Lightweight Implementations of Probabilistic Programming Languages Via Transformational Compilation

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Abstract

We describe a general method of transforming arbitrary programming languages into probabilistic programming languages with straightforward MCMC inference engines. Random choices in the program are "named" with information about their position in an execution trace; these names are used in conjunction with a database of randomness to implement MCMC inference in the space of execution traces. We encode naming information using lightweight source-to-source compilers. Our method enables us to reuse existing infrastructure (compilers, interpreters, etc.) with minimal additional code, implying fast models with low development overhead. We illustrate the technique on two languages, one functional and one imperative: Bher, a compiled version of the Church language which eliminates interpretive overhead of the original MIT-Church implementation, and Stochastic Matlab, a new open-source language.

1 INTRODUCTION

Probabilistic programming languages simplify the development of probabilistic models by allowing programmers to specify a stochastic process using syntax that resembles modern programming languages. These languages allow programmers to freely mix deterministic and stochastic elements, resulting in tremendous modeling flexibility. The resulting programs define prior distributions: running the (unconditional) program forward many times results in a distribution over execution traces, with each trace generating a sample of data from the prior. The goal of inference in such programs

is to reason about the posterior distribution over execution traces conditioned on a particular program output. Examples include BLOG [6], PRISM [13], Bayesian Logic Programs [4] Stochastic Logic Programs [7], Markov Logic [12], Independent Choice Logic [10], IBAL [9], Probabilistic Scheme [11], Λ_{\circ} [8], and HANSEI [5].

We present a general technique for turning any programming language into a probabilistic programming language with an accompanying universal Markov chain Monte Carlo inference engine. Our method allows the full use of all language constructs permitted by the original (non-probabilistic) language by leveraging native compilers (or interpreters). The key technical idea is to give each random choice of a fixed program a unique "name" that depends on its position in a given execution trace. We then convert stochastic functions into deterministic ones which use these names as indices into a database of random choices. By controlling the values in the database of randomness, execution traces of the program can be controlled, which allow us to construct the key operations needed for the Metropolis-Hastings algorithm (i.e., generating and scoring proposals).

The most obvious techniques for tracking random choices (such as the position in a random bit stream) result in states that are unsuitable for MCMC (a function that consumed the 10th bit in one execution may consume the 30th in another—but perhaps should be considered the same). We instead name random choices based on their structural position in the execution trace. We describe how naming information can be provided by a side computation that is constructed by a sourceto-source transformation at compile time: the original probabilistic program can be augmented with naming information to create a new program that makes appropriate random choice names available. The resulting new program can be executed at full speed, with only minimal overhead to track names and control the database of randomness, and no changes to the compiler or interpreter for the underlying language.

The resulting MCMC algorithm constitutes a new implementation of the Church MCMC algorithm [3], which is very similar to the algorithm used in BLOG [6]. However,

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we provide a much simpler and uniform approach to constructing such inference engines for new languages, with small, clean code bases.

The implementation technique for probabilistic programs closest to ours is that of Kiselyov and Shan [5], who share the goal of transforming standard languages into probabilistic versions with little or no interpretive overhead. They use delimited continuations to control random choices without explicit choice naming. However, their technique is only applicable (prima facie) to inference by enumeration or importance sampling, not MCMC.

We illustrate our technique on two languages, one functional and one imperative. Our first implementation is named "Bher," and is a compiler for the Church probabilistic programming language [3]. The current MIT-Church implementation uses a custom interpreter to reason about the distribution of execution traces, but since our method does not incur interpretive overhead, the resulting programs can run orders of magnitude faster. We also illustrate the technique on an imperative language by creating Stochastic Matlab, a new probabilistic programming language.

