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# Outline

- Setting Bayesian inference for simulation codes
  - numerous continuous parameters
  - expensive function evaluations
- Adjoint differentiation on basis of code
  - efficiently calculates gradient of computed scalar quantity
  - uses
  - methods of implementation
- Hybrid Markov Chain Monte Carlo
  - basic algorithm
  - requires gradient of minus-log-probability
- A method to test convergence of MCMC sequence
  - based on gradient of minus-log-probability

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# Uses of adjoint differentiation

- Fitting large-scale simulations to data (regression):
  - atmosphere and ocean models, fluid dynamics, hydrodynamic
- Reconstruction: imaging through diffusive media
- Hybrid Markov Chain Monte Carlo
- Uncertainty analysis of simulation code
  - sensitivity of uncertainty variance to each contributing cause
- Metropolis-Hastings MCMC calculations
  - sensitivity of efficiency (or acceptance fraction) wrt proposal distribution parameters

# Maximum likelihood estimation by optimization



- Optimization process is accelerated by using **gradient-based algorithms**; therefore need gradients of functional behavior of simulation process
- Adjoint differentiation facilitates efficient calculation of gradients, i.e. derivative of scalar output  $(\frac{1}{2}\chi^2)$  wrt parameters **x**

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#### Derivative calculation by finite differences

• Derivative for function defined as limit of ratio of finite differences:

$$\frac{df}{dx}\Big|_{x1} = \lim_{\Delta x \to 0} \frac{f(x1 + \Delta x) - f(x1)}{\Delta x}$$

Wish to estimate derivatives of calculated function for which there is no analytic relation between outputs and inputs

- Numerical estimation based on finite differences is problematical:
  - difficult to choose perturbation  $\Delta x$
  - # function evaluations ~# variables
- Estimation based on functionality implied by computer code is more reliable



Error in derivative of sin(x) vs.  $\Delta x$  at  $x = \pi/4$ 

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# Differentiation of sequence of transformations

$$\xrightarrow{\mathbf{X}} \mathbf{A} \xrightarrow{\mathbf{y}} \mathbf{B} \xrightarrow{\mathbf{z}} \mathbf{C} \xrightarrow{\boldsymbol{\phi}}$$

- Data-flow diagram shows sequence of transformations A->B->C that converts data structures x to y to z and to scalar φ (forward calculation)
- Desire derivatives of  $\varphi$  wrt all components of  $\mathbf{x}$ , *assuming* that  $\varphi$  is *differentiable*
- Chain rule applies:  $\frac{\partial \varphi}{\partial x_i} = \sum_{j,k} \frac{\partial y_j}{\partial x_i} \frac{\partial z_k}{\partial y_j} \frac{\partial \varphi}{\partial z_k}$
- Two choices for summation order:
  - doing *j* before *k* means derivatives follow data flow (forward calculation)
  - doing *k* before *j* means derivatives flow in reverse (adjoint) direction

# Adjoint Differentiation In Code Technique ADICT



- For sequence of transformations that converts data structure **x** to scalar  $\varphi$
- Derivatives  $\frac{\partial \varphi}{\partial \mathbf{x}}$  are efficiently calculated in the reverse (adjoint) direction
- **Code-based approach**: logic of adjoint code is based explicitly on the forward code or on derivatives of the forward algorithm
- Not based on the theoretical eqs., which forward calc. only approximates
- Only assumption is that  $\phi$  is a **differentiable function** of **x**
- CPU time to compute **all** derivatives is comparable to forward calculation

# Level of abstraction of implementation

- One can choose to differentiate forward calculation at various levels of abstraction
  - model based
    - e.g., differentiate partial differential equations and solve
    - not advised because forward codes only approximates model
  - module or algorithm based
    - differentiate each basic algorithm (Bayes Inference Engine)
  - code based
    - direct interpretation of computer code (FORTRAN, C, etc.)
    - automatic differentiation utilities produce derivative code(s)
  - instruction based
    - reverse the sequence of CPU instructions for any particular calc.

# Example of algorithm-based approach

- Bayes Inference Engine (BIE) created at LANL
  - modeling tool for interpreting radiographs
  - BIE programmed by creating data-flow diagram linking transforms, as shown here for 3D reconstruction problem

#### • Adjoint differentiation crucial to BIE success



# Adjoint differentiation in diffusion calculation

- Adjoint differentiation calculation precisely reverses direction of forward calculation
- Each forward data structure has an associated derivative

- where  $\mathbf{U}_{n}$  propagates forward,  $\frac{\partial \varphi}{\partial \mathbf{U}_{n}}$  goes backward  $(\varphi = \frac{1}{2}\chi^{2})$ 



Optical tomographic reconstruction – determine image of light diffusion characteristics from measurements of how will IR light passes through tissue

# Automatic differentiation tools

- Several tools exist for automatically differentiating codes; various capabilities, e.g., forward or reverse (adjoint) differentiation, handling of large codes, etc.
  - FORTRAN 77 (90 under development)
    - ADIFOR (reverse mode)
    - TAMC (reverse mode)
    - TAPENADE (reverse mode)
  - C (C++ under development)
    - ADIC
    - ADOL-C (reverse mode)
  - MATLAB
    - ADMAT
- Very active area of development

