# Parallel Computing Approaches for Model Comparison

Tutorial

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# OUTLINE

- 1. Introduction
- 2. Algorithms
- 3. Shared memory approach
- 4. GPU approach
- 5. Further suggestions and conclusion

Introduction

#### OVERVIEW

- · Model comparison is one level of Bayesian inference
- Model comparison algorithms can take advantage of parallel computing
- · Many parallel programming models and hardware platforms exist
- · Some problems can be better suited for certain software and hardware combinations
- Tutorial will cover three algorithm/programming model/hardware combinations. Starting place for exploring other combinations.

#### INTRODUCTION: MODEL COMPARISON

$$\underline{p(\boldsymbol{\Theta}_{i}|M_{i},\boldsymbol{D},I)}_{\text{Posterior}} = \underbrace{\frac{prior}{p(\boldsymbol{\Theta}_{i}|M_{i},I)}\underbrace{\frac{\text{Likelihood}}{p(\boldsymbol{D}|M_{i},\boldsymbol{\Theta}_{i},I)}}_{\text{Evidence or model likelihood}} (1)$$

#### INTRODUCTION: MODEL COMPARISON

$$O_{ji} = \frac{p(M_j | \mathbf{D}, I)}{p(M_i | \mathbf{D}, I)} = \underbrace{\frac{p(M_j | I)}{p(M_i | I)}}_{\text{Prior odds}} \times \underbrace{\frac{p(\mathbf{D} | M_j, I)}{p(\mathbf{D} | M_i, I)}}_{\text{Evidence ratio}} \tag{3}$$

If model priors are equal,

$$O_{ji} = \frac{p(\mathbf{D}|M_j, I)}{p(\mathbf{D}|M_i, I)} \tag{4}$$

$$p(\mathbf{D}|M_i, I) = \int_{\tilde{\boldsymbol{\Theta}}} p(\boldsymbol{\Theta}_i|M_i, I) p(\mathbf{D}|M_i, \boldsymbol{\Theta}_i, I) \, \mathrm{d}\boldsymbol{\Theta}$$
 (5)

#### INTRODUCTION: PARALLEL COMPUTING

#### Main idea

Break task into pieces that can be worked on concurrently

# Why?

- More efficient use of resources
- · Get results more quickly
- $\cdot$  Solve massive problems that are intractable otherwise

#### INTRODUCTION: PARALLEL COMPUTING

# Programming models (a.k.a., useful abstractions)

- Shared memory
- Distributed memory
- Hybrid
- · Single program, multiple data (SPMD)
- Multiple program, multiple data (MPMD)

# Introduction: Parallel computing



SGI cluster at MCSR

# INTRODUCTION: PARALLEL COMPUTING



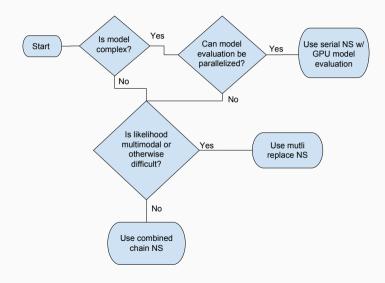
Intel Xeon Phi

# Introduction: Parallel computing



Nvidia Tesla GPU Accelerator

# INTRODUCTION: CHOOSING A METHOD



Algorithms

# **ALGORITHMS**

- Nested sampling
  - Combined chains
  - Multiple replacement
- Thermodynamic integration
- Reversible jump MCMC
- Sequential Monte Carlo

#### **ALGORITHMS**

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## **NESTED SAMPLING BASICS**

Nested sampling reparameterizes the model evidence integral

$$\mathcal{Z} = \int \pi(\boldsymbol{\Theta}) L(\boldsymbol{\Theta}) \, \mathrm{d}\boldsymbol{\Theta} \tag{6}$$

$$L(\boldsymbol{\Theta}) = \int_0^{L(\boldsymbol{\Theta})} d\mathcal{L} \tag{7}$$

$$\mathcal{Z} = \int \pi(\boldsymbol{\Theta}) \left[ \int_0^{L(\boldsymbol{\Theta})} d\mathcal{L} \right] d\boldsymbol{\Theta}$$
 (8)

