

Meta-Learning an Approximate Inference Algorithm for Low-Level Probabilistic Programs

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Abstract

We present a meta-algorithm for learning an approximate posterior-inference algorithm for low-level probabilistic programs that terminate. Our meta-algorithm takes a training set of probabilistic programs that describe models with observations, and attempts to learn an efficient method for inferring the posterior of a similar program. A key feature of our approach is the use of what we call a white-box inference algorithm that extracts information directly from model descriptions themselves, given as programs. Concretely, our white-box inference algorithm is equipped with multiple neural networks, one for each type of atomic command, and computes an approximate posterior of a given probabilistic program by analysing individual atomic commands in the program using these networks. The parameters of the networks are learnt from a training set of programs by our meta-algorithm. We empirically demonstrate that the learnt inference algorithm generalises well to programs that are new in terms of both parameters and model structures, and report important use cases where our approach, in combination with importance sampling (IS), achieves greater test-time efficiency than alternatives such as HMC. The overall results show the promise as well as remaining challenges.

1 Introduction

One key objective of probabilistic programming is to automate reasoning about probabilistic models from diverse domains (Ritchie et al., 2015; Perov & Wood, 2016; Baydin et al., 2019; Schaehtle et al., 2016; Cusumano-Towner et al., 2017; Saad & Mansinghka, 2016; Kulkarni et al., 2015; Young et al., 2019; Jäger et al., 2020). As a way to realize this goal, researchers have extensively worked on the development of posterior-inference or parameter-learning algorithms that are efficient and universal; the algorithms can be applied to all or nearly all models written in probabilistic programming languages (PPLs). This line of research has led to performant probabilistic programming systems (Goodman et al., 2008; Wood et al., 2014; Mansinghka et al., 2014; Minka et al., 2018; Narayanan et al., 2016; Salvatier et al., 2016; Carpenter et al., 2017; Tran et al., 2016; Ge et al., 2018; Bingham et al., 2018). Yet, it also revealed the difficulty of achieving efficiency and universality simultaneously, and the need for equipping PPLs with mechanisms for customising inference or learning algorithms to a given problem. Recent PPLs allow user interactions by, e.g., supporting constructs for specifying conditional independence in a model (Bingham et al., 2018) or for defining proposal distributions (Ritchie et al., 2015; Siddharth et al., 2017; Bingham et al., 2018; Tran et al., 2018; Cusumano-Towner et al., 2019), but oftentimes successful customisation is highly demanding.

In this paper, we provide a limited yet promising answer to the question: can an inference algorithm *learn by experience* and *generalise* to a similar but unseen probabilistic program? We present a meta-algorithm for learning an approximate posterior-inference algorithm itself from a given set of low-level probabilistic programs that terminate, which specifies a “class” of conventional (e.g., hierarchical) or non-conventional (e.g., Rosenbrock, and graphical with nonlinear operations) probabilistic models with observations. The meta-algorithm aims at constructing a customised inference algorithm for the given set, while ensuring universality to the extent that the constructed algorithm generalises within class: it works well for programs not in the training set, as long as the programs are similar to the ones in the set.

