Parallel Tempering and Population Annealing for Rough Free Energy Landscapes

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Melbourne, July 26, 2010

Collaborators

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Motivation

 How to sample equilibrium states of systems with rugged free energy landscapes, e.g. <u>spin</u> <u>glasses</u>, configurational glasses, proteins, NP-hard combinatorial optimization problems.



Problem



Problem



probability

 Markov chain Monte Carlo (MCMC) at a single temperature such as the Metropolis algorithm gets stuck in local minima.







- Introduction to <u>Parallel Tempering</u> (aka Replica Exchange Monte Carlo)
- Strengths and Weaknesses of <u>PT</u> (in several simple landscapes)
- Introduction to <u>Population Annealing</u>
- <u>PA</u> vs <u>PT</u>
- Conclusions

Parallel Tempering



- •*R* replicas at inverse temperatures $\beta_1 > \beta_2 > ... > \beta_R$ (each with the same couplings).
- •MCMC (e.g. Metropolis) on each replica
- •Exchange replicas with energies *E* and *E'* and temperatures β and β' , with probability:

$$p_{\text{swap}} = \min\left[1, e^{(\beta - \beta')(E - E')}\right]$$

Intuition

 Mixing is accelerated by "round trips" from low to high temperature and back.



Parallel tempering can be optimized by minimizing the equilibrium round trip time.

A simple landscape:

 Consider a model free energy landscape with two free energy minima separated by a high barrier.

– JM, PRE 80, 056706 (2009)



Two Well Model

Parallel tempering for the two-well model



- Assumptions:
 - -Fast equilibration within each well by standard MCMC.
 - –No transitions between wells except at β_c where each well is equally probable.
 - Energy is normally distributed in each state; from thermodynamics:

 $\langle E \rangle = -(\beta - \beta_c)(K + H\sigma)$ $\operatorname{Var}(E) = (K + H\sigma)$

Replica exchange probabilities

 For the two-well model, replica exchange transition probabilities can be computed exactly. For symmetric wells (*H*=0):

$$p_{\text{swap}} = \min\left[1, e^{(\beta - \beta')(E - E')}\right]$$
$$\langle E \rangle = -(\beta - \beta_c)K \text{, } \operatorname{Var}(E) = K$$
$$\downarrow$$
$$p_{\text{swap}} = \frac{1}{2} \operatorname{Erfc}\left((\beta - \beta')\sqrt{K}\right)$$





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$$\tau \sim R^2 \sim \beta F$$



Asymmetric wells (H>0)

- Replica exchange is biased toward moving replicas in the deep well to lower temperature.
- Drift and diffusion instead of diffusion:
 - Replicas in the deeper well drift toward lower temperature and vice versa.
- Faster equilibration than for the symmetric case!





 $\beta' > \beta$

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Equilibration time vs well asymmetry

 $\beta \delta F = -\frac{1}{2} (\beta - \beta_c)^2 H$





FIG. 2: The exponential autocorrelation time τ_{exp} for the fraction of sites in the deep well vs. the well depth parameter K for H = 10 (green diamonds), H = 5 (red squares) and H = 0 (blue circles). The lines are best power law fits, $\tau_{exp} \sim K^x$ with x = 0.76, 0.83 and 0.99 for H = 10, 5 and 0, respectively.

PT for the ferromagnetic Ising model



Exponential autocorrelation time for the magnetization for the low temperature Ising model with and without a temperature dependent external field.

Ising spin glass, Gaussian disorder, L=8, $T/T_c=0.2$



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Weakness of Parallel Tempering

- "First-order" free energy landscapes. The simplest and worst case is the "golf-course" potential:
 - $-e^{N}$ microstates,
 - Nearly all have energy $\varepsilon N > 0$
 - An small fraction, $e^{-\epsilon\beta_c N}$ are ground states with energy zero. There is a pseudo phase transition at β_c
 - Probability to be in a ground state:



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 $1 + e^{-(\beta - \beta_c)\epsilon N}$

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- Each replica independently searches for grounds states and sends them to low temperature when they are found.
- Efficiency exceeds that of single temperature MCMC only to the extent that there are more replicas looking for grounds states than cold replicas ($\beta < \beta_c$) needing them.
- Before equilibration round trip-time is misleading.

