

# Markov Chain Monte Carlo

## Metropolis-Hastings and related algorithms

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# Outline

- 1 Conjugate prior distributions
  - Gamma distribution
  - Dirichlet distribution
  - Normal-gamma distribution
  - Summary
- 2 MCMC in theory
  - Motivation
  - Metropolis-Hastings
  - Gibbs sampling
  - Jacobians
  - MCMC on continuous parameters
- 3 MCMC in bioinformatics
  - Alignments
  - Trees
  - Structures

# Motivation

- $P(\theta|x)$  and  $P(\theta)$  are **conjugate** if the posterior  $P(\theta|x)$  is of the same family as the prior  $P(\theta)$

$$P(\theta|x) = \frac{P(x|\theta)P(\theta)}{P(x)} = \frac{P(x|\theta)P(\theta)}{\int P(x, \theta')d\theta'}$$

- The denominator is  $P(x)$ , the **Bayesian evidence**, which does not depend on  $\theta$ .
- The dependence of  $P(\theta|x)$  on  $\theta$  is same as that of  $P(x|\theta)P(\theta)$ , but to get a normalized form for  $P(\theta|x)$ , we need to integrate out  $\theta$  to find the evidence  $P(x)$ .
- The parameters of the conjugate prior distribution are called **hyperparameters**.

# Motivation

$$\begin{aligned}L(\theta) &= P(\text{data } D | \text{params } \theta) &&= \text{likelihood} \\F(\theta, \alpha) &= P(\theta | \text{hyperparams } \alpha) &&= \text{prior} \\G(\theta) &= P(\theta | \alpha, D) &&= \text{posterior}\end{aligned}$$

$L$  and  $F$  are conjugate if  $G(\theta) = F(\theta, \alpha')$  where  $\alpha' \equiv \alpha'(\alpha, D)$

# Common conjugacies

Likelihood	Prior
Exponential	Gamma
Binomial	Beta
Multinomial	Dirichlet
Gaussian (fixed precision)	Gaussian
Gaussian (varying precision)	Gaussian + Gamma

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# Gamma distribution

- Exponential, Poisson, Gamma distributions
  - Exponential distribution: pdf for time to first event,  $T$ , given that mean event rate is  $\mu$

$$P(T|\mu) = \mu \exp(-\mu T)$$

- Poisson: probability distribution of number of events,  $n$ , in time  $T$ , given that mean event rate is  $\mu$

$$P(n|\mu) = \frac{(\mu T)^n \exp(-\mu T)}{n!}$$

(NB exponential distribution can be obtained by setting  $n = 0$  and differentiating w.r.t.  $T$ )

# Gamma distribution

- Poisson

$$P(n|\mu) = \frac{(\mu T)^n \exp(-\mu T)}{n!}$$

- Gamma, conjugate to Poisson.

Shape parameter  $\alpha$ , rate parameter  $\beta$ .

$$P(\mu|\alpha, \beta) = \frac{\mu^{\alpha-1} \beta^\alpha \exp(-\mu\beta)}{\Gamma(\alpha)}$$

where  $\Gamma$  is the **gamma function**

$$\Gamma(\alpha) = \int_0^\infty (\mu\beta)^{\alpha-1} \exp(-\mu\beta) d(\mu\beta)$$



# Gamma function

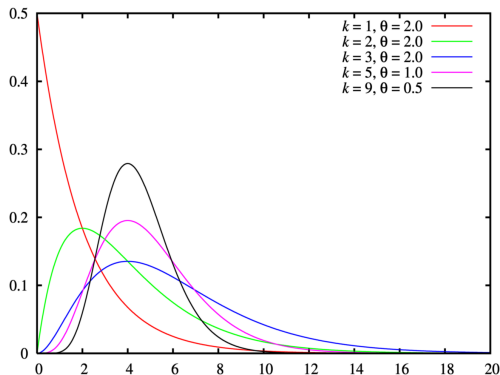
Changing variables in the gamma function integral

$$\Gamma(z) = \int_{u=0}^{\infty} u^{z-1} \exp(-u) du$$

Clearly  $\Gamma(1) = 1$ . Integrating by parts for positive integer  $z$ ,

$$\begin{aligned}\Gamma(z+1) &= \int_{u=0}^{\infty} u^z \exp(-u) du \\ &= [-u^z \exp(-u)]_{u=0}^{\infty} + z\Gamma(z) \\ &= z!\end{aligned}$$

# Gamma distribution



$$k = \alpha, \theta = 1/\beta$$

$\theta$  is called the scale parameter.