# **NESTED SAMPLING BASICS**

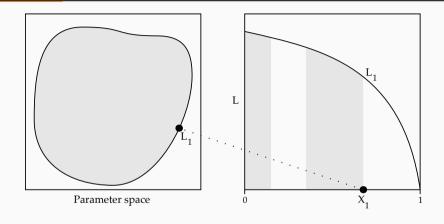
$$\mathcal{Z} = \int_0^\infty \left[ \int_{\{\boldsymbol{\Theta}: L(\boldsymbol{\Theta}) > \mathcal{L}\}} \pi(\boldsymbol{\Theta}) \, d\boldsymbol{\Theta} \right] \, d\mathcal{L}$$
 (9)

$$X(\mathcal{L}) = \int_{\{\boldsymbol{\Theta}: L(\boldsymbol{\Theta}) > \mathcal{L}\}} \pi(\boldsymbol{\Theta}) \, d\boldsymbol{\Theta}$$
 (10)

$$\mathcal{Z} = \int_0^\infty X(\mathcal{L}) \, \mathrm{d}\mathcal{L} \tag{11}$$

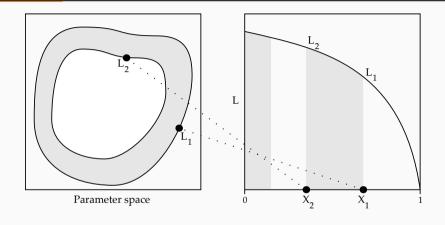
$$\mathcal{Z} = \int_0^1 \mathcal{L}(X) \, \mathrm{d}X. \tag{12}$$

# NESTED SAMPLING BASICS: PRIOR MASS AND LIKELIHOOD



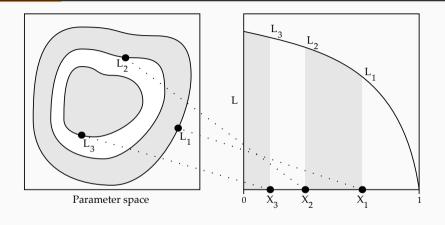
(From Skilling '06)

# NESTED SAMPLING BASICS: PRIOR MASS AND LIKELIHOOD



(From Skilling '06)

# NESTED SAMPLING BASICS: PRIOR MASS AND LIKELIHOOD

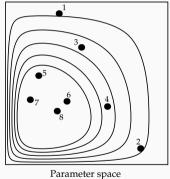


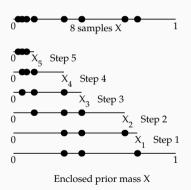
(From Skilling '06)

## **NESTED SAMPLING BASICS: PROCEDURE**

- 1. Generate N live samples from the unconstrained prior
- 2. Set Z=0 and  $X_0=1$
- 3. Record the lowest likelihood among the N live samples as  $L_i$
- 4. Estimate  $X_i$  corresponding to  $L_i$
- 5. Set  $w_i = X_{i-1} X_i$
- 6. Increment Z by  $L_i w_i$
- 7. (Optional) Save least-likelihood live sample in list of "dead" samples
- 8. Replace the least-likelihood live sample with one sampled from the prior, constrained by  ${\cal L}_i$
- 9. If halting condition is not met, go to 3

# **NESTED SAMPLING BASICS: PROCEDURE**





# Nested sampling basics: Estimating $X_i$

The likelihood  $L_i$  at a given parameter vector  $\pmb{\Theta}_i$  can be computed exactly, but the associated prior mass  $X_i$  cannot

At every step of nested sampling, the live samples' prior mass are distributed as U[0,X(L)]

Order statistics of the uniform distribution show

$$t_i = \frac{X_i}{X_{i-1}} \sim \text{Beta}(N, 1) \tag{13}$$

$$X_i = \prod_{k=1}^i t_k \tag{14}$$

# Nested sampling basics: Estimating $X_i$

$$E(\log t_i) = -1/N \tag{15}$$

$$X_i \approx \exp(-i/N)$$
 (16)

$$Z \approx \sum_{i=1}^{m} (X_{i-1} - X_i) L_i$$
 (17)