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mass ~  $\mathcal{N}(5, 10)$ ; // log of the mass of Milky Way
g1 ~  $\mathcal{N}(mass \times 2, 5)$ ; // for the first satellite galaxy
obs( $\mathcal{N}(g_1, 1), 10$ ); // vel1 = 10 for vel1 ~  $\mathcal{N}(g_1, 1)$ 
g2 ~  $\mathcal{N}(mass + 5, 2)$ ; // for the second satellite galaxy
obs( $\mathcal{N}(g_2, 1), 3$ ) // vel2 = 3 for vel2 ~  $\mathcal{N}(g_2, 1)$ 

```

Figure 1: Probabilistic program for a model for Milky Way and its two satellite galaxies. The `obs` statements refer to the observations of (unnamed) random variables vel_1 and vel_2 .

The distinguished feature of our approach is the use of what we call a white-box inference algorithm, which extracts information directly from model descriptions themselves, given as programs in a PPL. Concretely, our white-box inference algorithm is equipped with multiple neural networks, one for each type of atomic command in a PPL, and computes an approximate posterior for a given program by analysing (or executing in a sense) individual atomic commands in it using these networks. For instance, given the probabilistic program in Fig. 1, which describes a simple model on the Milky Way galaxy, the white-box inference algorithm analyses the program as if an RNN handles a sequence or an interpreter executes a program. Roughly, the algorithm regards the program as a sequence of the five atomic commands (separated by the “;” symbol), initialises its internal state $h \in \mathbb{R}^m$ with h_0 , and transforms the state over the sequence. The internal state h is the encoding of an approximate posterior at the current program point, which corresponds to an approximate filtering distribution of a state-space model. How to update this state over each atomic command is directed by the corresponding neural network. Our meta-algorithm trains the parameters of these networks by trying to make the inference algorithm compute accurate posterior approximations over a training set of probabilistic programs. One can also view our white-box inference algorithm as a message-passing algorithm in a broad sense where transforming the internal state h corresponds to passing a message, and understand our meta-algorithm as a method for learning how to pass a message for each type of atomic commands.

This way of exploiting model descriptions for posterior inference has two benefits. First, it ensures that even after customisation through the neural-network training, the inference algorithm does not lose its universality and can be applied to any probabilistic programs. Thus, at least in principle, the algorithm has a possibility to generalise beyond the training set; its accuracy degrades gracefully as the input probabilistic program diverges from those in the training set. Second, our way of using model descriptions guarantees the efficiency of the inference algorithm (although it does not guarantee the accuracy). The algorithm scans the input program only once, and uses neural networks whose input dimensions are linear in the size of the program. As a result, its time complexity is quadratic over the size of the input program. Of course, the guaranteed speed also indicates that the customisation of the algorithm for a given training set, whose main goal is to achieve good accuracy for probabilistic programs in the set, is a non-trivial process.

Our contributions are as follows: (i) we present a white-box posterior-inference algorithm, which works directly on model description and can be customised to a given model class; (ii) we describe a meta-algorithm for learning the parameters of the inference algorithm; (iii) we empirically analyse our approach with different model classes, and show the promise as well as the remaining challenges.

Related work The difficulty of developing an effective posterior-inference algorithm is well-known, and has motivated active research on learning or adapting key components of an inference algorithm. Techniques for adjusting an MCMC proposal (Andrieu & Thoms, 2008) or an HMC integrator (Hoffman & Gelman, 2014) to a given inference task were implemented in popular tools. Recently, methods for meta-learning these techniques themselves from a collection of inference tasks have been developed (Wang et al., 2018; Gong et al., 2019). The meta-learning approach also features in the work on stochastic variational inference where a variational distribution receives information about each inference task in the form of its dataset of observations and is trained with a collection of datasets (Wu et al., 2020; Gordon et al., 2019; Iakovleva et al., 2020). For a message-passing-style variational-inference algorithm, such as expectation propagation (Minka,

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u := 0; v := 5; w := 1; z1 ~ N(u, v); z2 ~ N(u, v);
z3 ~ N(u, w); μ3 := if (z3 > u) z1 else z2;
obs(N(μ3, w), -1.9); // x1 ~ N(μ3, w), x1 = -1.9
z4 ~ N(u, w); μ4 := if (z4 > u) z1 else z2;
obs(N(μ4, w), -2.2); // x2 ~ N(μ4, w), x2 = -2.2
z5 ~ N(u, w); μ5 := if (z5 > u) z1 else z2;
obs(N(μ5, w), 2.4); // x3 ~ N(μ5, w), x3 = 2.4
z6 ~ N(u, w); μ6 := if (z6 > u) z1 else z2;
obs(N(μ6, w), 2.2) // x4 ~ N(μ6, w), x4 = 2.2

```

Figure 2: Probabilistic program for a simple clustering model on four data points.

2001; Wainwright & Jordan, 2008), Jitkrittum et al. (2015) studied the problem of learning a mechanism to pass a message for a given *single* inference task. A natural follow-up question is how to meta-learn such a mechanism from a dataset of *multiple* inference tasks that can generalise to *unseen* models. Our approach provides a partial answer to the question; our white-box inference algorithm can be viewed as a message-passing-style variational inference algorithm that can meta-learn the representation of messages and a mechanism for passing them for given probabilistic programs.

Amortised inference and inference compilation (Gershman & Goodman, 2014; Le et al., 2017; Paige & Wood, 2016; Stuhlmüller et al., 2013; Kingma & Welling, 2013; Mnih & Gregor, 2014; Rezende et al., 2014; Ritchie et al., 2016; Marino et al., 2018) also attempt to learn a form of a posterior-inference algorithm, but the learnt algorithm has limited scope in terms of generalisation: their learnt algorithm only generalises to unseen inputs or observations of an assumed model, but not to new probabilistic programs. The distinction between theirs and ours is similar to the one between a compiled program and a compiler. Note that the distinction is language-independent; using a higher-level language in those approaches does not blur the distinction.

The idea of running programs with learnt neural networks also appears in the work on training neural networks to execute programs (Zaremba & Sutskever, 2014; Bieber et al., 2020; Reed & de Freitas, 2016). As far as we know, however, we are the first to frame the problem of learning a posterior-inference algorithm as the one of learning to execute.

2 Setup

Our results assume a low-level PPL with guaranteed termination. The PPL is low-level in the sense that it only supports basic forms of expression and conditional statement, and does not support loops and recursion; the design guarantees that there is no (common notion of) execution trace with infinite length. The syntax of the language is given by the following grammar, where r represents a real number, z and v_i variables storing a real, and p the name of a procedure taking two real-valued parameters and returning a real number:

$$\begin{aligned}
\text{Programs } C &::= A \mid C_1; C_2 \\
\text{Atom. Commands } A &::= z \sim \mathcal{N}(v_1, v_2) \mid \text{obs}(\mathcal{N}(v_0, v_1), r) \\
&\quad \mid v_0 := \text{if } (v_1 > v_2) v_3 \text{ else } v_4 \\
&\quad \mid v_0 := r \mid v_0 := v_1 \mid v_0 := p(v_1, v_2)
\end{aligned}$$

Programs in the PPL are constructed by composing atomic commands. The language supports six types of *atomic commands*. The first type is $z \sim \mathcal{N}(v_1, v_2)$, which draws a sample from the normal distribution with mean v_1 and variance v_2 , and assigns the sampled value to z . The second command, $\text{obs}(\mathcal{N}(v_0, v_1), r)$, states that a random variable is drawn from $\mathcal{N}(v_0, v_1)$ and its value is observed to be r . The next is a basic form of a conditional statement that selects one of v_3 and v_4 depending on the result of the comparison

$v_1 > v_2$. The following two commands are different kinds of assignments, one for assigning a constant and the other for copying a value from one variable to another. The last atomic command $v_0 := p(v_1, v_2)$ is a call to one of the known deterministic procedures, which may be standard binary operations such as addition and multiplication, or complex non-trivial functions that are used to build advanced, non-conventional models. When p is a standard binary operation, we use the usual infix notation and write, for example, $v_1 + v_2$, instead of $+(v_1, v_2)$.

