)}{Kf(X)}\right) &= \frac{\mathbf{P}\left(X \leq y, U \leq \frac{g(X)}{Kf(X)}\right)}{\mathbf{P}\left(U \leq \frac{g(X)}{Kf(X)}\right)} \\ &= \frac{\int_{-\infty}^y f(x) \frac{g(x)}{Kf(x)} dx}{\int_{-\infty}^{\infty} f(x) \frac{g(x)}{Kf(x)} dx} = \frac{\int_{-\infty}^y g(x) dx}{\int_{-\infty}^{\infty} g(x) dx} = \int_{-\infty}^y \pi(x) dx. \end{aligned}$$

- So, conditional on accepting, $X \sim \pi$. Good! iid!
- However, prob. of accepting may be very small, then get very few samples.
- 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. $h(x) = x^4$.
 - 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 $|x| \geq 2$, $Kf(x) = 8 \frac{1}{2} \exp(-|x|) \geq 8 \frac{1}{2} \exp(-x^2/2) \geq (2\pi)^{-1/2} \exp(-x^2/2) = \pi(x) = g(x)$.

END FRIDAY #1

- 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. (file “Rrej”)
- For Rejection Sampler, $P(\text{accept}) = \mathbf{E}[P(\text{accept} \mid X)] = \mathbf{E}\left[\frac{g(X)}{Kf(X)}\right] = \int \frac{g(x)}{Kf(x)} f(x) dx = \frac{1}{K} \int g(x) dx = \frac{1}{cK}$. (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 the 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? 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”)
- Can iid / importance / rejection / auxiliary sampling solve all 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.)
- Define a Markov chain (dependent random process) X_0, X_1, X_2, \dots in such a way that for large enough n , $X_n \approx \pi$.
- METROPOLIS ALGORITHM (1953):
 - Choose some initial value X_0 (perhaps random).
 - Then, given X_{n-1} , choose a proposal move $Y_n \sim \text{MVN}(X_{n-1}, \sigma^2 I)$ (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 multiplicative constants cancel.)
- Fact: Then, for large n , have $X_n \approx \pi$. (“[rwm.html](#)” Java applet)
- 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”) chosen large enough so $X_B \approx \pi$, and M chosen large enough to get good Monte Carlo estimates.

- Aside: if accepted all proposals, then would have a “random walk” Markov chain.
 - So, this is called the “random walk Metropolis” (RWM) algorithm.
- How large B ? Difficult to say! (Some theory ... active area of research [see e.g. [Rosenthal, “Quantitative convergence rates of Markov chains: A simple account”, Elec Comm Prob 2002](#), on instructor’s web page] ... usually use trial-and-error ...)
- What initial value X_0 ?
 - Virtually any one will do, but “central” ones best.
 - Ideal: “overdispersed starting distribution”, i.e. choose X_0 randomly from some distribution that “covers” the “important” part of the state space.
- 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. Important! (Soon.)
 - What if we change σ ? How does that affect estimate? plot? acf?
- 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 ... works ... gets between about 34 and 44 ... but large uncertainty ... (file “Rmet2”) (Mathematica gets 38.7044)

- Individual plots appear to have “good mixing” ...
- Joint plot shows fewer samples where $x_1 x_2 \approx (\pi/2)^2 \doteq 2.5$...
- e.g. if $\pi(x) = \exp\left(\sum_{i<j} |x_j - x_i|\right)$, then $\log(\pi(x)) = \sum_{i<j} |x_j - x_i|$.

END WEDNESDAY #2

- OPTIMAL SCALING:
 - Can change proposal distribution to $Y_n \sim MVN(X_{n-1}, \sigma^2 I)$ for any $\sigma > 0$.
 - 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 ... (“[rwm.html](#)” applet, files “Rmet”, “Rmet2”) ...
 - Some theory ... limited ... active area of research ...
 - General principle: the acceptance rate should be far from 0 and far from 1.
 - 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](#)]

MCMC STANDARD ERROR:

- What about standard error, i.e. uncertainty?
 - It’s usually larger than in iid case (due to 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 compute 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 $v \equiv \text{Var}\left(\frac{1}{M-B} \sum_{i=B+1}^M h(X_i)\right)$?
 - 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

$$\text{varfact} = 1 + 2 \sum_{k=1}^{\infty} \text{Corr}_\pi(h(X_0), h(X_k)) \equiv 1 + 2 \sum_{k=1}^{\infty} \rho_k = \sum_{k=-\infty}^{\infty} \rho_k$$

(“integrated auto-correlation time”).