#### NESTED SAMPLING: PARALLEL IMPLEMENTATIONS

- Combined chain nested sampling Combine multiple independent serial nested sampling results
- **Multiple replacement nested sampling** Discard and replace multiple samples at each likelihood threshold
- Parallel model evaluation Use serial nested sampling, but parallelize the model computation

#### **NESTED SAMPLING: COMBINED CHAIN**

# First parallel approach

- $\cdot$  Run M independent nested sampling processes, each using N live samples
- Resulting discarded samples can be combined and sorted by likelihood; prior mass estimate is same as in case of one nested sampling process with  $M \times N$  live samples
- Proofs of the correctness of this approach are available in our 2017 paper.
- Theoretical speed up of  $\mathcal{O}(M)$  over serial method
- · MIMD

#### NESTED SAMPLING: MULTIPLE REPLACEMENT

# Second parallel approach

- $\cdot$  Instead of discarding and replacing one sample, discard and replace R samples at each likelihood constraint
- Initially proposed by Burkoff, et al., in 2012
- Our 2014 paper found that in order to maintain the same level of precision in evidence estimate,  $\sqrt{R}N$  live samples must be used when R samples are discarded and replaced for each likelihood constraint
- Theoretical speed up of  $\mathcal{O}(\sqrt{R})$  over serial method
- MIMD

#### **NESTED SAMPLING: PARALLEL MODEL EVALUATION**

# Third parallel approach

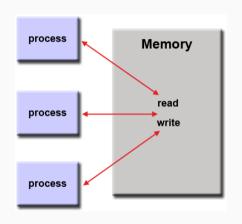
- · Use serial nested sampling
- · Parallelize model evaluation
- Ideal for cases with complex model equations
- Allows use of GPU for computation
- · SIMD

Shared memory approach

- · Multiple processors share same memory
- Examples
  - · Multi-core CPUs
  - · Intel Xeon Phi
- Advantages
  - · Processes can easily share data
  - · Relatively easy to modify programs to use
- Disadvantages
  - Without proper safeguards, data can be corrupted by competing processes
  - Scalability

#### No threads

- · Multiple processes used
- Very straightforward, but no standard approach across platforms



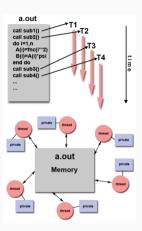
(From https://computing.llnl.gov/tutorials/parallel\_comp/)

#### With threads

pthreads Library based. Requires manual creation, starting, and synchronization of threads

OpenMP Compiler directive based.

Manual controls available, but simple complier directives can be used to parallelize some serial code.



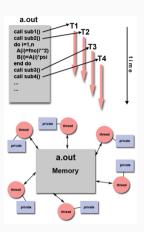
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 $(From \ https://computing.llnl.gov/tutorials/parallel\_comp/)$ 

#### SHARED MEMORY: XEON PHI

Two existing versions for general applications

# Knight's Corner (x100)

- · PCI express card, now discontinued
- Runs only on certain motherboards
- 4-way simultaneous multithreading per core
- 57-61 x86-64 cores
- · 512-bit SIMD units
- 512 KB L2 cache per core
- Supports offloading or native execution

# Knight's Landing (x200)

- · Bootable, mainboard chip
- Mostly available from workstation system builders. Expensive
- 64–72 Intel Atom cores
- Two 512-bit vector units per core
- AVX-512 SIMD instructions
- Native execution only

# COMBINED-CHAIN NS ON XEON PHI

- In combined-chain, most of the code can be made trivially parallel, so works well natively on the Phi
- This example code is in C++, but mostly uses C idioms
- Xeon Phi requires Intel C++ compiler. Free for students and in other limited cases

## COMBINED-CHAIN NS ON XEON PHI

```
Sample struct and static arrays
typedef struct
    declspec(align(64)) double theta[NUM PARAMS];
    declspec(align(64)) double logL;
    declspec(align(64)) double logWt;
} sample:
Dynamic arrays (Intel C++ compiler only)
#include <malloc.h>
sample ** samples = (sample**) mm malloc(SIZE * sizeof(sample). 64):
// Do some things
mm free(samples);
```

