Assignment 5: Monte Carlo with Python, Cython, and C++

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ABSTRACT

The goal of this assignment is to explore the performance of Monte Carlo methods in Python, Cython, and C++. By randomly sampling points within a unit square and determining the fraction that lie within an inscribed unit circle $(x^2 + y^2 < 1)$, we can approximate the area of the circle. This ratio, when multiplied by 4, provides an estimate π . The principle behind this method is that as the number of samples increases, the estimate of π converges to the true value, with the error scaling as $1/\sqrt{n}$. This exercise involves the following implementions: (1) pure Python using a for-loop, (2) Python with NumPy, (3) Python with numba, (4) Cython, and (5) C++. You will benchmark the performance of each implementation, generate a table comparing execution times and per-sample throughput, and run a larger job on Ganymede2 in C++ for $\sim 10^8$ or more samples. You will compile your results in a short PDF report discussing performance, memory usage, and solution accuracy relative to π .

Learning Objectives: Monte Carlo methods (demonstration of an embarrassingly parallel problem), performance comparison between implementations (Python, Cython, and C++), HPC workflows (building, running, and timing), plotting of results, and analysis of performance scaling and memory usage.

1. PURE PYTHON IMPLEMENTATION

1. **Serial For-Loop**: Create a file pi_python.py:

```
from numpy.random import rand
import sys

def calc_pi_loop(n):
    h = 0  # Number of hits inside the circle
    for _ in range(n):
        x, y = rand(), rand()  # Random points in [0, 1)
        if x*x + y*y < 1.:
            h += 1  # Successful hit
    return 4. * float(h) / float(n)  # Estimate pi

if __name__ == "__main__":
    n = int(sys.argv[1])  # Command-line argument
    pi_est = calc_pi_loop(n)
    print(f"n={n}, pi={pi_est}")</pre>
```

2. Timing:

```
time python pi_python.py 10000
```

Start with a small n to check correctness, then increase n to measure the performance.

3. Table of Results: Record the total time and the time per sample in a table, which will be used for comparison with other implementations. For best results, run each method with as large an n as possible (can be different for each method).

2. NUMPY IMPLEMENTATION

1. Vectorized Approach:

```
from numpy import sum
from numpy.random import rand
import sys

def calc_pi_numpy(n):
    h = sum(rand(n)**2 + rand(n)**2 < 1.)
    return 4. * float(h) / float(n) # Estimate pi

if __name__ == "__main__":
    n = int(sys.argv[1])
    pi_est = calc_pi_numpy(n)
    print(f"n={n}, pi={pi_est}")</pre>
```

2. Compare Performance: Time the execution for a range of n values. It should be faster than the Python loop, but the memory usage will be higher. Can you tell at large enough n?

3. NUMBA-ACCELERATED PYTHON

1. Numba-based Monte Carlo:

```
import random, sys
from numba import jit, njit, prange
@njit
def calc_pi_numba(n):
   h = 0
    for _ in range(n):
        x = random.uniform(0, 1)
        y = random.uniform(0, 1)
        if x*x + y*y < 1.:
           h += 1
    return 4. * h / n
@jit(nopython=True, nogil=True, parallel=True)
def calc_pi_parallel(n):
   h = 0
   for _ in prange(n):
        x = random.uniform(0, 1)
        y = random.uniform(0, 1)
        if x**2 + y**2 < 1:
            h += 1
    return 4. * h / n
if __name__ == "__main__":
    n = int(sys.argv[1])
    pi_est = calc_pi_numba(n)
    \# pi_est = calc_pi_parallel(n)
    print(f"n={n}, pi={pi_est}")
```

2. Compare Performance: Note that @njit compiles the function at runtime so there is some overhead with this method. Time the execution for a range of n values. At what point does the overhead become

negligible? Compare the parallel version with the serial version for large enough n, reporting the speedup in additin to the time and per-sample cost.