- Also $\text{varfact} = 2 (\sum_{k=0}^{\infty} \rho_k) - 1$.
- Then can estimate both iid variance, and varfact, from the sample run, as usual.
- 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, 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 $\text{varfact} \gg 1$; try to get “better” chains so varfact smaller.
 - Sometimes even try to design chain to get $\text{varfact} < 1$ (“antithetic”).

END FRIDAY #2

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 ?

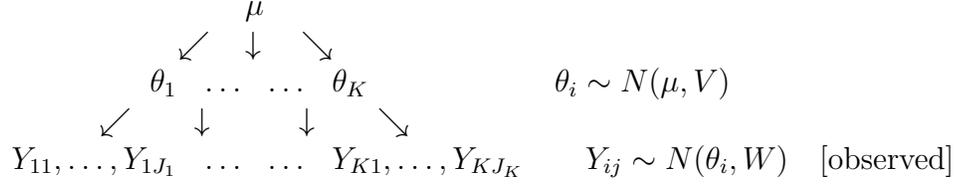
- Well, if have central limit theorem (CLT), then for large $M - B$, $e \approx N(u, v)$.
 - So $(e - u) v^{-1/2} \approx N(0, 1)$.
 - So, $\mathbf{P}(-1.96 < (e - u) v^{-1/2} < 1.96) \approx 0.95$.
 - So, $\mathbf{P}(-1.96 \sqrt{v} < e - u < 1.96 \sqrt{v}) \approx 0.95$.
 - i.e., with prob 95%, the interval $(e - 1.96 \sqrt{v}, e + 1.96 \sqrt{v})$ will contain u .
 - (Again, strictly speaking, should use “t” distribution, not 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.](#))
- So MCMC is more complicated than standard Monte Carlo.
 - Why should we bother?
 - Some problems too challenging for other methods. 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”):

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

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 $V, W > 0$):

$$\begin{aligned}
 & f(V, W, \mu, \theta_1, \dots, \theta_K, Y_{11}, Y_{12}, \dots, Y_{KJ_K}) \\
 &= C_1 \left(e^{-b_1/V} V^{-a_1-1} \right) \left(e^{-b_2/W} W^{-a_2-1} \right) \left(e^{-(\mu-a_3)^2/2b_3} \right) \times \\
 & \quad \times \left(\prod_{i=1}^K V^{-1/2} e^{-(\theta_i-\mu)^2/2V} \right) \left(\prod_{i=1}^K \prod_{j=1}^{J_i} W^{-1/2} e^{-(Y_{ij}-\theta_i)^2/2W} \right) \\
 &= C_2 e^{-b_1/V} V^{-a_1-1} e^{-b_2/W} W^{-a_2-1} e^{-(\mu-a_3)^2/2b_3} V^{-K/2} W^{-\frac{1}{2} \sum_{i=1}^K J_i} \times \\
 & \quad \times \exp \left[-\sum_{i=1}^K (\theta_i - \mu)^2/2V - \sum_{i=1}^K \sum_{j=1}^{J_i} (Y_{ij} - \theta_i)^2/2W \right].
 \end{aligned}$$

- Then

$$\begin{aligned}
 & \pi(V, W, \mu, \theta_1, \dots, \theta_K) \\
 &= C_3 \left(e^{-b_1/V} V^{-a_1-1} \right) \left(e^{-b_2/W} W^{-a_2-1} \right) \left(e^{-(\mu-a_3)^2/2b_3} \right) \times \\
 & \quad \times \left(\prod_{i=1}^K V^{-1/2} e^{-(\theta_i-\mu)^2/2V} \right) \left(\prod_{i=1}^K \prod_{j=1}^{J_i} W^{-1/2} e^{-(Y_{ij}-\theta_i)^2/2W} \right)
 \end{aligned}$$