We permit only the programs where a variable does not appear more than once on the left-hand side of the $:=$ and \sim symbols. This means that no variable is updated twice or more, and it corresponds to the so-called static single assignment assumption in the work on compilers. This protocol lets us regard variables updated by \sim as latent random variables. We denote those variables by z_1, \dots, z_n .

The choice of the target PPL is intended. The use of basic forms of expression in atomic commands (e.g., only variables as the arguments to a normal distribution or to a procedure p) allows the simple definition (and training) of our white-box inference algorithm; each neural network focuses on exploiting useful information from the kind of atomic command that it is defined for, without having to discharge the complexity that arises from general expressions. While having the important merit, the PPL can serve as an intermediate language for a high-level PPL; the compilation scheme in §3 of (van de Meent et al., 2018), for instance, translates a PPL with general conditional statements and for-loops into graphical models, and can be adopted to compile such programs into our target PPL and hence to make our approach accessible to the end user. Programs with while-loops and recursion cannot generally be translated into our PPL. See Appendix A for further discussion.

Fig. 2 shows a simple model for clustering four data points $\{-1.9, -2.2, 2.4, 2.2\}$ into two clusters, where the cluster assignment of each data point is decided by thresholding a sample from the standard normal distribution. The variables z_1 and z_2 store the centers of the two clusters, and z_3, \dots, z_6 hold the random draws that decide cluster assignments for the data points. See Appendix B for the Milky Way example in Fig. 1 compiled to a program in our language.

Probabilistic programs in the PPL denote unnormalised probability densities over \mathbb{R}^n for some n . Specifically, for a program C , if z_1, \dots, z_n are all the variables assigned by the sampling statements $z_i \sim \mathcal{N}(\dots)$ in C in that order and C contains m observe statements with observations r_1, \dots, r_m , then C denotes an unnormalised density p_C over the real-valued random variables z_1, \dots, z_n : $p_C(z_{1:n}) = p_C(x_{1:m} = r_{1:m} | z_{1:n}) \times \prod_{i=1}^n p_C(z_i | z_{1:i-1})$, where x_1, \dots, x_m are variables not appearing in C and are used to denote observed variables. This density is defined inductively over the structure of C . See Appendix C for details. The goal of our white-box inference algorithm is to compute efficiently the approximate posterior and marginal likelihood estimate accurately for a given C (that is, for the normalised version of p_C and the normalising constant of p_C), when p_C has a finite non-zero marginal likelihood and, as a result, a well-defined posterior density. We next describe how the algorithm attempts to achieve this goal.

3 White-box inference algorithm

Given a program $C = (A_1; \dots; A_k)$, our white-box inference algorithm views C as a sequence of its constituent atomic commands (A_1, A_2, \dots, A_k) , and computes an approximate posterior and a marginal likelihood estimate for C by sequentially processing the A_i 's. Concretely, the algorithm starts by initialising its internal state to $h_0 = \vec{0} \in \mathbb{R}^s$ and the current marginal-likelihood estimate to $Z_0 = 1$. Then, it updates these two components based on the first atomic command A_1 of C . It picks a neural network appropriate for the type of A_1 , applies it to h_0 and gets a new state $h_1 \in \mathbb{R}^s$. Also, it updates the marginal likelihood estimate to Z_1 by analysing the semantics of A_1 . This process is repeated for the remaining atomic commands A_2, A_3, \dots, A_k of C , and eventually produces the last state h_k and estimate Z_k . Finally, the state h_k gets decoded to a probability density on the latent variables of C by a neural network, which together with Z_k becomes the result of the algorithm.

Formally, our inference algorithm is built on top of three kinds of neural networks: the ones for transforming the internal state $h \in \mathbb{R}^s$ of the algorithm; a neural network for decoding the internal states h to probability densities; and the last neural network for approximately solving integration questions that arise from the

marginal likelihood computation in observe statements. We present these neural networks for the programs that sample n -many latent variables z_1, \dots, z_n , and use at most m -many variables (so $m \geq n$). Let \mathbb{V} be $[0, 1]^m$, the space of the one-hot encodings of those m variables, and \mathbb{P} the set of procedure names. Our algorithm uses the following neural networks:

$$\begin{aligned} nn_{\text{sa}, \phi_1} &: \mathbb{V}^3 \times \mathbb{R}^s \rightarrow \mathbb{R}^s, & nn_{\text{ob}, \phi_2} &: \mathbb{V}^2 \times \mathbb{R} \times \mathbb{R}^s \rightarrow \mathbb{R}^s, \\ nn_{\text{if}, \phi_3} &: \mathbb{V}^5 \times \mathbb{R}^s \rightarrow \mathbb{R}^s, & nn_{\text{:=}, \phi_4}^c &: \mathbb{V} \times \mathbb{R} \times \mathbb{R}^s \rightarrow \mathbb{R}^s, \\ nn_{\text{:=}, \phi_5}^v &: \mathbb{V}^2 \times \mathbb{R}^s \rightarrow \mathbb{R}^s, & nn_{p, \phi_p} &: \mathbb{P} \times \mathbb{V}^3 \times \mathbb{R}^s \rightarrow \mathbb{R}^s \quad \text{for every } p \in \mathbb{P}, \\ nn_{\text{de}, \phi_6} &: \mathbb{R}^s \rightarrow (\mathbb{R} \times \mathbb{R})^n, & nn_{\text{intg}, \phi_7} &: \mathbb{V}^2 \times \mathbb{R} \times \mathbb{R}^s \rightarrow \mathbb{R}, \end{aligned}$$