## COMBINED-CHAIN NS ON XEON PHI

Start and collect results from each NS instance

## Collecting result samples

```
int q = 0:
for (int i = 0; i < NUM CHAINS; i++) {</pre>
    for (int j = 0; j < total samples[i]; j++) {</pre>
        for (int k = 0; k < NUM PARAMS; k++) {
            samples[q].theta[k] = sample list 2d[i][i].theta[k];
        samples[q].logL = sample list 2d[i][j].logL;
        samples[q].logWt = 0.0:
        q++;
```

Sorting samples and computing evidence

```
gsort(samples. *sample count, sizeof(sample), comparator);
double mean log t = 1.0 / (NUM CHAINS * NUM LIVE SAMPLES):
double logwidth = log(1.0 - exp(-mean log t));
*logZ = -1.0 * std::numeric limits<double>::max():
*H = 0.0:
for (int i = 0; i < *sample count; i++) {</pre>
    samples[i].logWt = logwidth + samples[i].logL;
    logZnew = log plus(*logZ, samples[i].logWt):
    *H = exp(samples[i].logWt - logZnew) * samples[i].logL +
        exp(*logZ - logZnew) * (*H + *logZ) - logZnew;
    *logZ = logZnew;
    logwidth -= mean log t:
```

# Sample comparator function

```
int comparator(const void * lhs. const void * rhs)
    double diff:
    diff = ((sample *)lhs)->logL - ((sample *)rhs)->logL:
    if (diff > 0)
        return 1:
    else if (diff < 0)
        return -1;
    else
        return 0;
```

#### SIMD block of likelihood function

```
#pragma omp simd
#pragma vector aligned
for (int i = 0; i < NUM DATA; i++)
#pragma vector aligned
   for (int j = 0; j < NUM ATOMS; j++)
        mock[i] += A[i] * cos(2 * PI * f[i] * time[i]) +
            B[j] * sin(2 * PI * f[j] * time[i]);
    sq error[i] = (mock[i] - data[i]) * (mock[i] - data[i]);
```

## COMBINED-CHAIN NS ON XEON PHI

## Other implementation considerations

- main function loads data and sets nested sampling parameters, then calls manager function
- manager sets up temporary arrays for collecting results from each nested sampling function, then calls nested\_sampling within OpenMP
- Make sure that any temporary arrays used (in explore function, likelihood function, etc.) are declared with the correct alignment

## SHARED MEMORY: MULTI-CORE CPU

- · Common in consumer-grade desktops, laptops, and smartphones
- · One processor package, two or more independent processing units
- For development, I used an Intel Xeon E5-1603 v3
  - · 2.80 GHz
  - 4 cores
  - · 10 MB SmartCache

## MULTIPLE REPLACEMENT NS ON CPU

- $\boldsymbol{\cdot}$  More serial-only portions of this code, so CPU is perhaps better fit
- This example code is idiomatic C++

#### MULTIPLE REPLACEMENT NS ON CPU

```
Model class
class Model
    public:
        Model(std::vector<double> data_in);
        double compute log likelihood(std::vector<double> theta);
        void compute sq error(double * A, double * B, double * f,
                double * sq error):
        void set data(std::vector<double> data in);
        std::vector<double> get data();
    private:
        std::vector<double> data:
};
```

```
class Sample
    public:
        Sample(Model model):
        void set theta(std::vector<double> theta in);
        std::vector<double> get theta();
        void set logL(double logL in);
        double get logL();
        void set logWt(double logWt_in);
        double get logWt();
    private:
        std::vector<double> theta;
        double logL;
        double logWt;
```

#### MULTIPLE REPLACEMENT NS ON CPU

Within the nested\_sampling funcion:
std::vector<Sample> live\_samples;
for (int i = 0; i < NUM\_LIVE\_SAMPLES; i++)
{
 Sample live\_sample(model);
 live\_samples.push\_back(live\_sample);</pre>

```
Sort at each likelihood threshold
std::sort(live_samples.begin(), live_samples.end(), sample_comp);
Comparator function
double sample_comp(Sample a, Sample b)
{
    return (a.get_logL() < b.get_logL());
}</pre>
```