- After a bit of simplifying,

$$\pi(V, W, \mu, \theta_1, \dots, \theta_K)$$

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

- Dimension: $d = K + 3$, e.g. $K = 19$, $d = 22$.
- How to compute/estimate, say, $\mathbf{E}_\pi(W/V)$, or effect from changing b_1 ?
 - Numerical integration? No, too high-dimensional!
 - Importance sampling? Perhaps, but what “ f ”? Not very efficient!
 - Rejection sampling? What “ f ”? What “ K ”? Virtually no samples!
 - Perhaps MCMC can work!
 - But need clever, useful MCMC algorithms!
- COMMENT: For big complicated π , often better to work with the LOG-ARITHMS, i.e. accept if $\log(U_n) < \log(A_n) = \log(\pi(Y_n)) - \log(\pi(X_{n-1}))$.
 - Then only need to compute $\log(\pi(x))$, which might work better.
 - So, better to program on log scale: $\log \pi(V, W, \mu, \theta_1, \dots, \theta_K) = \dots$
- NOTE: Many applications of this model, e.g.:
 - Predicting success at law school (D. Rubin, JASA 1980), $K = 82$ schools.
 - Melanoma recurrence (http://www.mssanz.org.au/modsim07/papers/52_s24/Analysing_Clinicals24_Bartolucci.pdf), $K = 19$ 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. (data in file “Rdye”)

METROPOLIS-HASTINGS ALGORITHM:

- (Hastings [Canadian!], Biometrika 1970; see www.probability.ca/hastings)
- Previous Metropolis algorithm works provided proposal distribution is symmetric, i.e. $q(x, y) = q(y, x)$. But what if it isn’t?
- FACT: 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 it’s still valid (justification later); everything else remains the same.
 - i.e., 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 $q(y, x) > 0$.

- **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 $\mathbf{E}_\pi(h) \doteq 38.7044$.)
 - 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, $q(x, y) = C (1 + |x|^2)^{-2} \exp(-|y - x|^2 / 2 \sigma^2 (1 + |x|^2))$.
 - So, can run Metropolis-Hastings algorithm for this example. (file “RMH”)
 - Usually get between 34 and 43, with claimed standard error ≈ 2 . (Recall: Mathematica gets 38.7044.)
- **LANGEVIN ALGORITHM:**
 - $Y_n \sim MVN(X_{n-1} + \frac{1}{2} \sigma^2 \nabla \log \pi(X_{n-1}), \sigma^2 I)$.
 - Special case of Metropolis-Hastings algorithm.
 - 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 $\nabla \log \pi$. (Hard. Homework!)
 - For theory, see e.g. Roberts & Tweedie, Bernoulli **2(4)**, 341–363, 1996; [Roberts & Rosenthal, JRSSB 60, 255–268, 1998](#).

INDEPENDENCE SAMPLER:

- Recall: with “random-walk Metropolis”, propose $Y_n \sim MVN(X_{n-1}, \sigma^2 I_d)$, then accept if $U_n < A_n$ where $U_n \sim \text{Uniform}[0, 1]$ and $A_n = \pi(Y_n) / \pi(X_{n-1})$.
- One alternative (of many – later) is the “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).
- **EXAMPLE:** independence sampler with $\pi(x) = e^{-x}$ and $q(y) = k e^{-ky}$.
 - Then if $X_{n-1} = x$ and $Y_n = y$, then $A_n = \frac{e^{-y} k e^{-kx}}{e^{-x} k e^{-ky}} = e^{(k-1)(y-x)}$. (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 ?

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 case: “Metropolis-within-Gibbs”) or Metropolis-Hastings rule (general case: “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 “Rmwg”) or random-scan (file “Rmwg2”).

END WEDNESDAY #3

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.
 - Then, always accept. (Reason later.)
 - Can use either systematic or random scan, just like above.
- 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 everything except μ treated as constant.
- Recall: full joint density is:

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

- So, conditional density of μ is

$$C_2 e^{-(\mu-a_3)^2/2b_3} \exp \left[- \sum_{i=1}^K (\theta_i - \mu)^2 / 2V \right].$$

- This equals (verify this! HW!)

$$C_3 \exp \left(- \mu^2 \left(\frac{1}{2b_3} + \frac{K}{2V} \right) + \mu \left(\frac{a_3}{b_3} + \frac{1}{V} \sum_{i=1}^K \theta_i \right) \right).$$

- 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: $v = b_3 V / (V + Kb_3)$, and $m = (a_3 V + b_3 \sum_{i=1}^K \theta_i) / (V + Kb_3)$.
- 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:

$$C_4 e^{-b_1/V} V^{-a_1-1} V^{-K/2} \exp \left[- \sum_{i=1}^K (\theta_i - \mu)^2 / 2V \right], \quad V > 0.$$

- 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).
- So, in this case, the systematic-scan Gibbs sampler 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!