where $\phi_{1:7}$ and ϕ_p for $p \in \mathbb{P}$ are network parameters. The top six networks are for the six types of atomic commands in our language. For instance, when an atomic command to analyse next is a sample statement $z \sim \mathcal{N}(v_1, v_2)$, the algorithm runs the first network nn_{sa} on the current internal state h , and obtains a new state $h' = nn_{\text{sa}, \phi_1}(\bar{z}, \bar{v}_{1:2}, h)$, where \bar{z} and $\bar{v}_{1:2}$ mean the one-hot encoded variables z, v_1 and v_2 . The next nn_{de, ϕ_6} is a decoder of the states h to probability densities over the latent variables z_1, \dots, z_n , which are the product of n independent normal distributions. The network maps h to the means and variances of these distributions. The last nn_{intg, ϕ_7} is used when our algorithm updates the marginal likelihood estimate based on an observe statement $\text{obs}(\mathcal{N}(v_0, v_1), r)$. When we write the meaning of this observe statement as the likelihood $\mathcal{N}(r; v_0, v_1)$, and the filtering distribution for v_0 and v_1 under (the decoded density of) the current state h as $p_h(v_0, v_1)$,¹ the last neural network computes the following approximation: $nn_{\text{intg}, \phi_7}(\bar{v}_{0:1}, r, h) \approx \int \mathcal{N}(r; v_0, v_1) p_h(v_0, v_1) dv_0 dv_1$. See Appendix D for the full derivation of the marginal likelihood.

Given a program $C = (A_1; \dots; A_k)$ that draws n samples (and so uses latent variables z_1, \dots, z_n), the algorithm approximates the posterior and marginal likelihood of C as follows:

$$\begin{aligned} \text{INFER}(C) &= \text{let } h_0 = \vec{0} \in \mathbb{R}^s \text{ and } Z_0 = 1 \in \mathbb{R} \text{ in} \\ &\quad \text{let } (h_k, Z_k) = (\text{INFER}(A_k) \circ \dots \circ \text{INFER}(A_1))(h_0, Z_0) \text{ in} \\ &\quad \text{let } ((\mu_1, \sigma_1^2), \dots, (\mu_n, \sigma_n^2)) = nn_{\text{de}, \phi_6}(h_k) \text{ in return } \left(\prod_{i=1}^n \mathcal{N}(z_i \mid \mu_i, \sigma_i^2), Z_k \right), \end{aligned}$$

where $\text{INFER}(A_i) : \mathbb{R}^s \times \mathbb{R} \rightarrow \mathbb{R}^s \times \mathbb{R}$ picks an appropriate neural network based on the type of A_i , and uses it to transform h and Z :

$$\begin{aligned} \text{INFER}(z \sim \mathcal{N}(v_1, v_2))(h, Z) &= (nn_{\text{sa}}(\bar{z}, \bar{v}_{1:2}, h), Z), \\ \text{INFER}(\text{obs}(\mathcal{N}(v_0, v_1), r))(h, Z) &= (nn_{\text{ob}}(\bar{v}_{0:1}, r, h), Z \times nn_{\text{intg}}(\bar{v}_{0:1}, r, h)), \\ \text{INFER}(v_0 := \text{if } (v_1 > v_2) v_3 \text{ else } v_4)(h, Z) &= (nn_{\text{if}}(\bar{v}_{0:4}, h), Z), \\ \text{INFER}(v_0 := r)(h, Z) &= (nn_{\text{:=}}^c(\bar{v}_0, r, h), Z), \\ \text{INFER}(v_0 := v_1)(h, Z) &= (nn_{\text{:=}}^v(\bar{v}_{0:1}, h), Z), \\ \text{INFER}(v_0 := p(v_1, v_2))(h, Z) &= (nn_p(\bar{v}_{0:2}, h), Z). \end{aligned}$$

We remind the reader that $\bar{v}_{0:k}$ refers to the sequence of the one-hot encodings of variables v_0, \dots, v_k . For the update of the state h , the subroutine $\text{INFER}(A)$ relies on neural networks. For the computation of the marginal likelihood estimate, on the other hand, it exploits prior knowledge that non-observe commands do not change the marginal likelihood (except only indirectly by changing the filtering distribution), and keeps the input Z for those atomic commands without blindly neuralising the computation.

4 Meta-learning parameters

The parameters of our white-box inference algorithm are learnt from a collection of probabilistic programs in our language. Assume that we are given a training set of programs $\mathcal{D} = \{C_1, \dots, C_N\}$ such that each

¹The $p_h(v_0, v_1)$ is a filtering distribution, not prior.