4. CYTHON IMPLEMENTATION

2. Cython Code (calc_pi.pyx):

1. **Setup Directory**: Include the Cython code (calc_pi.pyx), a build script (setup.py), and a driver script (test_cython.py). In case you need to install Cython, use conda install cython (or pip if you prefer).

```
# cython: language_level=3
  cimport cython
  from libc.stdlib cimport rand, RAND_MAX
  @cython.boundscheck(False) # Disable bounds checking for performance
  @cython.wraparound(False) # Disable negative indexing for performance
  def calc_pi_cython(int n):
      cdef:
          int i, h = 0
          double x, y
      for i in range(n):
          x = rand() / RAND_MAX # Generate a random x coordinate
          y = rand() / RAND_MAX # Generate a random y coordinate
          # Check if the point (x, y) is inside the unit circle
          if x*x + y*y < 1.:
              h += 1
      # Estimate Pi using the ratio of points inside the circle to the total points
      return 4. * h / n
3. Setup Script (setup.py):
  from setuptools import setup
  from Cython. Build import cythonize
  setup(
      ext_modules=cythonize("calc_pi.pyx")
4. Driver Script (test.py):
  from calc_pi import calc_pi_cython
  from time import time
  logn = 8 # Number of samples
  n = 10**logn # Number of samples
  start = time() # Start timer
  pi = calc_pi_cython(n) # Calculate pi
  end = time() # End timer
  print(f"Estimated Pi: {pi} [Samples: 10^{logn}, Time: {end - start:.3f} s]")
```

5. Compile and Test:

```
python setup.py build_ext --inplace
python test_cython.py 1000000
```

6. **Performance Comparison**: Compare the Cython implementation with the previous methods. Note the speedup and memory usage. Does the Cython version incur any overhead for small n like numba? For large n, is it faster or slower than numba for this simple problem?

5. C++ IMPLEMENTATION

1. Code (main.cpp):

```
#include <iostream>
#include <random>
#include <cstdlib> // for atoi
int main(int argc, char* argv[]) {
   if (argc < 2) {
        std::cerr << "Usage: " << argv[0] << " N\n";
       return 1;
   }
   const int n = std::atoi(argv[1]);
   std::random_device rd; // Random seed from hardware
   std::mt19937 gen(rd()); // Mersenne twister engine
   std::uniform_real_distribution<> rand(0., 1.);
   int h = 0;
   for (int i = 0; i < n; ++i) {
        // Get random points
        const double x = rand(gen);
        const double y = rand(gen);
        // Check if point is inside the circle
       if (x*x + y*y \le 1.) h++;
   }
   double pi = 4. * double(h) / double(n);
   std::cout << "n=" << n << ", pi=" << pi << std::endl;
   return 0;
}
```

2. Compile and Run:

```
g++ -03 main.cpp -o mc_pi
time ./mc_pi 10000000
```

3. **Performance Comparison**: Compare the performance with the Python, Cython, and numba implementations. How does the C++ version stack up in terms of performance and memory usage? Can you run a job with $n = 10^8$ or more on Ganymede2? For very large n, you might spend several seconds to minutes for a single core run.

6. COLLECTING AND ANALYZING RESULTS

1. Benchmark Table: For the Python, NumPy, Numba, Cython (serial and parallel), and C++ implementations, determine the largest n that can be run in a reasonable time (~ 1 minute). Rank the speeds of the different methods in terms of the *samples per second*. Then for the fastest method only, run a range of $n \in \{10^3, 10^4, \dots, 10^8, 10^9\}$, record the following statistics:

Table: $n \mid \pi$ Estimate | Absolute Error | Standard Deviation | Runtime [s] | time/n | samples/sec Specifically, for each value of n, collect at least 10 estimates of π to compute the standard deviation. Save the results to a file to separate data collection from analysis. If you have time, use Ganymede2 to practice HPC job submission.

2. **Plots**:

- Mean Error: Plot the absolute error of the mean π estimate as a function of n.
- Standard Deviation: Plot the standard deviation of the π estimates as a function of n.
- Power-law Fit: Fit the standard deviation data to a power-law $\sigma \propto n^{-\alpha}$ to determine α .

3. Analysis:

- The value of α (and its uncertainty if you can). How does this compare to the expected $1/\sqrt{n}$ scaling?
- Based on α , estimate the number of samples (n) neded to achieve 12 digits of accuracy for π .
- Can you fit that many points in memory with the NumPy approach?
- Assuming perfect linear scaling, how long would it take to run that many samples on a single core?
- With perfect weak scaling, how many cores would you need to run that many samples in one year?

7. REPORTING GUIDELINES

1. Submission Format:

- Provide a short PDF (LaTeX or word processor) including:
 - (a) All relevant code modifications if needed.
 - (b) A benchmark table comparing each method.
 - (c) A plot of error and standard deviation vs. n.
 - (d) A power-law fit to the standard deviation data.
 - (e) Discussion of the analysis questions.
- 2. Evaluation Criteria: Correctness & Completeness, Clarity & Organization, and Discussion & Analysis.

This completes Assignment 5, practicing Monte Carlo methods in Python, Cython, and C++. Happy coding!