MCMC CONVERGENCE RATES THEORY:

- $\{X_n\}$: Markov chain on \mathcal{X} , with stationary distribution $\Pi(\cdot)$.
- Let $P^n(x, S) = \mathbf{P}[X_n \in S \mid X_0 = x]$.
 - 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. $x \in \mathcal{X}$.
- 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 $n \in \mathbf{N}$.
- 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.
 - Fact: 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 $(e - 1.96 \sqrt{v}, e + 1.96 \sqrt{v})$.
- So what do we know about ergodicity?
- Theorem (later): if chain is irreducible and aperiodic and $\Pi(\cdot)$ is stationary, then chain is ergodic.
- What about convergence rates of independence sampler?
 - By Thm, independence sampler is ergodic provided $q(x) > 0$ whenever $\pi(x) > 0$.
 - But is that sufficient?
 - No, e.g. previous “Rind” example with $k = 5$: ergodic (of course), but not geometrically ergodic, CLT does not hold, confidence intervals

often miss 1. (file “Rind2”)

- 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. $x \in \mathcal{X}$, in which case $D(x, n) \leq (1 - \delta)^n$ for π -a.e. $x \in \mathcal{X}$.
 - So, if $\pi(x) = e^{-x}$ and $q(x) = ke^{-kx}$ for $x > 0$, where $0 < k \leq 1$, then can take $\delta = k$, so $D(x, n) \leq (1 - k)^n$.
 - 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 not geometrically ergodic.
 - If $k > 2$, then no CLT (Roberts, J. Appl. Prob. **36**, 1210–1217, 1999).
 - 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”, MCMC 2011.]

END FRIDAY #3

- What about other chains (besides independence sampler)?
- FACT: if state space is finite, and chain is irreducible and aperiodic, then always geometrically ergodic. (See e.g. J.S. Rosenthal, SIAM Review **37**:387-405, 1995.)
- What about for “random-walk Metropolis algorithm” (RWM), i.e. where $\{Y_n - X_{n-1}\} \sim q$ 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 exponential 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]
- 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. $\sigma = 16$)
 - 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 $h(y) = y^2$ (file “Rheavy”),

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 exponential tails, so no CLT; estimates less stable, confidence intervals often miss 1.
- CASE #3: $\pi(y) = \frac{1}{\pi(1+y^2)}$ (Cauchy), and functional $h(y) = \mathbf{1}_{-10 < y < 10}$, so $\mathbf{E}_\pi(h) = \mathbb{P}(|X| < 10) = 2 \arctan(10)/\pi = 0.93655$. [$\mathbb{P}(0 < X < x) = \arctan(x)/\pi$] (file “Rcauchy”)
- Not geometrically ergodic.
- Confidence intervals often miss 0.93655.
- CASE #4: $\pi(y) = \frac{1}{\pi(1+y^2)}$ (Cauchy), and functional $h(y) = \min(y^2, 100)$. [Numerical integration: $\mathbf{E}_\pi(h) \doteq 11.77$] (file “Rcauchy2”)
- Again, not geometrically ergodic, and 95% CI often miss 11.77, though iid MC does better.
- **NOTE:** Even when CLT holds, it’s rather unstable, e.g. requires that chain has converged to Π , and might underestimate v .
 - So, estimate of v is very important!
 - “varfact” 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.,

JUSTIFICATION: WHY DOES MCMC 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 $i, j \in \mathcal{X}$ 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 $P(X_n \in A | X_0 = x) > 0$.
- 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.]
 - 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 $Y_n = X_{n-1} \pm 1 \pmod{4}$. But almost always holds.
- What about Π being a stationary distribution?
- Begin with DISCRETE CASE (e.g. [rwm.html](#)).
- Assume for simplicity that $\pi(x) > 0$ for all $x \in \mathcal{X}$.
 - Let $q(x, y) = \mathbf{P}(Y_n = y | X_{n-1} = x)$ be proposal distribution, e.g. $q(x, x+1) = q(x, x-1) = 1/2$. Always chosen to be symmetric, i.e. $q(x, y) = q(y, x)$.
 - Let $\alpha(x, y)$ be probability of accepting a proposed move from x to y , i.e.
$$\alpha(x, y) = \mathbf{P}(U_n < A_n | X_{n-1} = x, Y_n = y) = \mathbf{P}(U_n < \frac{\pi(y)}{\pi(x)}) = \min[1, \frac{\pi(y)}{\pi(x)}].$$
 - State space is \mathcal{X} , e.g. $\mathcal{X} \equiv \{1, 2, 3, 4, 5, 6\}$.
- Then, for $i, j \in \mathcal{X}$ with $i \neq j$,