# MCMC for simulations



- log(likelihood) distribution is result of calculation; function of parameters x
- Markov Chain Monte Carlo (MCMC) algorithm draws random samples of x from posterior probability p(x|Y)
- Produces plausible set of parameters  $\{x\}$ ; therefore model realizations

## MCMC - problem statement

- Parameter space of *n* dimensions represented by vector **x**
- Draw a set of samples {x<sub>k</sub>} from a given "arbitrary" target probability density function (pdf), q(x)
- Only requirement typically is that one be able to evaluate  $Cq(\mathbf{x})$  for any given  $\mathbf{x}$ , where C is an unknown constant; that is,  $q(\mathbf{x})$  need not be normalized
- Although focus here is on continuous variables, MCMC applies to discrete variables as well

# Uses of MCMC

• Permits evaluation of the expectation values of functions of **x**, e.g.,

$$\langle f(\mathbf{x}) \rangle = \int f(\mathbf{x}) q(\mathbf{x}) d\mathbf{x} \cong (1/K) \Sigma_k f(\mathbf{x}_k)$$

– typical use is to calculate mean  $\langle x\rangle$  and variance  $\langle (x$  -  $\langle x\rangle)^2\rangle$ 

- Useful for evaluating integrals, such as the partition function for properly normalizing the pdf
- Dynamic display of sequences provides visualization of uncertainties in model and range of model variations
- Automatic marginalization; when considering any subset of parameters of an MCMC sequence, the remaining parameters are marginalized over (integrated out)

# Metropolis Markov Chain Monte Carlo

Generates sequence of random samples from an arbitrary probability density function

- Metropolis algorithm:
  - draw trial step from symmetric pdf, i.e.,  $t(\Delta \mathbf{x}) = t(-\Delta \mathbf{x})$
  - accept or reject trial step
  - simple and generally applicable
  - requires only calculation of target pdf,  $q(\mathbf{x})$ , for any  $\mathbf{x}$



# Metropolis algorithm

- Select initial parameter vector **x**<sub>0</sub>
- Iterate as follows: at iteration number k
  - (1) create new trial position x\* = x<sub>k</sub> + Δx, where Δx is randomly chosen from t(Δx)
    (2) calculate ratio r = q(x\*)/q(x<sub>k</sub>)
    (3) accept trial position, i.e. set x<sub>k+1</sub> = x\* if r ≥ 1 or with probability r, if r < 1</li>

otherwise stay put,  $\mathbf{x}_{k+1} = \mathbf{x}_k$ 

- Only requires computation of  $q(\mathbf{x})$  (with arbitrary normalization)
- Creates Markov chain since  $\mathbf{x}_{k+1}$  depends only on  $\mathbf{x}_k$

# Gibbs algorithm

- Vary only one component of **x** at a time
- Draw new value of  $x_j$  from conditional pdf  $q(x_j | x_1 x_2 \dots x_{j-1} x_{j+1} \dots)$
- Cycle through all components

Probability( $x_1, x_2$ )



# Hybrid MCMC method

- Called hybrid method because it alternates Gibbs & Metropolis steps (better called "Hamiltonian" method?)
- Associate with each parameter  $x_i$  a momentum  $p_i$
- Define a Hamiltonian (sum of potential and kinetic energy):

 $H = \varphi(\mathbf{x}) + \sum p_i^2 / (2 m_i) ,$ where  $\varphi = -\log (q(\mathbf{x}))$ 

- Objective is to draw samples from new pdf:  $q'(\mathbf{x}, \mathbf{p}) \propto \exp(-H(\mathbf{x}, \mathbf{p})) = q(\mathbf{x}) \exp(-\sum p_i^2/(2m_i))$
- Samples {x<sub>k</sub>} from q'(x, p) represent draws from q(x)
   because p dependence marginalized out

# Hybrid algorithm



green path - trajectory with constant *H*, follow by Metropolis

# Hamiltonian algorithm

- Gibbs step: randomly sample momentum distribution
- Follow trajectory of constant *H* using leapfrog algorithm:

$$p_{i}(t + \frac{\tau}{2}) = p_{i}(t) - \frac{\tau}{2} \frac{\partial \varphi}{\partial x_{i}} \Big|_{\mathbf{X}(t)}$$
$$x_{i}(t + \tau) = x_{i}(t) + \frac{\tau}{m_{i}} p_{i}(t + \frac{\tau}{2})$$
$$p_{i}(t + \tau) = p_{i}(t + \frac{\tau}{2}) - \frac{\tau}{2} \frac{\partial \varphi}{\partial x_{i}} \Big|_{\mathbf{X}(t + \tau)}$$

where  $\tau$  is leapfrog time step.