Population Annealing

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K. Hukushima and Y. Iba, in *THE MONTE CARLO METHOD IN THE PHYSICAL SCIENCES: Celebrating the 50th Anniversary of the Metropolis Algorithm*, edited by J. E. Gubernatis (AIP, 2003), vol. 690, pp. 200–206.

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- Modification of simulated annealing for equilibrium sampling.
- A population of replicas is cooled according to an annealing schedule. Each replica is acted on by a MCMC (e.g. Metropolis) at the current temperature.
- During each temperature step, the population is differentially resampled according to Boltzmann weights to maintain equilibrium.

Population Annealing



Population annealing = simulated annealing with differential reproduction of replicas

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$$\tau_j(\beta, \beta') = \frac{\exp\left[-(\beta' - \beta)E_j\right]}{Q(\beta, \beta')}$$

$$Q(\beta, \beta') = \frac{\sum_{j=1}^{R} \exp\left[-(\beta' - \beta)E_j\right]}{R}$$

Replica *j* is reproduced n_j times where n_j is a Poisson random number with mean τ_j .

Temperature Step



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Strengths of Population Annealing

- Massively parallel
- Direct measurement of the free energy

$$-\beta_k F(\beta_k) = \sum_{\ell=k}^K \ln Q(\beta_{\ell+1}, \beta_\ell) + \beta_K F(\beta_K) \quad Q(\beta, \beta') = \frac{\sum_{j=1}^R \exp\left[-(\beta' - \beta)E_j\right]}{R}$$

 Comparable efficiency to parallel tempering for Ising spin glasses.

Weighted Averages

JM, arXive: 1006.0252

- Results using small number of replicas are biased.
- Results from independent runs can be combined and biases reduced using weighted averages.
- Observables from each run weighted by the exponential of the free energy estimator for that run.

$$\overline{A}(\beta) = \sum_{m=1}^{M} \tilde{A}_m(\beta) \omega_m(\beta) \qquad \omega_m(\beta) = \frac{e^{-\beta \tilde{F}_m(\beta)}}{\sum_{i=1}^{M} e^{-\beta \tilde{F}_i(\beta)}}$$

 Number of runs needed for unbiased results determined by variance of the free energy estimator--an intrinsic measure of equilibration.



Bias in the energy for the 1D Ising spin glass as a function of ensemble size.

Population Annealing	Parallel Tempering
Sequential Monte Carlo	Markov Chain Monte Carlo
# Replicas (R) 7 space	ی #Sweeps (t) parallel time
#Temperature steps (T) 🗲 parallel time	HReplicas (R) space

Compare PA and PT for the Two-Well Model







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Sequential Monte Carlo	Markov Chain Monte Carlo
# Replicas (R) space	#Sweeps (t) parallel time
#Temperature steps (T) 7	HReplicas (R) space
$(A-A_{eq}) \approx R_0/R$	$(A - A_{eq}) \approx \mathrm{e}^{(-t/\tau)}$
$R_0 \sim K^{1/2}$ $T \sim K^{1/2}$	$\tau \sim K R \sim K^{1/2}$



- Both parallel tempering and population annealing are efficient at finding equilibria between free energy minima with large basins of attraction separated by large barriers.
 - For PT efficiency is greatest when the minima are highly asymmetric
- Neither method is useful, except via brute force parallelism, for the case of free energy minima with small basins of attraction.
- Population annealing is comparably efficient to parallel tempering and has several features to recommend it:
 - Highly parallel
 - Direct measurement of free energy
 - Built-in equilibration criteria (variance of the free energy estimator)