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

- Follows that chain is “(time) reversible”: for all $i, j \in \mathcal{X}$, by symmetry,
$$\pi(i) P(i, j) = q(i, j) \min(\pi(i), \pi(j)) = q(j, i) \min(\pi(i), \pi(j)) = \pi(j) P(j, i).$$
 - (Intuition: if $X_0 \sim \pi$, i.e. $\mathbf{P}(X_0 = i) = \pi(i)$ for all $i \in \mathcal{X}$, then $\mathbf{P}(X_0 = i, X_1 = j) = \pi(i) P(i, j) = \mathbf{P}(X_0 = j, X_1 = i) \dots$)
- We then compute that if $X_0 \sim \pi$, i.e. that $\mathbf{P}(X_0 = i) = \Pi(i)$ for all $i \in \mathcal{X}$, then:

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

$$= \pi(j) \sum_{i \in \mathcal{X}} P(j, i) = \pi(j),$$

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 $i \in \mathcal{X}$. (applet “[rwm.html](#)”)
- 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. for Variance Components Model, $\mathcal{X} = (0, \infty) \times (0, \infty) \times \mathbf{R} \times \mathbf{R}^K \subseteq \mathbf{R}^{K+3}$.)
 - Let $q(x, y)$ be the proposal density for y given x . (So, in above case, $q(x, y) = (2\pi\sigma)^{-d/2} \exp(-\sum_{i=1}^d (y_i - x_i)^2 / 2\sigma^2)$.) 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 $P(x, S) = \mathbf{P}(X_1 \in S | X_0 = x)$ be the transition probabilities.
- Then if $x \notin S$, then

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

- Shorthand: for $x \neq y$, $P(x, dy) = q(x, y) \min[1, \pi(y)/\pi(x)] dy$.
- Then for $x \neq y$, $P(x, dy) \pi(x) dx = q(x, y) \min[1, \pi(y)/\pi(x)] dy \pi(x) dx = q(x, y) \min[\pi(x), \pi(y)] dy dx = P(y, dx) \pi(y) dy$. (symmetric)
- Follows that $P(x, dy) \pi(x) dx = P(y, dx) \pi(y) dy$ for all $x, y \in \mathcal{X}$. (“reversible”)
- Shorthand: $P(x, dy) \Pi(dx) = P(y, dx) \Pi(dy)$.
- How does “reversible” help? Just like for discrete chains!
- Indeed, suppose $X_0 \sim \Pi$, i.e. we “start in stationarity”. Then

$$\begin{aligned} \mathbf{P}(X_1 \in S) &= \int_{x \in \mathcal{X}} \mathbf{P}(X_1 \in S | X_0 = x) \pi(x) dx = \int_{x \in \mathcal{X}} \int_{y \in S} P(x, dy) \pi(x) dx \\ &= \int_{x \in \mathcal{X}} \int_{y \in S} P(y, dx) \pi(y) dy = \int_{y \in S} \pi(y) dy \equiv \Pi(S), \end{aligned}$$

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

- So, again, the Theorem applies.
- 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) $Y_n = X_{n-1} \pm 1$ with prob. $\frac{1}{2}$ each, etc.

JUSTIFICATION OF METROPOLIS-HASTINGS:

- Now that we understand the theory, we can consider more general algorithms too . . .
- Previous Metropolis algorithm works provided proposal distribution is symmetric, i.e. $q(x, y) = q(y, x)$. But what if it isn't?
- For Metropolis, key was that $q(x, y) \alpha(x, y) \pi(x)$ was symmetric (to make the Markov chain be reversible).
- 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:

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

So, still symmetric, even if q wasn't.

- So, for Metropolis-Hastings algorithm, 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 still reversible, and everything else remains the same: still accept if $U_n < A_n$, otherwise reject.
- Do require that $q(x, y) > 0$ iff $q(y, x) > 0$.
- 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.
 - Then $q(x, y) = q(y)$, depends only on y .
 - So, now $A_n = \frac{\pi(Y_n) q(X_{n-1})}{\pi(X_{n-1}) q(Y_n)}$. (files “Rind”, “Rind2” from before)
- VARIABLE-AT-A-TIME: The exact same justification works if we update the variables one-at-a-time (e.g. Metropolis-within-Gibbs, Metropolis-Hastings-within-Gibbs, etc.); each individual step is still reversible (for the same reason), so π is still stationary.

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.
 - Here $C(x^{(-i)})$ is the appropriate normalising constant (which depends on $x^{(-i)}$). (So $C(x^{(-i)}) = C(y^{(-i)})$.)
 - 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).