C_i samples n latent variables z_1, \dots, z_n and uses at most m variables. Let $\phi = (\phi_{1:n}, (\phi_p)_{p \in \mathbb{P}})$ be the parameters of all the neural networks used in the algorithm. We learn these parameters by solving the following optimisation problem:²

$$\arg \min_{\phi} \sum_{C \in \mathcal{D}} \text{KL}[\pi_C(z_{1:n}) \| q_C(z_{1:n})] + \frac{\lambda}{2} (N_C - Z_C)^2$$

where $\lambda > 0$ is a hyper-parameter, N_C is the marginal likelihood (or the normalising constant) $\int p_C(z_{1:n}) dz_{1:n}$ for p_C , the next $\pi_C(z_{1:n})$ is the normalised posterior $p_C(z_{1:n})/N_C$ for C , and the last q_C and Z_C are the approximate posterior and marginal likelihood estimate computed by the inference algorithm (that is, $(q_C(z_{1:n}), Z_C) = \text{INFER}(C)$). Note that q_C and Z_C both depend on ϕ , since INFER uses the ϕ -parameterised neural networks.

We optimise the objective by stochastic gradient descent. The key component of the optimisation is a gradient estimator derived as follows: $(\nabla_{\phi} \sum_{C \in \mathcal{D}} \text{KL}[\pi_C \| q_C] + \frac{\lambda}{2} (N_C - Z_C)^2) = (\sum_{C \in \mathcal{D}} \mathbb{E}_{z_{1:n} \sim \pi_C} [-\nabla_{\phi} \log q_C(z_{1:n})] - \lambda(N_C - Z_C) \nabla_{\phi} Z_C) \approx \sum_{C \in \mathcal{D}} -\widehat{L}_{C,\phi} - \lambda(\widehat{N}_C - Z_C) \nabla_{\phi} Z_C$. Here $\widehat{L}_{C,\phi}$ and \widehat{N}_C are sample estimates of $\mathbb{E}_{z_{1:n} \sim \pi_C} [\nabla_{\phi} \log q_C(z_{1:n})]$ and the marginal likelihood, respectively. Both estimates can be computed using standard Monte-Carlo algorithms. For instance, we can run the self-normalising IS with prior as proposal, and generate weighted samples $\{(w^{(j)}, z_{1:n}^{(j)})\}_{1 \leq j \leq M}$ for the unnormalised posterior p_C . Then, we can use these samples to compute the required estimates: $\widehat{N}_C = \frac{1}{M} \sum_{j=1}^M w^{(j)}$ and $\widehat{L}_{C,\phi} = \frac{1}{M} \sum_{j=1}^M (w^{(j)} \nabla_{\phi} \log q_C(z_{1:n}^{(j)})) / \widehat{N}_C$. Alternatively, we may run Hamiltonian Monte Carlo (HMC) (Duane et al., 1987) to generate posterior samples, and use those samples to draw weighted importance samples using, for instance, the layered adaptive IS (Martino et al., 2017). Then, we compute $\widehat{L}_{C,\phi}$ using posterior samples, and \widehat{N}_C using weighted importance samples. Note that neither π_C in $\mathbb{E}_{z_{1:n} \sim \pi_C} [-\nabla_{\phi} \log q_C(z_{1:n})]$ nor N_C depends on the parameters ϕ . Thus, for each $C \in \mathcal{D}$, N_C needs to be estimated only once throughout the entire optimisation process, and the posterior samples from π_C need to be generated only once as well. We use this fact to speed up the computation of each gradient-update step.

5 Empirical evaluation

An effective meta-algorithm should generalise well: the learnt inference algorithm should accurately predict the posteriors of programs unseen during training which have different parameters (§5.1) and model structures (§5.2), as long as the programs are similar to those in the training set. We empirically show that our meta-algorithm learns such an inference algorithm, and demonstrate important use cases where the learnt inference algorithm, when applied to IS as proposal, achieves higher test-time efficiency than alternatives such as HMC (Duane et al., 1987) (§5.3). We implemented our inference algorithm and meta-algorithm using `ocaml-torch` (Mazare, 2018) (Apache-2.0 License). For HMC, we used the NUTS sampler (Hoffman & Gelman, 2014) from the Python interface for Stan (Carpenter et al., 2017) (CC-BY 4.0 license). We used a Ubuntu server with Intel(R) Xeon(R) Gold 6234 CPU @ 3.30GHz with 16 cores, 32 threads, and 263G memory. Table 1 summarizes the full list of the model classes that we considered in our empirical evaluation. See Appendices E and F for the full details of our model classes and experimental setup.

5.1 Generalisation to new model parameters and observations

We evaluated our approach with six model classes: Gaussian models (`gauss`), two hierarchical models (`hierl` and `hierd`), clustering models (`cluster`), Milky Way models (`milky` and `milkyo`), and models (`rb`) using the Rosenbrock function,³ (see Table 1). The purpose of our evaluation is to show the feasibility of our approach, not to develop the state-of-the-art inference algorithm automatically, and also to identify the challenges of the approach. These models are chosen for this purpose. For instance, an inference algorithm should be able to reason about affine transformations and Gaussian distributions (for `gauss`), and dependency relationships

²Strictly speaking, we assume that the marginal likelihood of any $C \in \mathcal{D}$ is non-zero and finite.

³The function is often used to evaluate learning and inference algorithms (Goodman & Weare, 2010; Wang & Li, 2018; Pagani et al., 2019).

Table 1: Full list of the model classes in the empirical evaluation.