- Repeat leapfrog a predetermined number of times
- Metropolis step: accept or reject on basis of *H* at beginning and end of H trajectory

# Hybrid algorithm implementation

- Gibbs step easy because draws are from uncorrelated Gaussian
- H trajectories followed by several leapfrog steps permit long jumps in (**x**, **p**) space, with little change in *H* 
  - specify total time = T ; number of leapfrog steps =  $T/\tau$
  - randomize *T* to avoid coherent oscillations
  - reverse momenta at end of H trajectory to guarantee that it is symmetric process (condition for Metropolis step)
- Metropolis step no rejections if *H* is unchanged
- Adjoint differentiation efficiently provides gradient

#### 2D isotropic Gaussian distribution



Long H trajectories - shows ellipses when  $\sigma_1 = \sigma_2 = 1$ ,  $m_1 = m_2 = 1$  Randomize length of H trajectories to obtain good sampling of pdf

#### 2D correlated Gaussian distribution



- 2D Gaussian pdf with high correlation (r =0.95)
- Length of H trajectories randomized

### MCMC Efficiency

– Estimate of a quantity from its samples from a pdf q(v)

$$\widetilde{v} = \frac{1}{N_k} \Sigma v_k$$

- For *N* independent samples drawn from a pdf, variance in estimate:  $var(\tilde{v}) = \frac{var(v)}{N}$
- For *N* samples from an MCMC sequence with target pdf q(v)var(v)

$$\operatorname{var}(\widetilde{v}) = \frac{\operatorname{var}(v)}{\eta N}$$

where  $\eta$  is the sampling efficiency

- Thus,  $\eta^{-1}$  iterations needed for one statistically independent sample
- Let v = variance because aim is to estimate variance of target pdf

# n-D isotropic Gaussian distributions

- MCMC efficiency versus number dimensions
  - Hamiltonian method: drops little
  - Metropolis method:
     goes as 0.3/n
- Hybrid (Hamiltonian) method much more efficient at high dimensions
- Assumes gradient eval. costs same as function



• Variance integral

$$\operatorname{var}(x_i) = \int (x_i - \overline{x}_i)^2 p(\mathbf{x}) d\mathbf{x}$$

$$=\int_{-\frac{1}{3}}^{\frac{1}{3}} (x_i - \overline{x}_i)^3 \nabla \varphi(\mathbf{x}) p(\mathbf{x}) d\mathbf{x} + \frac{1}{3} (x_i - \overline{x}_i)^3 p(\mathbf{x}) \Big|$$

by integration by parts and  $\varphi(\mathbf{x}) = -\log(p(\mathbf{x}))$ 

- limits are typically  $\pm \infty$  and last term is usually zero
- thus, integrals are equal
- Form ratio of integrals, computed from samples  $\mathbf{x}^k$  from  $p(\mathbf{x})$

$$R = \frac{\sum (x_i^k - \overline{x}_i^k)^3 \frac{\partial \varphi}{\partial x_i^k}}{3\sum (x_i^k - \overline{x}_i^k)^2} , \qquad \overline{x}^k = \sum x_i^k$$

• *R* tends to be less than 1 when  $p(\mathbf{x})$  not adequately sampled

## 2D non-isotropic Gaussian distribution



- Nonisotropic Gaussian target pdf:  $\sigma_1 = 4$ ,  $\sigma_2 = 1$ ,  $m_1 = m_2 = 1$
- Randomize length of H trajectories to get random sampling
- Convergence; does sequence actually sample target pdf?

#### Convergence - 2D nonisotropic Gaussians



- Non-isotropic Gaussian target pdf:  $\sigma_1 = 4$ ,  $\sigma_2 = 1$ ,  $m_1 = m_2 = 1$ 
  - control degree of pdf sampling by using short leapfrog steps ( $\tau = 0.2$ ) and  $T_{max} = 2$
- Test statistic R < 1 when estimated variance is deficient June 5, 2002 Valencia 7

# 16D correlated Gaussian distribution



- 16D Gaussian pdf related to smoothness prior based on integral of L2 norm of second derivative
  - Efficiency/(function evaluation) = 2.2% with Hybrid (Hamiltonian) algorithm 0.11% or 1.6% with Metropolis; w/o & with covar. adapt.

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# MCMC - Issues

- Identification of convergence to target pdf
  - is sequence in thermodynamic equilibrium with target pdf?
  - validity of estimated properties of parameters (covariance)
- Burn in
  - at beginning of sequence, may need to run MCMC for awhile to achieve convergence to target pdf
- Use of multiple sequences
  - different starting values can help confirm convergence
  - natural choice when using computers with multiple CPUs
- Accuracy of estimated properties of parameters
  - related to efficiency, described above

# Conclusions

- Adjoint differentiation provides efficient calculation of gradient of scalar function of many variables
  - optimization (regression)
  - MCMC, especially hybrid method (other possible uses exist)
- Hybrid method
  - based on Hamiltonian dynamics
  - efficiency for isotropic Gaussians is about 7% per function evaluation, independent of number of dimensions
  - much better efficiency than Metropolis for large dimensions provided gradient can be efficiently calculated
- Convergence test based on gradient of -log(probability)
  - tests how well MCMC sequence samples full width of target pdf

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