2 OVERVIEW OF THE METHOD

We begin by outlining our setup. We define an unconditioned probabilistic program to be a parameterless function f with an arbitrary mix of stochastic and deterministic elements (hereafter, we will use the term function and program interchangeably). The function f may be written in any language, but our running example will be Matlab. We allow the function to be arbitrarily complex inside, using any additional functions, recursion, language constructs or external libraries it wishes. The only constraint is that the function must be self-contained, with no external side-effects which would impact the execution of the function from one run to another (in other words, no information or program state can be communicated from one run to the next).

The stochastic elements of f must come from a set of known, fixed *elementary random primitives*, or ERPs. Complex distributions are constructed compositionally, using ERPs as building blocks. In Matlab, ERPs may be functions such as rand (sample uniformly from [0,1]) or randn (sample from a standard normal). Higher-order random primitives, such as nonparametric distributions, may also be defined, but must be fixed ahead of time.

Formally, let \mathcal{T} be the set of ERP types. We assume that each type $t \in \mathcal{T}$ is a parametric family of distributions $p_t(x|\theta_t)$, where θ_t are the parameters of the distribution. As f is executed, it encounters a series of ERPs. We denote by f_k the k'th ERP, and x_k the value returned by f_k .

Algorithm 1 A simple Stochastic Matlab program representing a Gaussian-Gamma mixture model.

```
1: for i=1:1000

2: if (rand > 0.5)

3: X(i) = randn;

4: else

5: X(i) = gammarnd;

6: end;

7: end;
```

Note that the k'th ERP may change types (indeed, the total number K of ERPs may change) depending on previous x_k 's. By f_k , it should therefore be understood that we mean $f_{k|x_1,\cdots,x_{k-1}}$, but will suppress the conditional to keep notation clear.

Algorithm 1 shows an example of a simple f written in Matlab. Here, there are three syntactic ERPs: rand, randn, and gamma. During execution, depending on the return value of each call to rand, different ERPs are encountered. A total of 2000 random choices will be made when executing this program.

We denote by x all of the random choices which are made internally by f, meaning that f defines the probability distribution p(x). In our example, $x \in \mathbb{R}^{2000}$. The probability of any particular x is therefore the product of the probability of all of the ERP choices made:

$$p(x) = \prod_{k=1}^{K} p_{t_k}(x_k | \theta_{t_k})$$

again remembering that types and parameters may depend arbitrarily on previous random choices.

Generative functions as described above are, of course, easy to write. A much harder problem, and our goal in this paper, is to reason about the posterior conditional distribution $p(x_{\setminus c}|x_c)$, where we define x_c to be a subset of random choices which we condition on and $x_{\setminus c}$ to be the remaining random choices. For example, we may condition our example program on the values of the X(i)'s, and reason about the sequence of rand's most likely to generate the X(i)'s. Rejection sampling is one possible inference method: we may run f forward until x_c is generated, somehow recording $x_{\setminus c}$. Of course, such an approach is intractable for all but the simplest models.

We propose a different approach, based on the following intuition. Consider repeatedly running an unconditioned f. Every time an f_k is encountered, the program samples an x_k and continues execution. Running f multiple times will result in different execution paths through the code defining f. We term such an execution path a *trace*. Importantly, if two runs of f make all the same random choices (i.e., each f_k returns the same x_k), their execution trace will be the same. This suggests the following

Algorithm 2 An MCMC trace sampler.

```
1: Initialize: [ ll, \mathbb{D} ] = trace_update(\emptyset)
 2: Repeat forever:
         Select a random f_k via its name n
 3:
         Look up its current value (t, x, l, \theta_{db}) = \mathbb{D}(n).
 4:
         Propose a new value x' \sim \mathcal{K}_t(\cdot | x, \theta_{dh})
 5:
 6:
         Compute F = \log \mathcal{K}_t(x'|x, \theta_{db})
          Compute R = \log \mathcal{K}_t(x|x', \theta_{dh})
 7:
 8:
         Compute l' = \log p_t(x'|\theta_{dh})
         Let \mathbb{D}' = \mathbb{D}
 9:
         Set \mathbb{D}'(n) = (t, x', l', \theta_{db})
10:
          [ll', \mathbb{D}'] = \text{trace\_update}(\mathbb{D}');
11:
          if (\log(\text{rand}) < ll' - ll + R - F)
12:
13:
              // accept
              \mathbb{D} = \mathbb{D}'
14:
              ll = ll'
15:
16:
              // clean out unused values from D
17:
18:
              // reject; discard \mathbb{D}'
19:
          endif;
20: end repeat;
```

procedure for controlling the execution trace of f:

- 1. Name each f_k in every possible execution trace.
- 2. Rewrite the source code f to generate f', replacing random functions f_k with deterministic functions f_k' . When encountered in the execution trace, these functions f_k' deterministically use their name to look up a current value x_k in a database of randomness and return it (and if no value exists, they sample $x_k \sim p_{t_k}(\centerdot|\theta_{t_k})$., store it in the database, and then return it). Behind the scenes, these functions also accumulate the likelihood of each random choice.

Thus, the execution trace of f' can be controlled by manipulating the values in the database of randomness.

This makes MCMC inference possible by enabling proposals, scoring, and the ability to accept or reject proposals. We make a proposal to an execution trace by picking a random choice in the database, modifying it according to a proposal distribution, and then re-executing f' and computing the likelihood of the new trace. As f' is executed, some random choices may be reused, new randomness may be triggered (which is sampled from the prior), and randomness that was previously used may no longer be needed. In our example, we can make a proposal to, say, the 500th rand. This will change the execution path of the program, switching between the Gaussian branch and the gamma branch.

The foregoing outline leaves many questions unanswered, which we now address in detail. Section 2.1

Algorithm 3 Function trace_update(□)

```
1: Set ll = 0
 2: Execute f :
 3: For all random choices k:
        Run computation until choice k,
        determining n, t_c, \theta_c for k.
 5:
        Look up (t, x, l, \theta_{db}) = \mathbb{D}(n)
        if a value is found in the database and t = t_c
 6:
 7:
            if parameters match (ie, \theta_c == \theta_{dh})
                ll = ll + l
 8:
 9:
            else
10:
                // rescore ERP with new parameters
                compute l = \log p_{t_c}(x|\theta_c)
11:
12:
                store \mathbb{D}(n) = (t_c, x, l, \theta_c)
                ll = ll + l
13:
            endif
14:
        else
15:
            // sample new randomness
16:
17:
            sample x \sim p_t(.|\theta_c)
18:
            compute l = \log p_t(x|\theta_c)
            store \mathbb{D}(n) = (t_c, x, l, \theta_c)
19:
            ll = ll + l
20:
21:
        endif
        Set return value of f_k to x
22:
23: end for all
24: return [ll, \mathbb{D}]
```

discusses the overall MCMC algorithm, Section 2.2 discusses how random variables are named, and Section 2.3 discusses conditional inference.

2.1 MCMC IN TRACE SPACE

We now describe our MCMC algorithm. In this section, We will assume that some naming scheme for random choices has been defined, and that a mechanism for computing names of random choices has been implemented, but we will return to this important topic in the next section.

Let a database of randomness $\mathbb D$ be defined as a mapping $\mathbb N \to \mathcal T \times X \times \mathbb L \times \theta$, where $\mathbb N$ is the name of a random choice, $\mathcal T$ is its ERP type, X is its value, θ are ERP parameters, and $\mathbb L$ is this random value's likelihood. We allow missing entries in the database.

We can control the execution of f' by controlling the values in \mathbb{D} . When f' encounters an f'_k , it computes its name $n \in \mathbb{N}$, its parameters θ_c (where 'c' is a mnemonic for "current"), and its current type $t_c \in \mathcal{T}$. It then looks up $(t, x, l, \theta_{db}) = \mathbb{D}(n)$. If a value is found and the types match, we set the return value of f_k to be x. Otherwise, the value is sampled from the appropriate ERP $x \sim p_{t_c}(.|\theta_c)$, its likelihood is computed, and the corresponding entry in the database is updated. This is formalized

Algorithm 4 An illustration how the number and type of random choices can change.