## Choose surviving samples to evolve

```
std::vector<int> surviving idxs(NUM LIVE SAMPLES - NUM REP);
for (int i = NUM REP; i < NUM LIVE SAMPLES; i++)</pre>
    surviving idxs[i - NUM REP] = i;
std::random shuffle(surviving idxs.begin(). surviving idxs.end());
std::vector<int> copy idxs(NUM REP);
for (int i = 0; i < NUM REP; i++)
    copy idxs[i] = surviving idxs[i];
```

## Evolve each sample simultaneously

```
std::vector<std::vector<double> > theta in vec(NUM REP);
std::vector<std::vector<double> > theta out vec(NUM REP):
for (int i = 0: i < NUM REP: i++)
   theta in vec[i] = live samples[copy idxs[i]].get theta();
#pragma omp parallel for
for (int i = 0; i < NUM REP; i++)
   mcmc explore(theta in vec[i], logLstar, model, live samples,
        theta out vec[i]);
```

GPU approach

#### **OVERVIEW**

- Main idea: use hardware originally meant for graphics computations for general purpose computation
- · Relative to CPUs, GPUs have many more, slower cores
- Parallel portion of program needs to be SIMD for maximum performance
- Several languages/libraries exist for running code on GPUs
  - · Khronos Group's OpenCL
  - · Nivida's CUDA
  - · Apple's Metal
- We'll focus on OpenCL

## **OPENCL**

- OpenCL code runs on many device types (AMD and Nvidia GPUs, CPUs, even the Xeon Phi)
- · Kernels contain code that is run by each work item
- Work items are grouped into work groups
- · Each work item operates on a different piece of data stream
- · Branches within kernel are inefficient
- On Nvidia GPUs, CUDA can perform better for some code, but 1-to-1 comparison is difficult

## PARALLEL MODEL EVALUATION ON GPU

- Nested sampling portion of code is straightforward
- Complexity arises in problem-specific code
- Example code is idiomatic C++ and uses OpenCL
- There is a lot more work involved in adapting code for OpenCL, compared with the previous examples

```
Model class
class Model
    public:
        Model(std::vector<double> data in. cl::Buffer buffer A.
             cl::Buffer buffer B, cl::Buffer buffer f,
             cl::Buffer buffer data, cl::Buffer buffer sg error,
             cl::CommandQueue queue, cl::Context context,
             cl::Kernel kernel);
        double compute log likelihood(std::vector<double> theta):
        void set data(std::vector<double> data in);
        std::vector<double> get data();
```

#### PARALLEL MODEL EVALUATION ON GPU

## Model class, continued

```
private:
        std::vector<double> data:
        int num data:
        cl::Buffer buffer_A;
        cl::Buffer buffer B;
        cl::Buffer buffer f;
        cl::Buffer buffer data;
        cl::Buffer buffer sq error;
        cl::CommandQueue queue:
        cl::Context context;
        cl::Kernel kernel compute sq error:
};
```

# Setup

```
std::vector<cl::Platform> all_platforms;
cl::Platform::get(&all_platforms);
cl::Platform default_platform = all_platforms[0];
std::vector<cl::Device> all_devices;
default_platform.getDevices(CL_DEVICE_TYPE_ALL, &all_devices);
cl::Device default_device = all_devices[0];
cl::Context context({ default_device });
cl::Program::Sources sources;
```

#### Kernel

```
std::string kernel code =
  "kernel void compute sq error("
    "global const double * A, global const double * B,"
    "global const double * f, global const double * data,"
    "const int num data, const int num atoms,"
    "global double * sq error)"
 " { "
       double pi = 3.14159265359;"
       int i = get global id(0):"
       double mock = 0.0;"
       sq error[i] = 0.0;"
       double time = data[num data + i];"
```

## Kernel, continued

```
" for (int j = 0; j < num_atoms; j++)"
" {"
"         mock += A[j] * cos(2 * pi * f[j] * time) +"
"         B[j] * sin(2 * pi * f[j] * time);"
"     }"
" sq_error[i] = (mock - data[i]) * (mock - data[i]);"
"}";</pre>
```