Section	Model class	Description	Detail
§5.1	gauss	Gaussian models with a latent variable and an observation where the mean of the Gaussian likelihood is an affine transformation of the latent.	Appendix E.1.1
	hierl	Hierarchical models with three hierarchically structured latent variables.	Appendix E.1.2
	hierd	Hierarchical or multi-level models with both latent variables and data structured hierarchically where data are modelled as a regression of latent variables of different levels.	Appendix E.1.3
	cluster	Clustering models where five observations are clustered into two groups.	Appendix E.1.4
	milky and milkyo	Milky Way models (milky), and their multiple-observations extension (milkyo) where five observations are made for each satellite galaxy.	Appendix E.1.5
	rb	Models with the Rosenbrock function, which is expressed as an external procedure. The main purpose of <code>rb</code> is to show that our approach can in principle handle models with non-trivial computation blocks.	Appendix E.1.6
§5.2	ext1	Models with three Gaussian variables and one deterministic variable storing the value of the function $nl(x) = 50/\pi \times \arctan(x/10)$, where the models have 12 different types — four different dependency graphs of the variables, and three different positions of the deterministic <code>nl</code> variable for each of these graphs.	Appendix E.2.1 (and Fig. 9)
	ext2	Models with six Gaussian variables and one <code>nl</code> variable, which are grouped into five model types based on their dependency graphs.	Appendix E.2.2 (and Fig. 10)
§5.3	mulmod	Three-variable models where two latent variables follow normal distributions and the other stores the value of the function $mm(x) = 100 \times x^3/(10 + x^4)$. The models in this class are grouped into three types defined by their dependency graphs and the positions of <code>mm</code> in the programs.	Appendix E.3 (and Fig. 11)

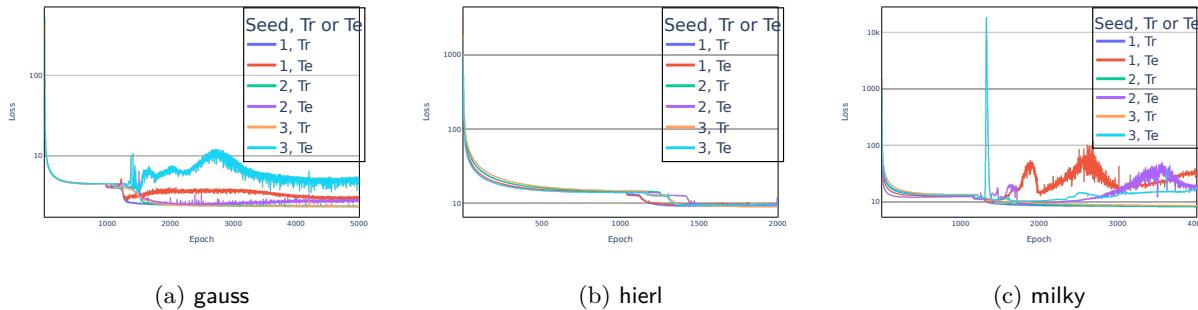


Figure 3: Average training and test losses under three random seeds. The y -axes are log-scaled. The increases in later epochs of Fig. 3c were due to only one or a few test programs out of 50.

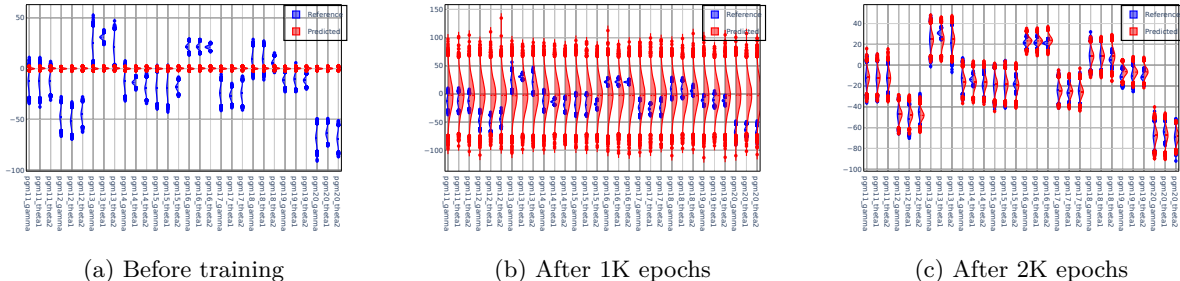


Figure 4: Comparisons of predicted (red) and reference (blue) marginal posteriors for the test programs, with the inference algorithm instantiated at three different training steps: at the initial step, after 1K epochs, and after 2K epochs.

among variables (for `hierl` and `hierd`) to compute a posterior accurately. Successful outcomes in the classes indicate that our approach learns an inference algorithm with such capacity in some cases.

Setup For each model class, we used 400 programs to meta-learn an inference algorithm, and then applied the learnt algorithm to 50 unseen test programs. We measured the average test loss over the 50 test programs, and checked if the loss also decreases when the training loss decreases. We also compared the marginal posteriors predicted by our learnt inference algorithm with the reference marginal posteriors that were computed analytically, or approximately by HMC. When we relied on HMC, we computed the marginal sample means and standard deviations using one of the 10 Markov chains generated by independent HMC runs. Each chain consisted of 500K samples after 50K warmups. We ensured the convergence of the chains using diagnostics such as \hat{R} (Gelman et al., 1992). All training and test programs were automatically generated by a random program generator. This generator takes a program class and hyperparameters (e.g., boundaries of the quantities that are used to specify the models), and returns programs from the class randomly (see Appendix E).