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 $q(x, y)$ still symmetric.
 - π stationary? Yes since reversible!
 - Irreducible? Yes since $P^n(x, dy)$ has positive density whenever $|y -$

$$|x| \leq n.$$

- 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 a Metropolis-Hastings algorithm.)

END WEDNESDAY #4

- ASIDE: Why does Theorem say “ π -a.e.” $X_0 = x$?
- Example: $\mathcal{X} = \{1, 2, 3, \dots\}$, and $P(1, \{1\}) = 1$, and for $x \geq 2$, $P(x, \{1\}) = 1/x^2$ and $P(x, \{x+1\}) = 1 - (1/x^2)$.
 - Stationary distribution: $\Pi(\cdot) = \delta_1(\cdot)$, i.e. $\Pi(S) = \mathbf{1}_{1 \in S}$ for $S \subseteq \mathcal{X}$.
 - Irreducible, 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 since $P(1, \{1\}) > 0$.
 - So, by Theorem, for π -a.e. X_0 , have $\lim_{n \rightarrow \infty} \mathbf{P}(X_n \in S) = \Pi(S)$, i.e. $\lim_{n \rightarrow \infty} \mathbf{P}(X_n = 1) = 1$.
 - But 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”)

OPTIMAL RWM PROPOSALS:

- 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 “targ20”):
 - $Y_n \sim MVN(X_{n-1}, 0.025 I_d)$. ($\text{acc} \approx 0.234$; $\text{varfact} \approx 400$ 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 = $((2.38)^2/d)\Sigma_{\text{target}}$.
 - 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:

- What if target covariance Σ_0 is unknown??
- Can estimate target covariance 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 Java applet “[adapt.html](#)”
- For Adaptive MCMC, could use proposal $Y_n \sim MVN(X_{n-1}, \Sigma_n)$ [or $Y_n \sim MVN(X_{n-1}, 1.4\Sigma_n)$, or $Y_n \sim MVN(X_{n-1}, ((2.38)^2/d)\Sigma_n)$].

END FRIDAY #4

- 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”, $s_n \equiv d \left(\sum_{i=1}^d \lambda_{in}^{-2} / \left(\sum_{i=1}^d \lambda_{in}^{-1} \right)^2 \right)$, 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 “[plotAMx200.png](#)” (dim=200). (beautiful!)
 - Works well, but it takes many iterations before the adaption is helpful.
- BUT IS “ADAPTIVE MCMC” A VALID ALGORITHM??
- Not in general: see e.g. “[adapt.html](#)”
- Algorithm now non-Markovian, doesn’t preserve stationarity at each step.
- However, still guaranteed to converge to Π under various technical conditions.
- For example, it suffices (see [Roberts & Rosenthal, “Coupling and Convergence of Adaptive MCMC” \(J. Appl. Prob. 2007\)](#)) that the adaption satisfies:
 - (a) Diminishing Adaptation: Adapt less and less as the algorithm proceeds. Formally, $\sup_{x \in \mathcal{X}} \|P_{\Gamma_{n+1}}(x, \cdot) - P_{\Gamma_n}(x, \cdot)\| \rightarrow 0$ in prob. [Can always be made to hold, since adaption is user controlled.]
 - (b) Containment: For all $\epsilon > 0$, the time to converge to within ϵ of stationary from $x = X_n$, if fix $\gamma = \Gamma_n$, remain bounded in probability as $n \rightarrow \infty$. [Technical condition, to avoid “escape to infinity”. Holds if e.g. the state space and adaption spaces are both compact, etc. And always seems to hold in practice.]
 - (This also guarantees WLLN for bounded functionals. Various other results about LLN / CLT under stronger assumptions.)

- There are various “checkable” sufficient conditions which guarantee Containment, e.g. [Y. Bai, G.O. Roberts, and J.S. Rosenthal, Adv. Appl. Stat. 2011](#) and [Craiu, Gray, Latusynski, Madras, Roberts, and Rosenthal, Ann. Appl. Prob. 2015](#) and [J.S. Rosenthal and J. Yang, Ergodicity of Discontinuous Adaptive MCMC Algorithms, submitted.](#)
- So, some “reasonable” theory, but you have to be careful!

