

Annealing

- Analogy with physical systems
- Relevant for optimization (not integration)
- Powers of probability distributions emphasize the peaks
- If we are looking for a maximum within a lot of distracting peaks, this can help.

Annealing

- Define a temperature T , and a cooling schedule (black magic part)
- Lower temperatures correspond to emphasized maximal peaks.
 - Hence we exponentiate by $(1/T)$.

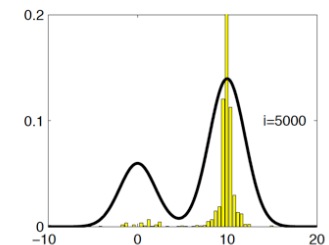
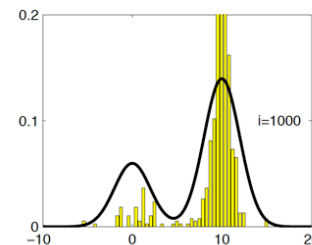
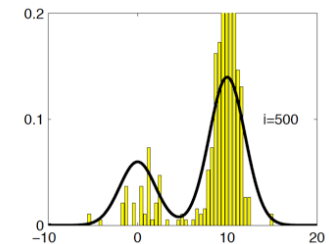
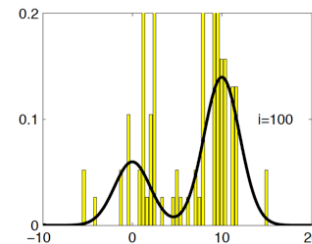
Annealing

1. Initialise $x^{(0)}$ and set $T_0 = 1$.
2. For $i = 0$ to $N - 1$
 - Sample $u \sim \mathcal{U}_{[0,1]}$.
 - Sample $x^* \sim q(x^* | x^{(i)})$.
 - If $u < \mathcal{A}(x^{(i)}, x^*) = \min \left\{ 1, \frac{p^{\frac{1}{T_i}}(x^*)q(x^{(i)} | x^*)}{p^{\frac{1}{T_i}}(x^{(i)})q(x^* | x^{(i)})} \right\}$

$$x^{(i+1)} = x^*$$
 - else

$$x^{(i+1)} = x^{(i)}$$
 - Set T_{i+1} according to a chosen cooling schedule.

(From Andrieu et al)



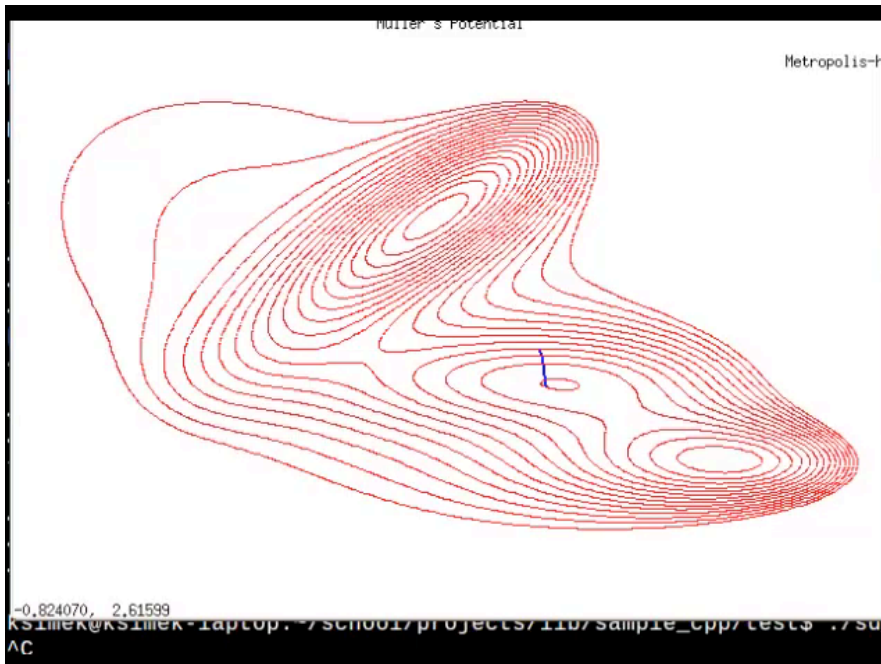
(From Andrieu et al)

Continuous versus discrete variables

- Derivatives of continuous distributions can tell you about the structure of your problem.
 - Opportunities for going much faster
- Naive approach is gradient ascent with added stochastic properties
 - Take a step, then perturb the result.
- Typical approach is to link the probability distribution to a potential energy function
 - Follow the system to find low energy (high probability)
 - Stochastic sampling via random momentum
 - An effective example method is Hybrid Monte Carlo

Hybrid Monte Carlo

- References include Andrieu et al. '01 and Neal'93.
- Link the probability distribution to a potential energy function
 - Alternate stochastic sampling with “dynamics”.
 - The dynamics follow the system to find low energy (high probability)
- HMC is an “auxiliary variable sampler”
 - Important trick
 - To sample $p(\mathbf{z})$ we sample $p(\mathbf{z},\mathbf{r})$ or $p(\mathbf{z},\mathbf{r}_1,\mathbf{r}_2,\dots)$
 - Ignore the auxiliary variables when we use the samples.



Hamiltonian Dynamics

$$p(\mathbf{z}) = \frac{1}{Z_p} \exp(-E(\mathbf{z}))$$

We equate \mathbf{z} with position, so $E(\mathbf{z})$ is the potential energy.