# Gamma distribution

- Properties of gamma distribution:  
mean of  $\mu$  is  $\alpha/\beta$  and variance is  $\alpha/\beta^2$ .
- Conjugacy:

$$P(n) = \frac{\Gamma(\alpha')}{\Gamma(n+1)\Gamma(\alpha)} \frac{T^n \beta^\alpha}{(\beta')^{(\alpha')}} \\ P(\mu|n) = \frac{\mu^{\alpha'-1} (\beta')^{\alpha'} \exp(-\mu\beta')}{\Gamma(\alpha')}$$

i.e. posterior for  $\mu$  is a gamma distribution with shape  $\alpha' = \alpha + n$  and rate  $\beta' = \beta + t$ .

- $\alpha$  and  $\beta$  are like a *pseudocount* and *pseudotime*.

# Gamma distribution

- Lengthier derivation:

$$\begin{aligned}
 P(n) &= \int_{\mu=0}^{\infty} P(n|\mu)P(\mu)d\mu \\
 &= \int_{\mu=0}^{\infty} \frac{(\mu T)^n \exp(-\mu T)}{\Gamma(n+1)} \frac{\mu^{\alpha-1} \beta^{\alpha} \exp(-\mu\beta)}{\Gamma(\alpha)} d\mu \\
 &= \frac{T^n \beta^{\alpha}}{\Gamma(n+1)\Gamma(\alpha)} \int_{\mu=0}^{\infty} \mu^{n+\alpha-1} \exp(-\mu(T+\beta)) d\mu \\
 &= \frac{T^n \beta^{\alpha}}{\Gamma(n+1)\Gamma(\alpha)} \frac{1}{(T+\beta)^{n+\alpha}} \int_{u=0}^{\infty} u^{n+\alpha-1} \exp(-u) du \\
 &= \frac{\Gamma(\alpha')}{\Gamma(n+1)\Gamma(\alpha)} \frac{T^n \beta^{\alpha}}{(\beta')^{(\alpha')}}
 \end{aligned}$$

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# Dirichlet distribution

- Multinomial, Dirichlet distributions
  - Multinomial:  $K$  possible outcomes, outcome probabilities  $\mathbf{p}$ , outcome frequencies  $\mathbf{n}$  in  $N = \sum_k n_k$  trials

$$P(\mathbf{n}|\mathbf{p}) = \frac{N!}{\prod_k n_k!} \prod_k p_k^{n_k}$$

- Conjugate prior: let  $\mathbf{a}$  be a vector of **pseudocounts**,  $a_1 \dots a_K$ . The **Dirichlet distribution** for  $\mathbf{p}$  is

$$P(\mathbf{p}|\mathbf{a}) = \frac{\prod_i p_i^{a_i-1}}{\mathcal{B}(\mathbf{a})} \delta \left( \sum_i p_i - 1 \right)$$

where  $\mathcal{B}()$  is the **type one Dirichlet integral** or the **multinomial beta function**

$$\mathcal{B}(\mathbf{a}) = \int \left( \prod_i p_i^{a_i-1} \right) \delta \left( \sum_i p_i - 1 \right) d\mathbf{p} = \frac{\prod_k \Gamma(a_k)}{\Gamma(\sum_k a_k)}$$

# Dirichlet distribution

- Properties: mean value of  $p_k$  is  $a_k / \sum_j a_j$ . Modal value is  $p_k = (a_k - 1) / (\sum_j a_j - 1)$ .
- Note the relationship between the multinomial coefficient and  $\mathcal{B}$

$$\frac{N!}{\prod_k n_k!} = \frac{1}{\mathcal{B}(\mathbf{n} + \mathbf{1})} \times \frac{\Gamma(N + 1)}{\Gamma(N + K)}$$

- Conjugacy:

$$P(\mathbf{n}|\mathbf{a}) = \frac{\mathcal{B}(\mathbf{a}')}{\mathcal{B}(\mathbf{a})} \frac{N!}{\prod_k n_k!}$$

$$P(\mathbf{p}|\mathbf{n}, \mathbf{a}) = \frac{\prod_i p_i^{a'_i - 1}}{\mathcal{B}(\mathbf{a}')} \delta \left( \sum_i p_i - 1 \right)$$

where  $\mathbf{a}' = \mathbf{a} + \mathbf{n}$  (hence the name “pseudocount”).

