Drawing 10e12 samples from a judgmental estimation model

Nuño Sempere, Jorge Sierra

December 13, 2024

Abstract

To draw $10\mathrm{e}12$ samples you need two things: a fast language, and a supercomputer.

Background and motivation

Estimators in the tradition of Fermi or Tetlock sometimes create models that seek to capture their subjective credences—as opposed to, for instance, fitting a model to an underlying set of samples. One might go about this is by expressing a distribution using the set of idioms that grew from the foretold io experimental forecasting platform into Squiggle—now a programming language for intuitive probabilitistic estimation. Some of the idioms in squiggle have been ported to Python (SquigglePy), to C (Squiggle.c), or to go.

However, those idioms favour relying on Monte Carlo estimation, that is, approximating a model by drawing samples rather than specifying the model exactly. This approach sometimes has difficulty modelling long-tail behaviour.

For instance, in our squiggle.c test suite, we draw 10M samples from various distributions, and then compare the mean and standard deviation of the samples to the mean and standard deviation of the underlying distributions. With 10M samples, this sometimes shows relative errors of 50% or more. For insance:

```
[-] Mean test for lognormal(0.644931, 4.795860) NOT passed.
Mean of lognormal(0.644931, 4.795860): 125053.904456, vs expected mean: 188163.894101 delta: -63109.989645, relative delta: -0.504662
[-] Std test for lognormal(0.644931, 4.795860) NOT passed.
Std of lognormal(0.644931, 4.795860): 39976300.711166, vs expected std: 18577298706.170452 delta: -18537322405.459286, relative delta: -463.707799
```

To explore those limits, and also just for the sake of the technical challenge, we describe how to draw 10e12 samples (an American trillion) from a judgmental model. In this case, the model is a very rough back-of-the-envelope estimate of

the altruistic impact of Sentinel—a foresight and emergency response team set up by one of the authors—in basis points of existential/catastrophic risk¹ mitigated or averted per million dollars. We show how summary statistics change as we draw more samples. We conclude discussing limitations and practical applications.

1. How to draw 10e9 samples

1.1. Use a fast language.

Our time to botec repository compares how fast drawing 10e6 samples from a simple back-of-the-envelope calculation runs in different languages. The comparison depends on our familiarity with the different languages: as we are more familiar with a language, we grow capable of writing better & faster code in it. Nonetheless, speeds to get 10e6 samples in one author's computer are as follows:

Language	Time
C	$6.20 \mathrm{ms}$
squiggle.c	$7.20 \mathrm{ms}$
go	$21.20 \mathrm{ms}$
Nim	$41.10 \mathrm{ms}$
Lua (LuaJIT)	$68.80 \mathrm{ms}$
Python (numpy)	$118 \mathrm{ms}$
OCaml (flambda)	$185 \mathrm{ms}$
Squiggle (bun)	$384 \mathrm{ms}$
Javascript (node)	$395 \mathrm{ms}$
SquigglePy (v0.27)	$1,542 \mathrm{ms}$
R (3.6.1)	$4{,}494\mathrm{ms}$
Python 3.9	11,909 ms
Gavin Howard's bc	16,170 ms

Readers might have expected this list to include languages such as Rust or zig. FORTRAN, Lisp, Haskell, Java, C#, C++, or Julia are also missing. So are faster options, like GPU or even assembly programming. This is because we are not very familiar with those languages, or because in our attempts to use them we found them too "clunky".

Still, the above table shows that the time it takes to run a simple Fermi model varies 3 to 4 orders of magnitude depending on the language choice (and skill at using it), and so language choice will be an important choice.

1.2. Use standard optimizations.

Once we have chosen a fast language like C, we can deploy some standard practices, like inlining functions, passing pointers not values, or packing structs

 $^{^{1}}$ with a conversion factor of existential risk being 100x worse than catastrophic risk.

into a single cache line. The difference between an rough first pass and a final version with more care can be notable.

Using compiler flags can also give a noticeable speed bump, as can profiling. However, we didn't take advantage of this fully for drawing 10e12 samples because the compiler in our local machines (gcc/clang) was different than the compiler in the supercomputer we used for our final run (icc patched for usage with MPI).

1.3. Directly implementing fast randomness primitives helps with speed and clarity.

Back in our javascript days, we were frustrated at stlib.io for having needless complexity because of the need to support incompatible types of javascript modules, and for being slow. When writting squiggle.c, we initially looked for a library to rely on, and chose the GNU Scientific Library for supporting a good range of distributions.

However, ultimately, we ended up writting our own randomness primitives, mostly following the work of Marsaglia. Languages like Rust or zig have some well-optimized randomness primitives that have sometimes been a joy to read. But C doesn't, and C has other virtues, like being ubicuitous and having powerful parallelism and distributed computing libraries.