For each training program, our meta-algorithm used 2^{15} samples from the analytic (for `gauss`) or approximate (for the rest, by HMC) posterior distribution for the program.⁴ Similarly, our meta-algorithm computed the marginal likelihood analytically (for `gauss`) or approximately (for the rest) using layered adaptive IS (Martino et al., 2017) where the proposals were defined by an HMC chain. We performed mini-batch training; a single gradient update was done with a training program and a mini batch of size 2^{12} (out of 2^{15} samples for the program). We used Adam (Kingma & Ba, 2015) with its hyperparameters $\{\beta_1 = 0.9, \beta_2 = 0.999, \text{weight_decay} = 0\}$, and the initial learning rate was set to 0.001. When the average training loss

⁴Except for `rb`; see the discussion on Rosenbrock models in Appendix H.

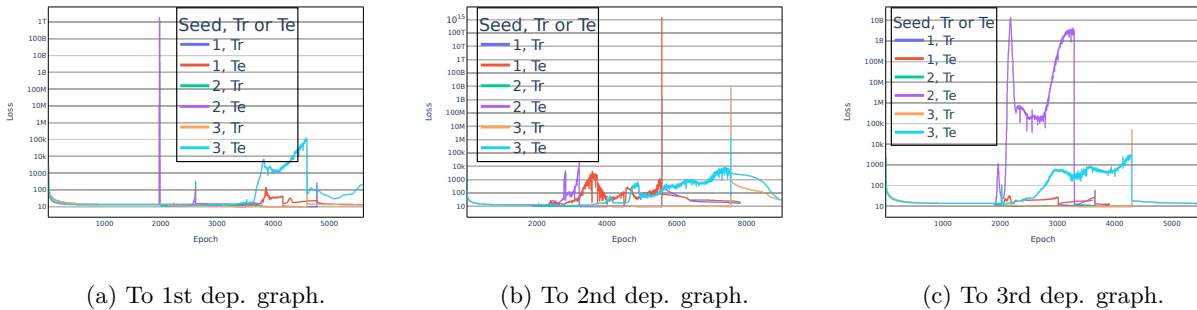


Figure 5: Average losses for generalisation to dependency graphs in ext1. The y-axes are log-scaled.

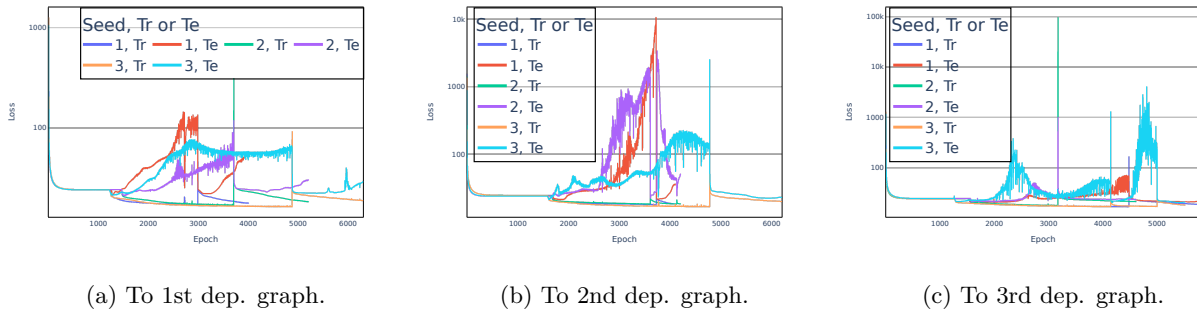


Figure 6: Average losses for generalisation to dependency graphs in ext2. The y-axes are log-scaled.

converged enough, the training stopped. We repeated the same experiments three times using different random seeds.

Results Fig. 3 shows the training and test losses for `gauss`, `hierl`, and `milky` under three random seeds. The losses for the other model classes are in Appendix G. The training loss was averaged over the training set and 8 batch updates, and the test loss over the test set. The training losses in all three experiments decreased rapidly, and more importantly, these decreases were accompanied by the downturns of the test losses, which shows that the learnt parameters generalised to the test programs well. The later part of Fig. 3c shows cases where the test loss increases. This was because the loss of only a few programs in the test set (of 50 programs) became large. Even in this situation, the losses of the rest remained small.

Fig. 4 compares, for 10 test programs in `hierl`, the reference marginal posteriors (blue) and their predicted counterparts (red) by the learnt inference algorithm instantiated at three different training epochs. The predicted marginals were initially around zero (Fig. 4a), evolved to cover the reference marginals (Fig. 4b), and finally captured them precisely in terms of both mean and standard deviation for most of the variables (Fig. 4c). The results show that our meta-algorithm improves the parameters of our inference algorithm, and eventually finds optimal ones that generalise well. We observed similar patterns for the other model classes and random seeds, except for `cluster` and `rb`; programs from these classes often have multimodal posteriors, and we provide an analysis for them in Appendix H.

5.2 Generalisation to new model structures

We let two kinds of model structure vary across programs: the dependency (or data-flow) graph for the variables of a program and the position of a nonlinear function in the program. Specifically, we considered two model classes: (1) models (`ext1`) with three Gaussian variables and one deterministic variable storing the value of the function $nl(x) = 50/\pi \times \arctan(x/10)$, where the models have 12 different types — four different dependency graphs of the variables, and three different positions of the deterministic `nl` variable for each of

Table 2: ESS per sec for the 60 test programs by HMC vs. IS-pred vs. IS-prior.