## Dirichlet distribution

- Special case of {Multinomial, Dirichlet} when  $K = 2$  is {Binomial, Beta}, and  $\mathcal{B}$  is the **beta function**.
- The following snippets from Mathworld point to a more rigorous solution of the type one Dirichlet integral
  - The beta integral may be obtained by writing  $m!n!$  as a product of gamma functions with integrands  $u, v$ , transforming to  $(x, y) = (\sqrt{u}, \sqrt{v})$  and then to polar co-ordinates  $(x, y) = r(\cos \theta, \sin \theta)$ . This yields

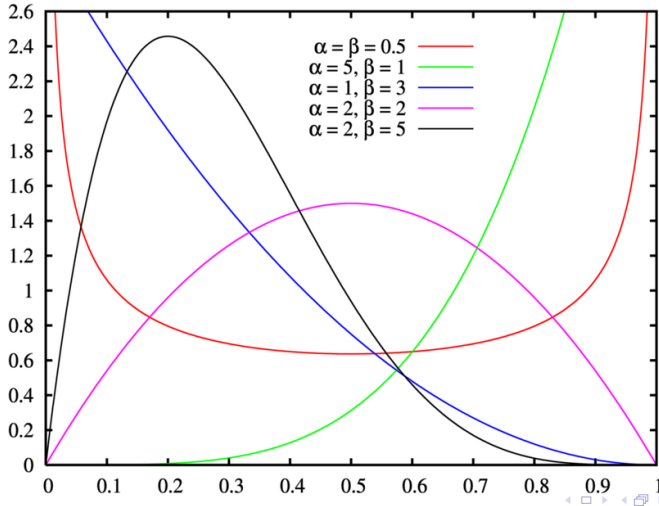
$$\mathcal{B}(m+1, n+1) = 2 \int_0^{\pi/2} \cos^{2m+1} \theta \sin^{2n+1} \theta d\theta = \frac{m!n!}{(m+n+1)!}$$

See [mathworld.wolfram.com/BetaFunction.html](http://mathworld.wolfram.com/BetaFunction.html) for details.

- More info on Dirichlet type one integral at [mathworld.wolfram.com/DirichletIntegrals.html](http://mathworld.wolfram.com/DirichletIntegrals.html)

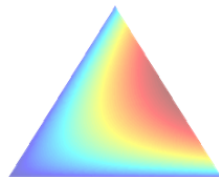
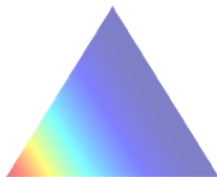
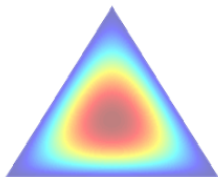


# Beta distribution



# Dirichlet distribution

- For  $K = 3$ , the probabilities  $(p_x, p_y, p_z)$  satisfy  $p_x + p_y + p_z = 1$
- Like 3D points lying on the 2D triangle with corners at  $(0, 0, 1)$ ,  $(0, 1, 0)$  and  $(1, 0, 0)$



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# Normal-gamma distribution

- Normal (Gaussian) and Normal-gamma distributions
  - Mean  $\mu$ , precision  $\tau$  (precision is reciprocal of variance).  
Data  $\{x_i\}$  with moments  $m_k = \sum_i x_i^k$

$$\begin{aligned} P(\mathbf{x}|\mu, \tau) &= \prod_i \sqrt{\frac{\tau}{2\pi}} \exp\left(-\frac{\tau}{2}(x_i - \mu)^2\right) \\ &= \left(\frac{\tau}{2\pi}\right)^{m_0/2} \exp\left(-\frac{\tau}{2}(m_2 - 2\mu m_1 + \mu^2 m_0)\right) \end{aligned}$$

