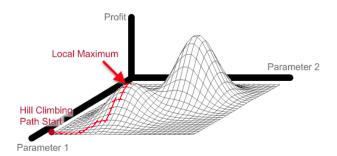
# Biostatistics 615/815 Lecture 20: Simulated Annealing Gibbs Sampling

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#### Local minimization methods

The problem with hill climbing is that it gets stuck on "local-maxima"

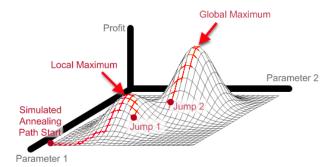


Images by Max Dama from

http://maxdama.blogspot.com/2008/07/trading-optimization-simulated.html

#### Global minimization with Simulated Annealing

Simulated Annealing can escape local minima with chaotic jumps



Images by Max Dama from

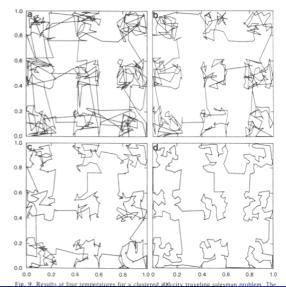
Simulated Annealing 

http://maxdama.blogspot.com/2008/07/trading-optimization-simulated.html

### **Example Applications**

- The traveling salesman problem (TSP)
  - Salesman must visit every city in a set
  - Given distances between pairs of cities
  - Find the shortest route through the set
- No polynomial time algorithm is known for finding optimal solution
- Simulated annealing or other stochastic optimization methods often provide near-optimal solutions.

# Examples of simulated annealing results



# Update scheme in Simulated Annealing

- Random walk by Metropolis criterion (1953)
- Given a configuration, assume a probability proportional to the Boltzmann factor

$$P_A = e^{-E_A/T}$$

• Changes from  $E_0$  to  $E_1$  with probability

$$\min\left(1, \frac{P_1}{P_0}\right) = \min\left(1, \exp\left(-\frac{E_1 - E_0}{T}\right)\right)$$

Given sufficient time, leads to equilibrium state

# Using Markov Chains

#### Markov Chain Revisited

The Markovian property

$$\Pr(q_t|q_{t-1}, q_{t-2}, \dots, q_0) = \Pr(q_t|q_{t-1})$$

Transition probability

$$\theta_{ij} = \Pr(q_t = j | q_{t-1} = i)$$

#### Simulated Annealing using Markov Chain

- Start with some state  $q_t$ .
- Propose a changed  $q_{t+1}$  given  $q_t$
- Decide whether to accept change based on  $\theta_{q_tq_{t+1}}$ 
  - Decision is based on relative probabilities of two outcomes

#### Key requirements

- Irreducibility: it is possible to get any state from any state
  - There exist t where  $\Pr(q_t = j | q_o = i) > 0$  for all (i, j).
- Aperiodicity: return to the original state can occur at irregular times

$$\gcd\{t : \Pr(q_t = i | q_0 = i) > 0\} = 1$$

These two conditions guarantee the existence of a unique equilibrium distribution

### Equilibrium distribution

- Starting point does not affect results
- The marginal distribution of resulting state does not change
- Equilibrium distribution  $\pi$  satisfies

$$\pi = \lim_{t \to \infty} \Theta^{t+1}$$
$$= (\lim_{t \to \infty} \Theta^t) \Theta$$
$$= \pi \Theta$$

• In Simulated Annealing,  $\Pr(E) \propto e^{-E/T}$ 

## Simulated Annealing Recipes

- 1 Select starting temperature and initial parameter values
- Randomly select a new point in the neighborhood of the original
- 3 Compare the two points using the Metropolis criterion
- 4 Repeat steps 2 and 3 until system reaches equilibrium state
  - In practice, repeat the process N times for large N.
- **5** Decrease temperature and repeat the above steps, stop when system reaches frozen state