MONTE CARLO IN FINANCE [brief]:

- 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.
 - 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 $\Phi = \text{cdf of } N(0, 1)$:

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

- 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 $M = 1000$)

- Note that it's iid replications, so $\text{varfact} \equiv 1$.
- 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 there is no simple formula . . . but can still simulate! (file “RAO”)

MONTE CARLO OPTIMISATION Example – 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 irreducible, aperiodic, reversible.
- So, converges (quickly!) to correct answer, breaking the code.
- We later extended this to transposition ciphers etc.
- References: S. Conner (2003), “Simulation and solving substitution codes”. P. Diaconis (2008), “The Markov Chain Monte Carlo Revolution”. J. Chen and J.S. Rosenthal (2010), “[Decrypting Classical Cipher Text Using Markov Chain Monte Carlo](#)” (*Statistics and Computing* **22**(2), 397–413, 2011).

MONTE CARLO OPTIMISATION – Pattern Detection:

- Try it out: [faces.html](#)
- 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!)
- 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.](#)
- In both of these examples, wanted to MAXIMIZE (i.e., OPTIMIZE) π , rather than SAMPLE from π .
 - General method?

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 $\tau < 1$.
 - 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.

END WEDNESDAY #5

- 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”)

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 $\mathbf{E}_\pi(h)$.
- HOW TO FIND THE TEMPERED DENSITIES π_τ ?
- 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:

$$\min\left(1, \frac{\pi_{\tau'}(x)}{\pi_\tau(x)}\right) = \min\left(1, \frac{c_{\tau'}(\pi(x))^{(1/\tau')-(1/\tau)}}{c_\tau}\right).$$
 - This depends on the c_τ , which are usually unknown – bad.
- What to do?

PARALLEL TEMPERING:

- (a.k.a. Metropolis-Coupled MCMC, or MCMCMC)
- Alternative to tempered MCMC.
- Instead, 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 $\min\left(1, \frac{\pi_\tau(Y_{n,\tau})}{\pi_\tau(X_{n-1,\tau})}\right)$.

- And, 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.

- EXAMPLE: suppose again that $\Pi_\tau = \frac{1}{2} N(0, \tau^2) + \frac{1}{2} N(20, \tau^2)$, for $\tau = 1, 2, \dots, 10$.
 - Can run parallel tempering ... works pretty well. (file “Rpara”)

TRANSDIMENSIONAL MCMC [skip?]:

- (a.k.a. “reversible-jump MCMC”: Green, Biometrika 1995)
- What if the state space is a union of parts of different dimension?
 - Can we still apply Metropolis-Hastings then??
- EXAMPLE: autoregressive process: suppose $Y_n = a_1 Y_{n-1} + a_2 Y_{n-2} + \dots + a_k Y_{n-k}$, but we don’t know what k should be.

END FRIDAY #5

- EXAMPLE: suppose $\{y_j\}_{j=1}^J$ are known data which are assumed to come from a mixture distribution: $\frac{1}{k}(N(a_1, 1) + N(a_2, 1) + \dots + N(a_k, 1))$.
- Want to estimate the unknown k, a_1, \dots, a_k .
 - Here the number of parameters is also unknown, i.e. the dimension is unknown and variable, which makes MCMC more challenging!
- The state space is $\mathcal{X} = \{(k, a) : k \in \mathbf{N}, a \in \mathbf{R}^k\}$.
- Prior distributions: $k - 1 \sim \text{Poisson}(2)$, and $a|k \sim \text{MVN}(0, I_k)$ (say).
- Define a reference measure λ by: $\lambda(\{k\} \times A) = \lambda_k(A)$ for $k \in \mathbf{N}$ and (measurable) $A \subseteq \mathbf{R}^k$, where λ_k is Lebesgue measure on \mathbf{R}^k .
 - i.e., $\lambda = \delta_1 \times \lambda_1 + \delta_2 \times \lambda_2 + \delta_3 \times \lambda_3 + \dots$
- Then the posterior density (with respect to λ) is:

$$\pi(k, a) = C \frac{e^{-2} 2^{k-1}}{(k-1)!} (2\pi)^{-k/2} \exp\left(-\frac{1}{2} \sum_{i=1}^k a_i^2\right) (2\pi)^{-J/2} \prod_{j=1}^J \left(\sum_{i=1}^k \frac{1}{k} \exp\left(-\frac{1}{2}(y_j - a_i)^2\right)\right).$$

- So, on a log scale,

$$\log \pi(k, a) = \log C + \log \frac{e^{-2} 2^{k-1}}{(k-1)!} - \frac{k}{2} \log(2\pi) - \frac{1}{2} \sum_{i=1}^k a_i^2 - \frac{J}{2} \log(2\pi) + \sum_{j=1}^J \log \left(\sum_{i=1}^k \frac{1}{k} \exp \left(-\frac{1}{2} (y_j - a_i)^2 \right) \right).$$

(Can ignore $\log C$ and $\frac{J}{2} \log(2\pi)$, but not $\frac{k}{2} \log(2\pi)$.)