We chose the xorshift pseudorandomness generation, on account of its speed, and keeping in mind that we didn't at all need our PRNG to be cryptographically secure.

```
static uint64_t xorshift64(uint64_t* seed)
{
    uint64_t x = *seed;
    x ^= x << 13;
    x ^= x >> 7;
    x ^= x << 17;
    return *seed = x;
}</pre>
```

The seed is fed as a pointer, rather than it being a global variable, to allow for parallelism.

In some sense, drawing 10e12 samples from such a simple generator feels like stretching it too far. That said, we aren't aware f anything we are doing that's specifically wrong with it, and the algorithm's period is $2^{64} - 1$. Regardless, if you can draw 1T samples at all, you can draw 100K, 1M, 10M, 100M samples very fast.

Readers are encouraged to browse the squiggle.c README.md, FOLK_WISDOM.md and library code for a longer treatment on the virtue and pitfalls of our fast Monte Carlo primitives.

1.4. Use single-core and multi-code parallelism

We used OpenMP (for an 8x to 16x speedup if using that many thread on our local machines, but 64x if using the cores in the Finisterrae supercomputer); and MPI, for a speedup that ended up being roughly 4x (4 nodes, minus a bit of overhead) but which could be much scaled. For fast multi-core orchestration, our sense is that there aren't that many alternatives to MPI.

We could get another 10x speedup by spending 10x as much time (60 hours instead of 6 hours) or 10x more cores. However, that approach quickly doesn't scale. One could also get a final speedup by using a more recent generation computer, but we didn't pursue that option, or it wasn't available to us.

When doing parallelism, one interesting trick is to structure arrays such that each thread is accessing different parts of an array simultaneously, but without cache conflicts and misses. One can do this by starting each thread far enough, but one can also pad the elements of an array such that cache conflicts can't happen, as follows:

```
#define CACHE_LINE_SIZE 64
typedef struct seed_cache_box_t {
    uint64_t seed;
    char padding[CACHE_LINE_SIZE - sizeof(uint64_t)];
    // Cache line size is 64 *bytes*, uint64_t is 64 *bits* (8 bytes). Different units!
} seed_cache_box;
// ...
```

seed_cache_box* cache_box = (seed_cache_box*)malloc(sizeof(seed_cache_box) * (size_t)n_threa

In our project, we used C macros sparingly, but we did use them to allow MPI and non-MPI code to exist side by side, such that we could run the OpenMP

side on our local machine, with an MPI "world" of only one machine. Readers can find our OpenMP and MPI code in the samples.c file in this repos-

itory, as well as recipes to actually deploy it in our makefile. TODO: add link.

1.5. Design for speed then compromise, rather than the reverse

Throughout this project, we didn't always make the speed-maximalist decision all the time. Some notable places where we didn't are as follows:

squiggle.c has a sample_mixture function which takes an array of function pointers:

```
double weights[] = { 0.3, 0.4, 0.3 };
double (*samplers[])(uint64_t*) = { sample_dist_1, sample_dist_2, sample_dist_3 };
double x = sample_mixture(samplers, weights, n_dists, seed);
```

This adds a level of indirection, and would be slower than directly writing:

```
double p = sample_uniform(0,1, seed);
double x;
if(p < p1){
  return sample_dist_1(seed);
} else if (p < p1 + p2) {
  return sample_dist_2(seed);
} else {
  return sample_dist_3(seed);
}</pre>
```

We further didn't use algebraic manipulations. For instance, multiplying two lognormal distributions, or two beta distributions together, produces a result in the same family. However, when expressing a judgmental model, it isn't parsimonious to group betas and lognormals together to allow for algebraic manipulations: doing so would make the judgmental model much less readable, and the underlying code messier and less general. We took a $\sim 2\text{-}3x$ slowdown hit because of this.

Ultimately, there s a sweet spot in the (high user accessibility, high complexity, low speed) end of the spectrum: tools like causal, guesstimate squiggle. There is also a sweet spot in the (low user accessibility, low complexity, high speed): the project we are exploring. With some of the lessons from writting squiggle.c, one of the authors created fermi, a command line read-evaluate-print loop (REPL) in go in which it is much faster to write small models, at the cost of being slower (while still being able to draw 100K samples instantaneuosly).

The punchline is that by paying attention to speed from the beginning and then dialing it down for some expressivity, usability gains, we end up with a much faster result, one that is able to better explore the long tail of outcomes due to being able to draw enough samples to meaningfully draw from it.

2. Results as samples increase

2.1. Mean and standard deviation

2.2. Minimum and maximum

Conclusion

Rethink Priorities' models puny struggled (https://forum.effectivealtruism.org/posts/pniDWyjc9vY5sjGre/retheriorities-cross-cause-cost-effectiveness-model) to reach more than 150k samples. Eventually, with some optimizations, they moved to the billions. Much more is possible with a focus on speed.

Thin line between Pascal's wager and vNM. Kosonen?

Other titles: The virtue of fast Monte Carlo primitives: Drawing 1T samples and a REPL for rapid fermi iteration