# Final setup

```
sources.push back(
    std::make pair(kernel code.c str(), kernel code.length()));
cl::Program program(context, sources);
program.build(all devices):
cl::Buffer buffer A(context, CL MEM READ WRITE, sizeof(double) * NA);
cl::Buffer buffer B(context, CL MEM READ WRITE, sizeof(double) * NA);
cl::Buffer buffer f(context, CL MEM READ WRITE, sizeof(double) * NA);
cl::Buffer buffer data(context, CL MEM READ WRITE,
    sizeof(double) * (2 * num data + 1));
cl::Buffer buffer_sq_error(context, CL MEM READ WRITE.
    sizeof(double) * num data);
```

## Final setup, continued

## Log-likelihood function

```
double Model::compute log likelihood(std::vector<double> theta)
    // Scale the parameters before sending them to the device
    double A[NUM ATOMS]:
    double B[NUM ATOMS]:
    double f[NUM ATOMS];
    const int param per atom = NUM PARAMS / NUM ATOMS;
    for (int i = 0; i < NUM ATOMS; i++)
        A[i] = (AMAX - AMIN) * theta[i * param per atom + 0] + AMIN;
        B[i] = (AMAX - AMIN) * theta[i * param per atom + 1] + AMIN;
        f[i] = (FMAX - FMIN) * theta[i * param per atom + 2] + FMIN:
```

#### PARALLEL MODEL EVALUATION ON GPU

# Log-likelihood function, continued

```
queue.enqueueWriteBuffer(buffer A, CL TRUE, 0,
    sizeof(double) * NUM ATOMS. A):
queue.enqueueWriteBuffer(buffer B, CL TRUE. 0.
    sizeof(double) * NUM ATOMS. B):
queue.engueueWriteBuffer(buffer f, CL TRUE, 0,
    sizeof(double) * NUM ATOMS, f);
kernel compute sq error.setArg(0, buffer A);
kernel compute sq error.setArg(1, buffer B);
kernel compute sq error.setArg(2. buffer f):
kernel compute sq error.setArg(3, buffer data);
kernel compute sq error.setArg(4, num data);
kernel compute sq error.setArg(5, NUM ATOMS):
kernel compute sq error.setArg(6, buffer sq error);
```

#### PARALLEL MODEL EVALUATION ON GPU

## Log-likelihood function, continued

# Log-likelihood function, continued

```
double q2;
double logL:
const double sigma = 0.5;
q2 = 0.0;
for (int i = 0; i < num data; i++)</pre>
    q2 += sq error[i];
logL = -1 * q2 / (2 * sigma * sigma);
return logL:
```

# Further suggestions and conclusion

## SUGGESTION: CLOUD COMPUTING

If your access to parallel computing hardware is insufficient, consider cloud computing

- Google Cloud Platform
- Amazon Elastic Cloud Compute (EC2)
- · Microsoft Azure

Each platform has a free trial with limited access to HPC resources

Example: NVIDIA Tesla K80, in Europe, costs \$0.49 per hour per die

## SUGGESTION: CLOUD COMPUTING

#### Common features

- · Create VMs with a variety of resource configurations
- Connect using SSH to administer and run code
- · Create instances with GPUs
- Create clusters
- Only pay for what you use

If you want to write HPC code on a minimal system like a Chromebook, Amazon has a somewhat minimal cloud IDE, Cloud9

#### CONCLUSION

- · Model comparison comprises an important class of inference problems
- Existing techniques, e.g., nested sampling, can be parallelized in a variety of ways using a variety of hardware
- It is critically important to match algorithm to hardware
- Examples have been shown that address implementation on the Xeon Phi, CPUs, and GPUs
- Cloud computing platforms can provide an attractive alternative to dedicated hardware

#### RESOURCES

- Lawrence Livermore National Lab's "Introduction to Parallel Computing" https://computing.llnl.gov/tutorials/parallel\_comp/
- Best practice guide for older Xeon Phi models (Knight's Corner): http://www.prace-ri.eu/best-practice-guide-intel-xeon-phi-html/
- Best practice guide for newer Xeon Phi models (Knight's Landing): http://www. prace-ri.eu/best-practice-guide-knights-landing-january-2017/
- Introduction to OpenCL: http://www.drdobbs.com/parallel/ a-gentle-introduction-to-opencl/231002854