# Normal-gamma distribution

- Normal

$$P(\mathbf{x}|\mu, \tau) = \left(\frac{\tau}{2\pi}\right)^{m_0/2} \exp\left(-\frac{\tau}{2}(m_2 - 2\mu m_1 + \mu^2 m_0)\right)$$

- Conjugate prior

$$P(\mu, \tau) = P(\tau)P(\mu|\tau)$$

$$P(\tau) = \frac{e^{-\beta\tau} \tau^{\alpha-1} \beta^\alpha}{\Gamma(\alpha)}$$

$$P(\mu|\tau) = \sqrt{\frac{\lambda\tau}{2\pi}} \exp\left(-\frac{\lambda\tau}{2}(\mu - \epsilon)^2\right)$$

- $P(\tau)$  is gamma (shape  $\alpha$ , rate  $\beta$ )
- $P(\mu|\tau)$  is Normal (mean  $\epsilon$ , precision  $\lambda\tau$ )
- Full set of hyperparameters is  $\{\alpha, \beta, \epsilon, \lambda\}$

# Normal-gamma distribution

- Conjugacy:

$$P(\mathbf{x}) = (2\pi)^{-m_0/2} \frac{\Gamma(\alpha')}{\Gamma(\alpha)} \frac{\beta^\alpha}{\beta'^{\alpha'}} \sqrt{\frac{\lambda}{\lambda'}}$$

$$P(\mu, \tau | \mathbf{x}) = \frac{e^{-\beta' \tau} \tau^{\alpha' - 1} \beta'^{\alpha'}}{\Gamma(\alpha')} \times \sqrt{\frac{\lambda' \tau}{2\pi}} \exp\left(-\frac{\lambda' \tau}{2} (\mu - \epsilon')^2\right)$$

where  $\epsilon' = \frac{\lambda\epsilon + m_1}{\lambda + m_0}$ ,  $\lambda' = \lambda + m_0$ ,  $\alpha' = \alpha + \frac{m_0}{2}$  and  $\beta' = \beta + \frac{1}{2} \left( \epsilon^2 + m_2 - \frac{(\lambda\epsilon + m_1)^2}{\lambda + m_0} \right)$ .

- NB if we regard  $\tau$  as fixed, then Gaussian is auto-conjugate.

## $\chi^2$ distribution

$$P(\mu, \tau | \mathbf{x}) = \frac{e^{-\beta' \tau} \tau^{\alpha' - 1} \beta'^{\alpha'}}{\Gamma(\alpha')} \times \sqrt{\frac{\lambda' \tau}{2\pi}} \exp\left(-\frac{\lambda' \tau}{2} (\mu - \epsilon')^2\right)$$

- Rescale the posterior distribution for  $\tau$ : divide by  $(m_2/m_0 - m_1^2/m_0)^{-1}$  (the posterior mean estimate for  $\tau$ )
- Resulting gamma distribution with shape  $m_0/2$  and scale  $1/2$  is the “ $\chi^2$  distribution”.
- Canonical definition of the quantity  $\chi^2$  is in the special case where  $\mu = 0$  and  $\tau = 1$ , in which case  $\chi^2 = m_2$

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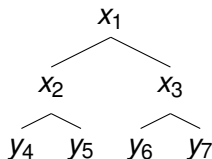
# Common conjugacies

- Exponential/Poisson  $\leftrightarrow$  Gamma
- Binomial  $\leftrightarrow$  Beta
- Multinomial  $\leftrightarrow$  Dirichlet
- Normal  $\leftrightarrow$  Normal-gamma

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# Motivation



$$P(X|Y) = \frac{P(X,Y)}{P(Y)} = \frac{P(X,Y)}{\sum_{X'} P(X',Y)}$$

- Let  $A$  = alphabet size
- Felsenstein's pruning algorithm for  $P(Y)$ 
  - Time  $\mathcal{O}(NA^2)$
  - Memory  $\mathcal{O}(NA)$
- Exponentiating an  $A \times A$  matrix  $N$  times
  - Time  $\mathcal{O}(NA^3)$
- OK: bases ( $A = 4$ ), amino acids ( $A = 20$ ), codons ( $A = 64$ )
  - Painful: Gene Ontology ( $A = 33,587$ ), GO-Slim ( $A = 127$ )