#### Practical issues

- The maximum temperature
- Scheme for decreasing temperature
- Strategy for proposing updates
  - For mixture of normals, suggestion of Brooks and Morgan (1995) works well
  - Select a component to update, and sample from within plausible range

```
#ifndef __TRAVERSE_2D_H
#define TRAVERSE 2D H
#include <vector>
#include <algorithm>
#include <cstdlib>
#include <cmath>
class Traverse2D {
protected:
  double distance;
  bool stale;
public:
  std::vector<double> xs;
  std::vector<double> vs;
  std::vector<int> order;
```

```
Traverse2D() : distance(-1), stale(true) {}
Traverse2D(std::vector<double>& _xs, std::vector<double>& _ys)
           : xs(xs), ys(ys), stale(true) {
 int n = (int)xs.size():
 if ( n != ys.size() ) abort();
 for(int i=0: i < n: ++i) {
   order.push back(i);
int numPoints() { return (int)order.size(); }
void addPoint(double x, double y) {
 xs.push back(x);
 ys.push back(y);
 order.push back((int)order.size());
}
```

```
void initOrder() {
 stale = true:
  std::sort( order.begin(), order.end() );
bool nextOrder() {
 stale = true;
 return std::next permutation( order.begin(), order.end() );
}
void shuffleOrder() {
 stale = true;
 std::random shuffle( order.begin(), order.end() );
void swapOrder(int x, int y) {
 stale = true;
 int tmp = order[x]:
 order[x] = order[y];
 order[v] = tmp;
```

```
double getDistance() {
    if ( stale ) {
      int n = (int)order.size();
      distance = 0:
      for(int i=1; i < n; ++i) {</pre>
        distance += sqrt(
            (xs[order[i]]-xs[order[i-1]])*(xs[order[i]]-xs[order[i-1]])
            + (ys[order[i]]-ys[order[i-1]])*(ys[order[i]]-ys[order[i-1]]) );
      stale = false:
    return distance;
};
#endif // TRAVERSE 2D H
```

# Implementing TSP : main()

```
int main(int argc, char** argv) {
  if ( argc != 2 ) {
    std::cerr << "Usage: TSP [infile]" << std::endl;</pre>
    return -1;
  }
  Matrix615 < double > xy(argv[1]);
  int n = xy.rowNums();
  if ( xy.colNums() != 2 ) {
    std::cerr << "Input matrix does not have exactly two columns" << std::endl;</pre>
    return -1;
  // build graph from file
  Traverse2D graph:
  for(int i=0; i < n; ++i) {</pre>
    graph.addPoint(xy.data[i][0], xy.data[i][1]);
  }
```

## Implementing TSP : main()

```
int start = 0, finish = 0, nperm = 0;
double duration = 0, minDist = DBL MAX, maxDist = 0, sumDist = 0;
std::vector<int> minOrder;
start = clock();
graph.initOrder(); // initialize order
do {
  double d = graph.getDistance();
  sumDist += d; ++nperm;
  if ( d > maxDist ) maxDist = d;
  if ( d < minDist ) {</pre>
    minDist = d;
    minOrder = graph.order:
} while ( graph.nextOrder() );
finish = clock();
duration = (double)(finish-start)/CLOCKS PER SEC;
```

## Implementing TSP : main()

```
std::cout << "-----" << std::endl:
std::cout << "Minimum distance = " << minDist << std::endl:</pre>
std::cout << "Maximum distance = " << maxDist << std::endl;</pre>
std::cout << "Mean distance = " << sumDist/nperm << std::endl;</pre>
std::cout << "Fxhaustive search duration = " << duration << " seconds"
        << std::endl;
std::cout << "-----" << std::endl:
start = clock();
runTSPSA(graph, 1e-6); // run Simulated Annealing
finish = clock();
duration = (double)(finish-start)/CLOCKS PER SEC;
std::cout << "SA distance = " << graph.getDistance() << std::endl;</pre>
std::cout << "SA search Duration = " << duration << " seconds" << std::endl;</pre>
std::cout << "-----" << std::endl:
return 0;
```

}

#### Implementing TSP : runTSPSA()

```
#define MAX TEMP 1000
#define N ITER 1000
double runTSPSA(Traverse2D& graph, double eps) {
  srand(std::time(0));
  graph.shuffleOrder();
  double temperature = MAX TEMP;
  double prevDist = graph.getDistance();
  int n = graph.numPoints();
  while( temperature > eps ) {
    for(int i=0: i < N ITER: ++i) {</pre>
      int i1 = (int)floor( rand()/(RAND MAX+1.) * n);
      int i2 = (int)floor( rand()/(RAND MAX+1.) * n);
      graph.swapOrder(i1,i2);
      double newDist = graph.getDistance();
      double diffDist = newDist-prevDist;
```

### Implementing TSP: runTSPSA()

```
if ( diffDist < 0 ) {</pre>
    prevDist = newDist;
  else {
    double p = rand()/(RAND_MAX+1.);
    if ( p < exp(0-diffDist/temperature) ) {</pre>
      prevDist = newDist;
    else {
      graph.swapOrder(i1,i2);
temperature *= 0.90;
```