- 1: m = poissrnd();
- 2: for i=1:m
- X(i) = gamma;
- 4: end;
- 5: for i=m+1:2*m
- 6: X(i) = randn;
- 7: end;

in the trace_update procedure, which we define in Algorithm 3.

We use trace_update to define our overall MCMC algorithm, shown in Algorithm 2. Given a current trace x and score p(x), we proceed by reconsidering one random choice x_k . We equip each ERP type with a proposal kernel $\mathcal{K}_t(x'|x,\theta)$, which we use to generate proposals to x_k . After a proposal, we call trace_update to generate a new trace x', and compute its likelihood p(x'), which is the product of any reused random choices, and any new randomness that was sampled. This gives us an overall score, which is used as an MH accept ratio

$$\alpha = \min \left\{ 1, \frac{p(x') \mathcal{K}_t(x|x', \theta)}{p(x) \mathcal{K}_t(x'|x, \theta)} \right\}$$

There are a few subtleties to the implementation. Even if an x for an f_k' is found in \mathbb{D} , we must compare the parameters θ_c and θ_{db} to make sure they are equal; if they are not, we may still be able to re-use x, but must re-score it under the new parameters θ_c . If a proposal is accepted, we also remove stale values from the database. This ensures that new randomness is sampled from the prior, and allows us to avoid reversible jump corrections.

2.2 NAMING RANDOM VARIABLES

What properties do we desire in a naming scheme? We build around a simple idea: when we propose a new x_k' and update the trace, we wish to reuse as many x_k 's as possible. This will give us a new execution trace "close" to the original, which is important to ensure a high acceptance rate in our MCMC sampler (recall that if an f_k' cannot reuse a value from $\mathbb D$, one is sampled from the prior, which is unlikely to be good).

How might different naming schemes affect reuse of randomness? Algorithm 4 shows a simple program f which we will use to illustrate different approaches. As f executes, it first encounters a Poisson, and then a sequence of gamma and Gaussian variables. One might be tempted to name random variables sequentially, according to the order in which they are encountered during execution—that is, use k itself as the name. Figure 1 (top) illustrates why this is a bad idea. Two traces are shown schemat-

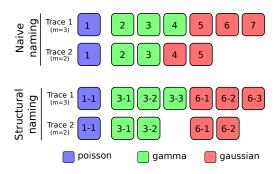


Figure 1: An illustration of two naming schemes. Variables are boxes with names inside. Top: the problem with naive variable naming. Notice the type mismatch at time 4 as well as the general sequence misalignment. Bottom: structural naming yields the desired behavior, reusing the maximum number of random choices.

ically, illustrating the f_k' 's encountered. Suppose that in trace #1, the value of m is 3. A proposal is made, changing m to 2, yielding trace #2. If variables are named sequentially, then x_1 and x_2 can be reused without problem, but when f_3' looks itself up in $\mathbb D$, it discovers that its type has changed, and must sample x_3 from the prior. The rest of the sequence is now "misaligned," and must probably be discarded. More sophisticated versions of this problem complicate this naming scheme to the point of unworkability, especially with complex models.

What is an alternative? Examining Alg. 4, we see that would like to name variables according to the way they are used at a more semantic level. For example, we might want to name a variable "the third gamma variable in the loop on lines 2-4."

Our key technical contribution formalizes this intuition. We name random choices according to their *structural position* in the execution trace, which we define in a way roughly analogous to a stack address: a variable's name is defined as list of the functions, their line numbers, and their loop indices that preceed it in the call stack. Thus, in Alg. 4, variables are named "1" (the initial Poisson), "3-1,", "3-2," ... (the gamma variables sampled on line 3), and "6-1,", "6-2," ... (the Gaussian variables sampled on line 6). Figure 1(bottom) illustrates this on Alg. 4: we see that the maximum number of random variables are reused.