- How to “explore” this posterior distribution??
- For fixed k , can move around \mathbf{R}^k in usual way with RWM (say).
- But how to change k ?
- Can propose to replace k with, say, $k' = k \pm 1$ (prob $\frac{1}{2}$ each).
- Then have to correspondingly change a . One possibility:
 - If $k' = k + 1$, then $a' = (a_1, \dots, a_k, Z)$ where $Z \sim N(0, 1)$ (“elongate”).
 - If $k' = k - 1$, then $a' = (a_1, \dots, a_{k-1})$ (“truncate”).
- Then accept with usual probability, $\min \left(1, \frac{\pi(k', a') q((k', a'), (k, a))}{\pi(k, a) q((k, a), (k', a'))} \right)$.
 - Here if $k' = k + 1$, then $q((k', a'), (k, a)) = \frac{1}{2}$, while $q((k, a), (k', a')) = \frac{1}{2} \frac{1}{\sqrt{2\pi}} e^{-(a'_k)^2/2}$.
 - Or, if $k' = k - 1$, then $q((k, a), (k', a')) = \frac{1}{2}$, while $q((k', a'), (k, a)) = \frac{1}{2} \frac{1}{\sqrt{2\pi}} e^{-(a_k)^2/2}$.
- Seems to work okay; final k usually between 5 and 9 ... (file “Rtrans”)
- ALTERNATIVE method for the “correspondingly change a ” step:
 - If $k' = k + 1$, then $a' = (a_1, \dots, a_{k-1}, a_k - Z, a_k + Z)$ where $Z \sim N(0, 1)$ (“split”).
 - If $k' = k - 1$, then $a' = (a_1, \dots, a_{k-2}, \frac{1}{2}(a_{k-1} + a_k))$ (“merge”).
 - What about the densities $q((k', a'), (k, a))$?
 - Well, if $k' = k + 1$, then $q((k', a'), (k, a)) = \frac{1}{2}$, while roughly speaking,

$$q((k, a), (k', a')) = \frac{1}{2} \frac{1}{\sqrt{2\pi}} e^{-z^2/2} = \frac{1}{2} \frac{1}{\sqrt{2\pi}} e^{-(\frac{1}{2}(a'_k - a_k))^2/2}.$$

- One subtle additional point: The map $(a, Z) \mapsto a' = (a_1, \dots, a_{k-1}, a_k - Z, a_k + Z)$ has “Jacobian” term:

$$\det \left(\frac{\partial a'}{\partial (a, Z)} \right) = \det \begin{pmatrix} I_{k-1} & 0 & 0 \\ 0 & 1 & -1 \\ 0 & 1 & 1 \end{pmatrix} = 1 - (-1) = 2,$$

i.e. the split moves “spread out” the mass by a factor of 2.

- So by Change-of-Variable Thm, actually

$$q((k, a), (k', a')) = \frac{1}{2} \frac{1}{\sqrt{2\pi}} e^{-\frac{1}{2}(a' - a_k)^2/2} / 2.$$

- Similarly, if $k' = k - 1$, then $q((k, a), (k', a')) = \frac{1}{2}$, while

$$q((k', a'), (k, a)) = \frac{1}{2} \frac{1}{\sqrt{2\pi}} e^{-\frac{1}{2}(a_k - a_{k'})^2/2} / 2.$$

- Algorithm still seems to work okay ... (file “Rtrans2”)
 - For more complicated transformations, need to include more complicated “Jacobian” term (but above it equals 1 or 2).
 - Check: if we start the algorithms with, say, $k = 24$, then they don’t manage to reduce k enough!
 - They might be trying to remove the “wrong” a_i .
 - So, try another MODIFICATION, this time where any coordinate can be added/removed, not just the last one.
 - While we’re at it, change “new a_i distribution” from $Z \sim N(0, 1)$ to $Z \sim \text{Uniform}(-20, 30)$, with corresponding change to the $q((k, a), (k', a'))$ formulae.
 - file “Rtrans3” – now works well even if started with $k = 24$.
 - Seems to settle on $k = 6$ regardless of starting value.
 - This seems to indicate rapid mixing – good!
 - SUMMARY: Monte Carlo can be used for nearly everything!
-