# MCMC: general idea

- Want to sample from some pdf  $P(X) = \frac{f(X)}{Z}$  where  $X \in \mathcal{X}$
- Problem: set  $\mathcal{X}$  is large; computing  $Z = \sum_{X \in \mathcal{X}} f(X)$  is hard
- Construct a stochastic process  $X_1, X_2, X_3, X_4, \dots$  such that the equilibrium distribution is  $P(X)$
- If the process is ergodic, then  $X_t \sim P(X)$  for “large”  $t$ 
  - How large must  $t$  be? The **burn-in time** or **mixing time**

# MCMC: construction

- Want a Markov process  $X_1, X_2, \dots$  w/equilibrium  $P(X)$
- Let  $Q(i, j) = P(X_{t+1} = j | X_t = i)$  be the **transition matrix**
- One way to make  $P(X)$  the equilibrium distribution is to force  $Q$  to satisfy **detailed balance**

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

- If  $P(X) = \frac{f(X)}{Z}$  we can drop the  $Z$

$$f(i)Q(i, j) = f(j)Q(j, i)$$

- Note this process is **reversible** (MCMC doesn't have to be, but usually is)

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# Metropolis-Hastings algorithm: the proposal

- One way to construct a  $Q$  that satisfies detailed balance

$$f(i)Q(i, j) = f(j)Q(j, i)$$

- Start with a **proposal distribution**

$$U(i, j) = P(X_{t+1} = j | X_t = i)$$

- Metropolis/Teller (1953): **symmetric** proposal,  
 $U(i, j) = U(j, i)$
- Hastings (1970) generalized to **reversible** proposal

$$g(i)U(i, j) = g(j)U(j, i)$$

- Note  $g \neq f$ : proposal does **not** have to converge on  $f$
- Now we adapt this  $U$  to construct  $Q$

# Metropolis-Hastings algorithm: accept/reject

- We have a proposal density,  $U$ , satisfying (for some  $g$ )

$$g(i)U(i, j) = g(j)U(j, i)$$

- Process:

- Given  $X_n$ , **propose a move** by sampling  $X'$  from

$$P(X' = j | X_n = i) = U(i, j)$$

- Calculate the **Hastings ratio**

$$h(X, X') = \frac{f(X')}{f(X)} \frac{g(X)}{g(X')} = \frac{f(X')}{f(X)} \frac{U(X', X)}{U(X, X')}$$

- If  $h(X, X') \geq 1$ , then **accept the move**: set  $X_{n+1} \leftarrow X'$
    - If  $h(X, X') < 1$ , then
      - Sample an r.v.  $\alpha$  from  $U(0, 1)$  (uniform distribution)
      - If  $\alpha < h(X, X')$ , then accept the move
      - If  $\alpha \geq h(X, X')$ , then **reject the move**: set  $X_{n+1} \leftarrow X_n$



## Metropolis-Hastings: proof of detailed balance

$$\begin{aligned}h(i, j) &= \frac{f(j)}{f(i)} \frac{U(j, i)}{U(i, j)} \\Q(i, j) &= U(i, j) \times \min(1, h(i, j))\end{aligned}$$

- Suppose (with no loss of generality) that  $h(i, j) \leq 1$

$$\begin{aligned}Q(i, j) &= U(i, j)h(i, j) \\&= U(i, j) \frac{f(j)}{f(i)} \frac{U(j, i)}{U(i, j)} \\&= \frac{f(j)}{f(i)} U(j, i)\end{aligned}$$