We now discuss naming schemes for imperative and functional languages.

Imperative. For an imperative language, such as Matlab, we define the structural position with an abstract stack trace that is augmented with a combination of function identifiers, line numbers, and loop iteration numbers. The formal specification is inductive, as shown in Figure 2. In an interpreted imperative language, such as Python, Matlab or Ruby, this naming infor-

- Begin executing f with empty function, line, and loop stacks.
- · When entering a new function,
 - push a unique function id on the function stack.
 - o push a 0 on the line stack.
- When moving to a new line, increment the last value on the line stack
- When starting a loop, push a 0 on the loop stack.
- When iterating through a loop, increment the last value on the loop stack.
- When exiting a loop, pop the line stack
- When exiting a function, pop the function stack

Figure 2: The imperative naming specification. The name of an f_k is the state of the three stacks when f_k is encountered.

mation could potentially be acquired via dynamic stack examination, assuming suitable libraries exist. Alternatively, this information can be provided by augmenting the source code for f, as we shortly discuss.

Functional. For a functional language, such as Lisp, we do not need to worry about loops or line numbers: everything is a function, so we may define the structural position with a stack trace that is augmented with function identifiers. The specification is shown in Figure 3.

There is no guarantee that this naming scheme is optimal in any sense, but it does capture our desiderata, is simple and fast, and incurs minimal overhead.

2.3 CONDITIONAL INFERENCE

In principle, a program can be conditioned on any expression that evaluates to a Boolean value, not just on a fixed value assignment for a subset of the random variables of the program. The meaning of such a conditioned program is defined by rejection sampling: Sample an execution trace from the unconditioned program. If the value of the conditioning expression is true, return this trace. Otherwise, retry.

In practice, a MCMC implementation that allows general conditions requires that for any valid condition, we can find an initial assignment of random variables to values that satisfies this condition. If conditioning is not restricted to random variables (Stochastic Matlab), then this is a search problem that needs to be solved. Possible approaches include constraint propagation (MIT-Church), annealing (Bher), and rejection sampling (Bher). We leave the analysis of this problem to future research. In our experimental results, we exclude the time it takes to find an initial assignment from the infer-

Figure 4: Laziness transform \mathcal{L} and definition of the function force referenced in the transform.

ence time comparison in order to evaluate MCMC performance more directly.

3 IMPLEMENTATION AND EXPERIMENTS

We now present two implementations of our technique. Bher is a compiled implementation of Church, and Stochastic Matlab is a new, open-source language. Both illustrate the virtues of our approach: small code bases, high speed, and language flexibility.

3.1 EXAMPLE: BHER

Bher is an implementation of the Church probabilistic programming language that compiles Church code to native Scheme code. Like MIT-Church, the original implementation of the Church language, Bher is written in R6RS Scheme. However, MIT-Church is an *interpreter*—not based on the lightweight implementation principle—and thus provides a useful point of comparison.

In the following, we first discuss the addressing transform that names random choices in Bher, then compare performance between Bher and MIT-Church on Hidden Markov Models and Latent Dirichlet Allocation.

3.1.1 Syntactic Transformations

The syntactic transform that provides addresses is shown in Figure 3. The address that is built at runtime consists of a list of symbols. The transform is designed to change the program code such that a new symbol is added to the front of this list whenever the stack is extended.