High probability \Leftrightarrow Low energy

$$E(\mathbf{z}) = -\log(Z_p) - \log(p(\mathbf{z})).$$

Hamiltonian Dynamics

Recall that the gradient, ∇ , is the vector of partial derivatives.

Recall from physics that force is the negative gradient of potential energy

From before $E(\mathbf{z}) = -\log(Z_p) - \log(p(\mathbf{z}))$

So force is $-\nabla E(\mathbf{z}) = \nabla(\log(p(\mathbf{z})))$

Hamiltonian Dynamics

$$p(\mathbf{z}) = \frac{1}{Z_p} \exp(-E(\mathbf{z})) \quad \text{and} \quad \nabla E(\mathbf{z}) = \nabla(-\log(p(\mathbf{z})))$$

Let \mathbf{r} be the momentum vector for the system. Denote the kinetic energy by $K(\mathbf{r})$.

$$K(\mathbf{r}) = \frac{1}{2} \|\mathbf{r}\|^2 = \frac{1}{2} \sum_i r_i^2 \quad (\text{We assume that mass is one}).$$

Hamiltonian Dynamics

$$H(\mathbf{z}, \mathbf{r}) = E(\mathbf{z}) + K(\mathbf{r}) \quad (\text{conserved})$$

Our distribution with auxiliary variables is

$$p(\mathbf{z}, \mathbf{r}) = \frac{1}{Z} \exp(-H(\mathbf{z}, \mathbf{r}))$$

Hamiltonian Dynamics

$$H(\mathbf{z}, \mathbf{r}) = E(\mathbf{z}) + K(\mathbf{r}) \quad (\text{conserved})$$

We follow \mathbf{z} according to H with a random \mathbf{r}

This can rapidly transport us towards a local minimum (but not to stay indefinitely) thus avoiding random walk.

To follow H , we observe that \mathbf{z} (position) changes proportional to \mathbf{r} (momentum), and \mathbf{r} changes proportion to force ($-\nabla E$).

$$\Delta \mathbf{r} \propto -\nabla E = \nabla(\log(p(\mathbf{z})))$$

$$\Delta \mathbf{z} \propto \mathbf{r}$$

Following Dynamics

In HMC we follow the dynamics for L time steps of size τ (tunable parameters).

In the "leap frog" method for each τ .

1. Take 1/2 step in \mathbf{r} .
2. Take a full step in \mathbf{z} .
3. Take 1/2 step in \mathbf{r} .

Following Dynamics

For L leap frog steps we have.

1. Take 1/2 step in \mathbf{r} .
2. $(L-1)$ times take a full steps in \mathbf{z} , then \mathbf{r} .
3. Take a full step in \mathbf{z} .
4. Take 1/2 step in \mathbf{r}

Following Dynamics

To take a full step in \mathbf{z} .

$$\mathbf{z}(\tau+1) = \mathbf{z}(\tau) + \varepsilon \cdot (\mathbf{r}(\tau))$$

(ε is the step size).

Following Dynamics

To take 1/2 step in \mathbf{r} .

$$\mathbf{r}\left(\tau + \frac{1}{2}\right) = \mathbf{r}(\tau) + \frac{1}{2}\varepsilon \cdot (-\nabla E(\mathbf{z}))$$

Following Dynamics

- After L steps of size t , we are at a new point with some bias of being at a lower potential energy (higher probability) and higher momentum.
- Momentum allows us to jump out of wells.

HMC dynamics step acceptance

- If our integration is perfect (i.e., in the limit as $t \rightarrow 0$) then energy is conserved.
 - Thus the value of distribution $p(\mathbf{z}, \mathbf{r})$ is the same after the dynamics.
- If we assume no integration errors, we simply accept this step
- If we want to account for error accumulation, we accept the result according to:

$$\min\left(1, \frac{p(\mathbf{z}^*, \mathbf{r}^*)}{p(\mathbf{z}, \mathbf{r})}\right) = \min\left(1, \exp\left(H(\mathbf{z}, \mathbf{r}) - H(\mathbf{z}^*, \mathbf{r}^*)\right)\right)$$

HMC stochastic step

- Typical instantiations sample the momentum variable
- Two common strategies
 - Sample the \mathbf{r} independently from a Gaussian
 - Sample \mathbf{r} from a Gaussian using Gibbs
- Note that in both of these cases the proposals are always accepted.

Putting it all together (A typical vision lab sampler)

- Discrete variables are sampled using (reversible jump) Metropolis Hastings.
- Continuous variables are sampled using stochastic dynamics (essentially hybrid Monte Carlo).
- Discrete variables typically control topology or components
 - The number of components and their type (block, cylinder)
 - How components are connected (branches from a stem)

A typical vision lab sampler

- Randomly proposing structure is too expensive because of the high rejection rate.
- Solution (part one) is to use data driven sampling
 - Proposals are conditioned on distributions computed before we begin using the data
 - For example, the probability of a corner being present in each point in the image.
- Solution (part two) is to delay acceptance
 - Adjust continuous parameters using stochastic dynamics so that the proposed structure is a good fit to the data.

A typical vision lab sampler

- We thus alternate between
 - (1) data driven proposals for new structure (or to switch or kill existing structure)
 - (2) exploring the continuous parameters of the structure
- Additional gains in optimization through having multiple samplers running in parallel exchange information