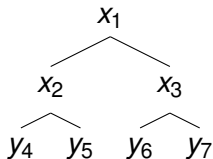
- Since  $h(j, i) \geq 1$ , we must have  $Q(j, i) = U(j, i)$ , and so

$$f(i)Q(i, j) = f(j)Q(j, i) \quad \square$$

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# Gibbs sampling



- The  $X$  in  $P(X)$  is  $N$ -dimensional,  $X = (x_1, x_2 \dots x_N)$
- **Gibbs Sampling** proposal distribution is as follows
  - Pick a dimension  $m$ , where  $1 \leq m \leq N$
  - Conceptually, fix all of the  $\{x_n\}$  except for  $x_m$ ; sample  $x_m$  from its marginal distribution conditional on all the others
  - Proposal distribution is

$$\begin{aligned} U(X, X') &= U(\{x_n\}, \{x'_n\}) \\ &= P(x'_m | \{x_n : n \neq m\}) \times \prod_{n \neq m} \delta(x'_n = x_m) \end{aligned}$$

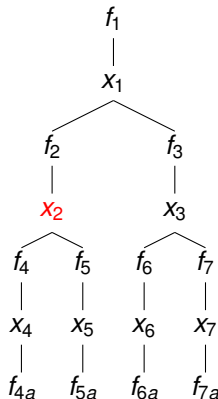
## Gibbs sampling: the marginal distribution

Let  $X_{\text{other}} = \{x_n : n \neq m\}$

$$\begin{aligned} P(x'_m | X_{\text{other}}) &= \frac{P(x'_m, X_{\text{other}})}{P(X_{\text{other}})} \\ &= \frac{P(x'_m, X_{\text{other}})}{\sum_k P(x'_m = k, X_{\text{other}})} \end{aligned}$$

If  $P(X)$  is a product of functions of sparse subsets of  $X$  (c.f. factor graphs), then many of these functions will cancel in numerator and denominator

# Gibbs sampling: factor graph example



$$P(x'_2 | X_{\text{other}}) = \frac{f_2(x_1, x'_2) f_4(x'_2, x_4) f_5(x'_2, x_5)}{\sum_k f_2(x_1, k) f_4(k, x_4) f_5(k, x_5)}$$

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# The Jacobian matrix and its determinant

- Function  $F : \mathbb{R}^n \rightarrow \mathbb{R}^m$
- Let  $(y_1 \dots y_m) = F(x_1 \dots x_n)$
- The **Jacobian matrix** of  $F$  is

$$J_F(x_1, \dots, x_n) = \begin{bmatrix} \frac{\partial y_1}{\partial x_1} & \dots & \frac{\partial y_1}{\partial x_n} \\ \vdots & \ddots & \vdots \\ \frac{\partial y_m}{\partial x_1} & \dots & \frac{\partial y_m}{\partial x_n} \end{bmatrix}$$

- For square matrices ( $m = n$ ), the magnitude of the **Jacobian determinant**,  $|\det(J_F)|$ , gives the factor by which the function  $F$  expands or shrinks volumes in  $\mathbf{x}$

# Multidimensional change of variables

- If  $p(x)$  is pdf of  $x$ , and  $y = F(x)$ , what is pdf  $q(y)$  of  $y$ ?
  - Now with multidimensional  $x = (x_1 \dots x_n)$  and  $y = (y_1 \dots y_n)$
- Derivation:

$$\begin{aligned} dy_1 dy_2 \dots dy_n &= |\det(J_F)| dx_1 dx_2 \dots dx_n \\ q(y) dy_1 dy_2 \dots dy_n &= p(x) dx_1 dx_2 \dots dx_n \\ q(y) &= p(x) / |\det(J_F)| \\ &= p(F^{-1}(y)) |\det(J_{F^{-1}})| \end{aligned}$$

where the last line uses the **inverse function theorem**,

$$J_{F^{-1}} = (J_F)^{-1} \quad \dots \text{ c.f. } \textbf{chain rule: } J_{F \circ G}(x) = J_F(G(x)) J_G(x)$$



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# MCMC on continuous variables

- Let  $x$  be a continuous state
- Suppose the proposal generates  $x'$  by sampling some parameter  $u$  from a p.d.f.  $g(u)$ , so  $x' = x'(x, u)$ 
  - Let  $u' = u'(x, u)$  be the **inverse** parameter that takes us back from  $x' \rightarrow x$
  - For convenience, define the function  $F(x, u) = (x', u')$
- The Hastings ratio then depends on the Jacobian  $J_F$

$$h(x, x') = \frac{P(x')}{P(x)} \frac{g(u')}{g(u)} |\det(J_F)|$$

- Interpretation: we are moving through  $(x, u)$ -space rather than just  $x$ -space
- Green (1995, 2003)

# There's much more

- Summarizing results of an MCMC run
- Diagnosing MCMC convergence
  - Replicates
  - Intra- and inter-chain variance

Not covered in depth here.