The transform has two central parts: First, change each function definition such that the function takes another argument, the address. Second, change each function

```
 \begin{split} &\mathcal{A}^{top} \llbracket E \rrbracket = ((\text{lambda } (\text{addr}) \ \mathcal{A} \llbracket E \rrbracket) \ '(\text{top})) \\ &\mathcal{A} \llbracket ((\text{lambda } (I_{i=1}^n) \ E_{body}) \rrbracket = (\text{lambda } (\text{addr } . I_{i=1}^n) \ \mathcal{A} \llbracket E_{body} \rrbracket) \\ &\quad \text{where } \mathcal{S} \text{ is a globally unique symbol.} \\ &\mathcal{A} \llbracket (\text{mem } E) \rrbracket = ((\text{lambda } (\text{maddr } f) \ (\text{lambda } (\text{addr } . \text{args}) \ (\text{apply } f \ (\text{cons args maddr}) \ \text{args}))) \ \text{addr } \mathcal{A} \llbracket E \rrbracket) \\ &\mathcal{A} \llbracket (\text{begin } E_{i=1}^n) \rrbracket = (\text{begin } \mathcal{A} \llbracket E_i \rrbracket_{i=1}^n) \\ &\mathcal{A} \llbracket (\text{letrec } ((I_i \ E_i)_{i=1}^n) \ E_{body}) \rrbracket = (\text{letrec } ((I_i \ \mathcal{A} \llbracket E_i \rrbracket)_{i=1}^n) \ \mathcal{A} \llbracket E_{body} \rrbracket) \\ &\mathcal{A} \llbracket (\text{if } E_t \ E_c \ E_a) \rrbracket = (\text{if } \mathcal{A} \llbracket E_t \rrbracket \ \mathcal{A} \llbracket E_a \rrbracket) \\ &\mathcal{A} \llbracket (\text{define } I \ E) \rrbracket = (\text{define } I \ \mathcal{A} \llbracket E \rrbracket) \\ &\mathcal{A} \llbracket (\text{quote } E) \rrbracket = (\text{quote } E) \\ &\mathcal{A} \llbracket (E_{op} \ E_{i=1}^n) \rrbracket = (\mathcal{A} \llbracket E_{op} \rrbracket \ (\text{cons } '\mathcal{S} \ \text{addr}) \ \mathcal{A} \llbracket E_i \rrbracket_{i=1}^n) \\ &\mathcal{A} \llbracket E \rrbracket = E, \text{ otherwise.} \end{split}
```

Figure 3: The naming specification for Bher is given by a syntactic transform \mathcal{A}^{top} .

application such that (1) the current address is extended with a symbol that uniquely identifies this (syntactic) application within the program and (2) pass on this modified address as an argument when the function is applied.

On the top level of the program, the address variable addr is initialized to '(top). Except for memoization (mem) and quote, all remaining syntax cases are handled by applying the addressing transform to all proper subexpressions. Primitive functions like + and cons need to be redefined to take (and ignore) the address argument, e.g. to (lambda (addr . args) (apply prim args)). In Bher, all globally free variables are treated as primitives and redefined in this way.

In deterministic programs, memoization is an optimization technique that does not affect semantics. In stochastic programs, we can distinguish mem as a semantic construct from mem as an optimization technique [3]. The distribution defined by (eq? (flip) (flip)) is very different from the distribution defined by (let ([mf (mem flip)]) (eq? (mf) (mf))).

When addresses are available, semantic memoization is a simple syntactic transform. In a program that has been transformed to provide addresses, random variables get their values by looking up the value that is stored in the global database at their address. Thus, to enforce that two random choices always have the same value, we make sure that the address they receive is the same. To achieve this, the mem case of the addressing transform builds a function that captures within its closure the address of the mem creation and uses this address (extended by any function arguments) instead of the address provided to the function when it is applied.

The Church program shown in Algorithm 5 and its transformed version (Algorithm 6) illustrate the address computation that is induced by the syntactic transform. The program defines a simple geometric distribution with a potentially unbounded number of random choices, each of which gets a unique address. When we propose to change the value of the flip that controls the recur-

Algorithm 5 A Church program that samples from a geometric distribution.

Algorithm 6 Transformed version of the Bher program shown in Algorithm 5.

sion depth, we want to reuse as much of the existing trace as possible. An example of such a trace is shown in Figure 5. Note that apply nodes (shown in green) are the only place where addresses change and that nodes that refer to the same *syntactic* place (e.g. the two blue (flip p) nodes) have different addresses when they occur in different places in the trace. A change to the return value of such a flip does not affect the addresses of the choices "above" it. Thus, any randomness stored for these choices persists across the change.

