

- cuRAND uses GPU to generate pseudorandom numbers
- Host and Device API
- Host API like cuBLAS but for random numbers
- Device API can generate numbers while in kernel, more complicated

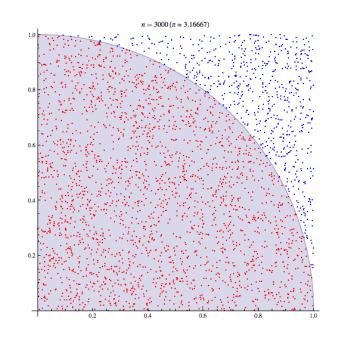
Cuda with random numbers is very useful for Monte Carlo

Repeated random sampling of the same system to get results

Each repetition is independent -> parallelizable!

Simple example: numerically approximate pi

randomly scatter points, in a region, and calculate the fraction of points inside the circle



Parametric definition of a circle makes it easy to tell which points are in the circle

$$x^2 + y^2 < 1$$
 : inside unit circle

Generate points uniformly distributed in {(0, 1], (0, 1]}

float* dev_points;

curandGenerator_t gen;

curandCreateGenerator(&gen,CURAND_RNG_PSEUDO_DEFAULT); curandGenerateUniform(gen, dev points, numPoints * 2);

Check bounds and reduce

__global___ void countPointsInCircle(float* points, int n) {

```
extern int sharedMem[];
int idx = threadIdx.x + blockIdx.x * blockDim.x;
float x = points[idx];
float y = points[idx + n];
sharedMem[threadIdx.x] = (x * x + y * y < 1);
//syncthreads and perform summing reduction
```

Just use our count to calculate the value we want

sum/numPoints is pi/4

Stochastic simulation of dynamic systems

Individual transitions are memoryless, if nothing happens for a long time, the system does not "remember" this

Simulation step:

Calculate the probability of each transition being the next event, calculate the distribution of the time the next event will occur at

Minimum of exponential random variables if X1, X2, X3... are exponential random variables

min(X1, X2, X3...) is an exponential random variable with $\lambda = \lambda 1 + \lambda 2 + \lambda 3...$

So we can calculate our transition distributions P(event i happens next) = λi /sum(all λ) Distribution of times

T : exponential with rate constant = sum(all λ)

We can sample these random variables using uniform distributions taken from cuRAND

A biological molecule is produced by a large reservoir of reactants, and decays at a rate proportional to concentration

reactants -> X -> 0

Nonstochastic model d[X]/dt = k - γ * [X]

Only accounts for average behavior What is the variance of [X] at equilibrium?

Reaction propensitiesX++kX--γ * X