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# Sampling ungapped sequence alignments

- Example: ungapped sequence alignments (Lawrence *et al*)
  - Suppose you have  $K$  sequences,  $\{S^{(1)} \dots S^{(K)}\}$ , and that  $S_i^{(k)}$  is the  $i$ 'th residue of the  $k$ 'th sequence.
  - Let  $x_k$  be the indentation of the  $k$ 'th sequence. The motif (of length  $M$ ) runs from  $x_k$  to  $x_k + M - 1$ .

# Sampling ungapped alignments

- Let  $\mathcal{U}$  denote the null hypothesis that the sequences are unrelated. Then

$$\frac{P(\mathbf{x})}{P(\mathcal{U})} = \prod_{m=0}^{M-1} \frac{f(\mathbf{n}(m))}{\prod_{\omega} q(\omega)^{n_{\omega}(m)}}$$

where  $q(\omega)$  is a background distribution over alphabet symbols  $\omega$ , and  $n_{\omega}(m)$  is the number of times symbol  $\omega$  appears in column  $m$  of the motif

$$n_{\omega}(m) = \sum_{k=1}^K \delta(S_{x_k+m} = \omega)$$

# Sampling ungapped alignments

- The  $f(\mathbf{n})$  function should reward conserved columns. Lawrence *et al* use an entropy-like measure

$$f(\mathbf{n}) = \prod_{\omega} p_{\omega}^{n_{\omega}} = \exp(-KS[p])$$

where  $p_k = n_k/K$ . Note that this is the distribution  $\mathbf{p}$  that maximizes  $\prod_{\omega} p_{\omega}^{n_{\omega}}$  for a given  $\mathbf{n}$ . Due to this implicit maximization, the above  $f(\mathbf{n})$  is not strictly a probability distribution for  $\mathbf{n}$ .

- Typically a vector of pseudocounts  $\mathbf{a}$  is added to  $\mathbf{n}$ . This suggests the Dirichlet evidence as an alternative form for  $f$  that does not involve any implicit maximization:

$$f(\mathbf{n}) = \frac{\mathcal{B}(\mathbf{n} + \mathbf{a})}{\mathcal{B}(\mathbf{a})\mathcal{B}(\mathbf{n})}$$

# Sampling ungapped alignments

- Another way to formulate this problem is as a chain over  $(\mathbf{x}, \{\mathbf{p}(m)\})$  with Gibbs-sampling steps that alternate between resampling one of the  $x_k$ 's and resampling all the  $\mathbf{p}$ 's.
- For good mixing it's also useful to allow moves that slide the entire motif window, i.e.  $\mathbf{x} \leftarrow \mathbf{x} \pm \mathbf{1}$



# Outline

- 1 Conjugate prior distributions
  - Gamma distribution
  - Dirichlet distribution
  - Normal-gamma distribution
  - Summary
- 2 MCMC in theory
  - Motivation
  - Metropolis-Hastings
  - Gibbs sampling
  - Jacobians
  - MCMC on continuous parameters
- 3 MCMC in bioinformatics
  - Alignments
  - Trees
  - Structures

# Sampling trees

- Priors: coalescent
- Moves:
  - Topology: prune-and-graft operations, swapping nodes & branches
  - Branch lengths: rescaling, sliding
- Equivalent ML moves:
  - quartet-puzzling (Strimmer & von Haeseler)
  - pplacer's "pendant branch length" (Matsen)

# Extensions

- Co-sampling alignment and tree (and secondary/3D structure, ...)
- Models and approximations that sum out alignment, ancestral sequence, tree

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# Molecular dynamics

- Folding
- Docking
- Transport
- Kinetics
- etc.