A major virtue of source-to-source transformations is simplicity and compositionality. To illustrate this point, we now sketch a complementary transformation, eager style to lazy style, that can be used to avoid unnecessary computation. As shown in Figure 4, we delay expressions that must not necessarily get evaluated—for example, ar-

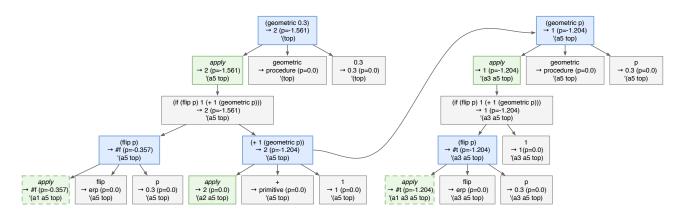


Figure 5: Example of an execution trace for the geometric sampler defined by Algorithm 5. Each node is annotated with its expression, value, log probability, and address. Applications (blue nodes) have subtraces for operator, operands, and for the application of operator to operands (*apply*, green nodes; dashed border indicates a random choice).

guments to compound functions—and force expressions that are always evaluated—operators, for example. To delay an expression E, we wrap it in (mem (lambda () E)). Thus, we ensure that it always returns the same result when evaluated multiple times. To force a value, we check whether it is delayed and if it is, we simply apply the memoized function stored in the value and recurse.

This difference in simplicity manifests itself in the code base of the two projects. While the MIT-Church implementation has about 11,119 lines of Scheme code, Bher consists of only 1,658 lines of code.

3.1.2 Experiments

We have tested inference performance in MIT-Church and Bher on Hidden Markov Models and Latent Dirichlet Allocation (Figure 6). In both cases, we conditioned on an observation sampled from the model. For each HMM, we generated a sentence of length 15. For the LDA models—each with 10 topics, 10 words per document, and 30 possible words—we varied the number of documents (bags of words) that we conditioned on. We have run all experiments using the optimizing compiler Ikarus [2].

On the HMM model, Bher outperforms the MIT-Church interpreter. On the LDA model, Bher shows better performance at smaller problem sizes whereas MIT-Church scales better to larger instances. This difference in scaling is due (1) reuse of cached computation in mem and (2) computational short-circuiting in MIT-Church. In the process of updating a trace after a proposal has been made, MIT-Church can detect that the information flowing into a function application has not changed; in this case, it does not recompute the return value. Since mem is transformed away by the compiler, we can ask whether there is a general transformation that results in reuse of computation with effects similar to those achieved in MIT-Church. This is an area of future research.

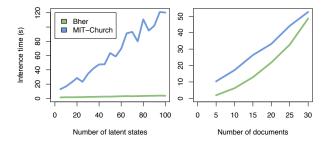


Figure 6: Inference performance for 10,000 samples from Hidden Markov Models and 1,000 samples from Latent Dirichlet Allocation models.

3.2 EXAMPLE: STOCHASTIC MATLAB

We now turn to an imperative language: Stochastic Matlab. Because Matlab is such a popular programming language among CS researchers, we felt that it would make a good testbed for a new language using our technique. Because we do not have deep language support (Matlab is a closed language), we use the transformational compilation technique to provide ERP names.

3.2.1 Syntactic Transformations

To rewrite f, we used the imperative ERP naming specification as outlined in Fig. 2. We implemented the rewriter using a parser developed with code from the open-source Octave project [1]. ERP names were rewritten to deterministic functions using a MEX-based database. Remaining code was implemented in pure Matlab. Figure 7 (top left) shows a small model and its rewritten equivalent.