\$ cat tsp.10.in.txt

```
-2.30963348991357 0.0773267767084084

-1.13260001198939 0.194723763831079

-0.47887704546568 -1.49043206086804

-1.14183413926286 -0.386463669289195

-0.0684871826034848 0.362329163828058

-1.28322395967065 -0.173892955683618

-0.684913927794102 0.0967915142130205

1.87577059887638 -0.229129514295367

-0.796217725319515 1.77563911372358

0.936967861258253 -0.103803298997143
```

```
$ cat tsp.11.in.txt
-0.636066544886696 2.25053338615707
0.0860940972604061 0.231139523090642
0.219459494449743 -0.518180472158068
0.0566391380933713 -1.10184323809265
-0.300676076997908 -0.765625163407885
2.64204087640419 1.29479579271570
0.152911487506204 0.228909136397270
-0.933319389247532 -0.846940788411644
-0.447908403019059 -1.16451734926683
1.61047052169711 1.66393401261582
-1.16737084487488 1.04729096252209
```

```
$ ./TSP tsp.11.in.txt

Minimum distance = 9.14731

Maximum distance = 28.1806

Mean distance = 20.3772

Exhaustive search duration = 192.85 seconds

SA distance = 9.14731

SA search Duration = 1.78 seconds

SA distance = 3.52509

SA search Duration = 0.514433 seconds
```

### Simulated Annealing for Gaussian Mixtures

```
class normMixSA {
public:
  int k;
                             // # of components
  int n:
                             // # of data
  std::vector<double> data: // observed data
  std::vector<double> pis; // pis
  std::vector<double> means: // means
  std::vector<double> sigmas; // sds
  double llk;
                              // current likelihood
  normMixSA(std::vector<double>& data, int k); // constructor
  void initParams();
                                                  // initialize parameters
  double updatePis(double temperature);
  double updateMeans(double temperature, double lo, double hi);
  double updateSigmas(double temperature, double sdlo, double sdhi);
                                                  // run Simulated Annealing
  double runSA(double eps);
  static int acceptProposal(double current, double proposal, double temperature);
};
```

# **Evaluating Proposals in Simulated Annealing**

### Updating Means and Variances

- Select component to update at random
- Sample a new mean (or variance) within plausible range for parameter
- Decide whether to accept proposal or not

## **Updating Means**

```
double normMixSA::updateMeans(double temperature, double min, double max) {
  int c = randn(0,k) // select a random integer between 0..(k-1)
  double old = means[c]: // save the old mean for recovery
  means[c] = randu(min, max); // update mean and evaluate the likelihood
  double proposal = 0-NormMix615::mixLLK(data, pis, means, sigmas);
  if ( acceptProposal(llk, proposal, temperature) ) {
   11k = proposal;
                  // if accepted, keep the changes
 else {
   means[c] = old;
                            // if rejected, rollback the changes
  return 11k;
```

## **Updating Component Variances**

```
double normMixSA::updateSigmas(double temperature, double min, double max) {
 int c = randn(0,k) // select a random integer between 0..(k-1)
 double old = sigmas[c]: // save the old mean for recovery
 sigmas[c] = randu(min, max); // update a component and evaluate the likelihood
 double proposal = 0-NormMix615::mixLLK(data, pis, means, sigmas);
 if ( acceptProposal(llk, proposal, temperature) ) {
   else {
   sigmas[c] = old;
                          // if rejected, rollback the changes
 return 11k;
```

# **Updating Mixture Proportions**

- Mixture proportions must sum to 1.0
- When updating one proportion, must take others into account
- Select a component at random
  - Increase or decrease probability by up to 25%
  - Rescale all proportions so they sum to 1.0

## Updating Mixture Proportions

```
double normMixSA::updatePis(double temperature) {
  std::vector<double> pisCopy = pis; // make a copy of pi
  int c = randn(0,k);  // select a component to update
  pisCopy[c] *= randu(0.8,1.25); // update the component
  // normalize pis
  double sum = 0.0;
  for(int i=0: i < k: ++i)
    sum += pisCopv[i];
  for(int i=0; i < k; ++i)</pre>
    pisCopv[i] /= sum:
  double proposal = 0-NormMix615::mixLLK(data, pisCopy, means, sigmas);
  if ( acceptProposal(llk, proposal, temperature) ) {
    11k = proposal:
    pis = pisCopy; // if accepted, update pis to pisCopy
  return 11k;
```

### Initializing parameters

```
void normMixSA::initParams() {
  double sum = 0, sqsum = 0;
  for(int i=0; i < n; ++i) {</pre>
    sum += data[i];
    sqsum += (data[i]*data[i]);
  double mean = sum/n;
  double sigma = sqrt(sqsum/n - sum*sum/n/n);
  for(int i=0; i < k; ++i) {</pre>
    pis[i] = 1./k;
                    // uniform priors
    means[i] = data[rand() % n]; // pick random data points
    sigmas[i] = sigma;  // pick uniform variance
```