3.2.2 Case Study

We now consider a case study showing why our technique is useful. We will briefly go through a complex

A layer model of rock porosity

```
function dp_render_rock( data )
  num_layers = poissrnd();
  dp_sample_layer = dpmem( @sample_layer, 1.0 );
  for i=1:num_layers
    layer(i,:) = dp_sample_layer();
  end;
  rock_volume = render_layers( layers );
  data = rock_volume + randn( size(rock_volume) );
  return;
```

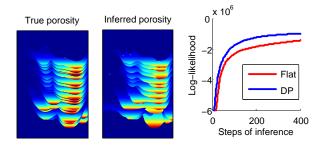


Figure 7: Results on modeling rock porosity.

Matlab model which relies on a deterministic GPU-based MEX function. The point of this example is twofold: first, to illustrate how easy it is to mix arbitrary deterministic elements with stochastic ones—reusing complex external libraries—and second, to illustrate how easy it can be to create complex model with a few simple commands.

Our example is a small (but real) application of inverse geological modeling. Layers of rock and sediment are deposited over millions of years to create a volumetric cube of rock with porosity that varies according to the geometries of the layers. Given this forward model, we wish to condition on a given volume, and recover a description of the individual layers that built up the volume.

Figure 7 shows the model, which generates a rock volume by first sampling parametric descriptions for each layer (the details are not shown; layers are described as a weighted set of basis functions), and then "rendering" them into a porosity volume (with Gaussian noise added at each voxel). The render_layers function encapsulates the MEX/GPU-based renderer. Given this model, we perform inference using tempered MCMC, conditioning on the data shown in Fig. 7. A sample from the posterior shows a reconstructed volume (also shown in Fig. 7) that closely matches the true volume—the program has successfully inferred a parametric form for each layer.

The program calls a function named dpmem, which is a stochastic memoizer similar to that found in Church. dpmem accepts a function representing a base measure (in this case, the function sample_layer) and a concentration parameter, and returns a new function which samples according to a Dirichlet process (DP). We use

The rewritten layer model with naming information.

```
function dp_render_rock( data )
 FUNCTON_PUSH( 1 )
 LINE_PUSH( 0 )
 LINE_SET( 1 )
 num_layers = poissrnd();
 LINE_SET(2)
 dp_sample_layer = dpmem( @sample_layer, 1.0 );
 LINE_SET(3)
 LOOP_PUSH( 0 )
 for i=1:num_layers
   LOOP_INC()
   LINE_SET(4)
   layer(i,:) = dp_sample_layer();
 end:
 LOOP_POP()
 LINE_SET(6)
 rock_volume = render_layers( layers );
 LINE_SET( 7 )
 data = rock_volume + randn( size(rock_volume) );
 FUNCTON_POP()
 LINE_POP()
return:
```

this to create a DP in "layer space," where parameters of layers can be shared. Fig. 7(lower left) shows a comparison of inference with and without the dpmem: shared layers enable the inference engine to find higher-likelihood models faster. Hierarchical DPs are equally easy: we could (for example) let f=dpmem(@randn,1.0), and g=dpmem(@f,1.0) to create an HDP with a Gaussian base measure. We consider such flexibility in defining models—combined with the rich library and type support of Matlab—to be the major virtue of our approach.

4 CONCLUSION

We have described a lightweight implementation technique for probabilistic programming languages. The method is simple and fast, and the resulting languages permit a great deal of flexibility in the specification of distributions. Not only can arbitrary stochastic and deterministic elements be mixed, but arbitrary language features (such as objects, inheritance, operator overloading, closures, recursion, libraries, etc.) may be used. Our example implementations demonstrate compact code bases and reasonable inference speed.

Fortunately, source-to-source transformations are often compositional. It is possible that additional transformations could be applied after the ERP naming transformation to enrich the performance and capabilities of these languages. For example, transformations may exist which automatically reduce redundant computation between traces, or which implement constraint propagation for initializing conditioners. Future work will investigate these issues, as well as the possibility of new languages, such as stochastic Python.

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