# Putting things together

```
double normMixSA::runSA(double eps) {
 initParams();
                                     // initialize parameter
 llk = 0-NormMix615::mixLLK(data, pis, means, sigmas); // initial likelihood
 double lo = min(data), hi = max(data); // min(), max() can be implemented
 double sd = stdev(data):
                                  // stdev() can also be implemented
 double sdhi = 10.0 * sd, sdlo = 0.1 * sd;
 while( temperature > eps ) {
   for(int i=0: i < 1000: ++i) {
     switch( randn(0,3) ) { // generate a random number between 0 and 2
     case 0:
                           // update one of the 3*k components
       11k = updatePis(temperature); break;
     case 1:
       11k = updateMeans(temperature, lo, hi); break;
     case 2:
       11k = updateSigmas(temperature, sdlo, sdhi); break;
     }
   temperature *= 0.90; // cool down slowly
 return 11k;
```

### Running examples

```
user@host:~/> ./mixSimplex ./mix.dat
Minimim = 3043.46, at pi = 0.667271,
between N(-0.0304604,1.00326) and N(5.01226,0.956009)

user@host:~/> ./mixEM ./mix.dat
Minimim = -3043.46, at pi = 0.667842,
between N(-0.0299457,1.00791) and N(5.0128,0.913825)

user@host:~/> ./mixSA ./mix.dat
Minimim = 3043.46, at pi = 0.667793,
between N(-0.030148,1.00478) and N(5.01245,0.91296)
```

# Comparisons

#### 2-component Gaussian mixtures

- Simplex Method : 306 Evaluations
- E-M Algorithm : 12 Evaluations
- Simulated Annealing :  $\sim 100,000$  Evaluations

#### For higher dimensional problems

- Simplex Method may not converge, or converge very slowly
- E-M Algorithm may stuck at local maxima when likelihood function is multimodal
- Simulated Annealing scale robustly with the number of dimensions.

# Optimization Strategies

- Single Dimension
  - Golden Search
  - Parabolic Approximations
- Multiple Dimensions
  - Simplex Method
  - E-M Algorithm
  - Simulated Annealing
  - Gibbs Sampling

### Gibbs Sampler

- Another MCMC Method
- Update a single parameter at a time
- Sample from conditional distribution when other parameters are fixed

# Gibbs Sampler Algorithm

- **1** Consider a particular choice of parameter values  $\lambda^{(t)}$ .
- 2 Define the next set of parameter values by
  - Selecting a component to update, say *i*.
  - Sample value for  $\lambda_i^{(t+1)}$ , from  $p(\lambda_i|x,\lambda_1^{(t)},\cdots,\lambda_{i-1}^{(t)},\lambda_{i+1}^{(t)},\cdots,\lambda_k^{(t)})$ .

## An alternative Gibbs Sampler Algorithm

- **1** Consider a particular choice of parameter values  $\lambda^{(t)}$ .
- 2 Define the next set of parameter values by
  - Sample value for  $\lambda_1^{(t+1)}$ , from  $p(\lambda_1|x,\lambda_2,\lambda_3,\cdots,\lambda_k)$ .
  - Sample value for  $\lambda_2^{(t+1)}$ , from  $p(\lambda_2|x,\lambda_1,\lambda_3,\cdots,\lambda_k)$ .
  - . . . .
  - Sample value for  $\lambda_k^{(t+1)}$ , from  $p(\lambda_k|x,\lambda_1,\lambda_2,\cdots,\lambda_{k-1})$ .

# Gibbs Sampling for Gaussian Mixture

#### Using conditional distributions

- Observed data :  $\mathbf{x} = (x_1, \dots, x_n)$
- Parameters :  $\lambda = (\pi_1, \cdots, \pi_k, \mu_1, \cdots, \mu_k, \sigma_1^2, \cdots, \sigma_k^2)$ .
- Sample each  $\lambda_i$  from conditional distribution not very straightforward

#### Using source of each observations

- Observed data :  $\mathbf{x} = (x_1, \dots, x_n)$
- Parameters :  $\mathbf{z} = (z_1, \dots, z_n)$  where  $z_i \in \{1, \dots, k\}$ .
- Sample each  $z_i$  conditioned by all the other **z**.

# Update procedure in Gibbs sampler

$$\Pr(z_j = i | x_j, \lambda) = \frac{\pi_i \mathcal{N}(x_j | \mu_i, \sigma_i^2)}{\sum_l \pi_l \mathcal{N}(x_j | \mu_l, \sigma_l^2)}$$

- Calculate the probability that the observation is originated from a specific component
- For a random  $j \in \{1, \dots, n\}$ , sample  $z_j$  based on the current estimates of mixture parameters.

#### Initialization

- Must start with an initial assignment of component labels for each observed data point
- A simple choice is to start with random assignment with equal probabilities

### The Gibbs Sampler

- Select initial parameters
- Repeat a large number of times
  - Select an element
  - Update conditional on other elements