	HMC			IS-pred			IS-prior		
	GM	Q1	Q3	GM	Q1	Q3	GM	Q1	Q3
ESS	204.8K	4.1K	4.6M	4.2K	2.2K	13.8K	2.8K	1.1K	9.5K
Time	48.2s	27.4s	82.3s	22.7ms	21.4ms	23.0ms	23.1ms	22.1ms	24.0ms
ESS / sec	4.3K	124	127.7K	196.5K	102.4K	646.5K	123.8K	52.6K	436.1K

these graphs; and (2) models (`ext2`) with six Gaussian variables and one nl variable, which are grouped into five types based on their dependency graphs. The evaluation was done for `ext1` and `ext2` independently as in §5.1, but here each of `ext1` and `ext2` itself has programs of multiple (12 for `ext1` and 5 for `ext2`) model types. See Fig. 9 and 10 in the appendix for visualisation of the different model types in `ext1` and `ext2`.

Setup For `ext1`, we ran seven different experiments. Three of them evaluated generalisation to unseen positions of the nl variable, and the other four to unseen dependency graphs. Let $T_{i,j}$ be the type in `ext1` that corresponds to the i -th position of nl and the j -th dependency graph, and $T_{-i,*}$ be all the types in `ext1` that correspond to any nl positions except the i -th and any of four dependency graphs. For generalisation to the i -th position of nl ($i = 1, 2, 3$), we used programs from $T_{-i,*}$ for training and those from $T_{i,*}$ for testing. For generalisation to the j -th dependency graph ($j = 1, 2, 3, 4$), we used programs from $T_{*,j}$ for training and those from $T_{*,j}$ for testing. For `ext2`, we ran five different experiments where each of them tested generalisation to an unseen dependency graph after training with the other four dependency graphs. All these experiments were repeated three times under different random seeds. So, the total numbers of experiment runs were 21 ($= 7 \times 3$) and 15 ($= 5 \times 3$) for `ext1` and `ext2`, respectively.

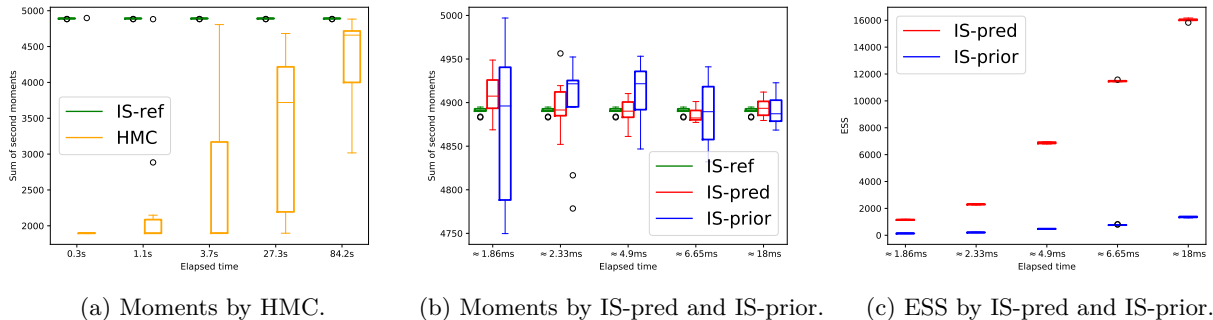
In each experiment run for `ext1`, we used 720 programs for training, and 90 (when generalising to new graphs) or 100 (when generalising to new positions of the nl variable) unseen programs for testing. In each run for `ext2`, we used 600 programs for training and tested the learnt inference algorithm on 50 unseen programs. We ran HMC to estimate posteriors and marginal likelihoods, and used 200K samples after 10K warmups to compute reference posteriors. We stopped training after giving enough time for convergence within a limit of computational resources. The rest was the same as in §5.1.

Results Fig. 5 shows the average training and test losses for generalisation to the first three dependency graphs in `ext1`. Fig. 6 shows the losses for generalisation to the first three dependency graphs in `ext2`. The losses for generalisation to the last dependency graph and to all positions of the nl variable in `ext1`, and those for generalisation to the 4th and 5th dependency graphs in `ext2` are in Appendices I and J. In 17 runs (out of 21) for `ext1`, the decrease in the training losses eventually stabilised or reduced the test losses, even when the test losses were high and fluctuated in earlier training epochs. In 8 runs (out of 15) for `ext2`, the test losses were stabilised as the training losses decreased. In 4 runs out of the other 7, the test losses increased only slightly. In terms of predicted posteriors, we observed highly accurate predictions in 8 runs for `ext1`. For `ext2`, the predicted posteriors were accurate in 7 runs. We also measured the improvements in $\text{KL}[p'(z)||q(z)]$ for all the test programs, where p' is the marginal posterior for a latent variable z ; we refer the reader to Appendix K for the details. Overall, the learnt algorithms generalised to unseen types of models well or fairly well in many cases.

5.3 Application to IS: test-time efficiency in comparison with alternatives

The learnt inference algorithm is a standalone artifact, but another important use case is its application to IS as proposal. In fact, application of variational techniques to samplers such as IS Stites et al. (2021); Jerfel et al. (2021) and MCMC Liang et al. (2021) is an active research, and we demonstrate the inference-time benefits of our application in the setup of generalisation across probabilistic programs.

Specifically, we studied a class (`mulmod`) of three-variable models where two latent variables follow normal distributions and the other stores the value of the function $\text{mm}(x) = 100 \times x^3 / (10 + x^4)$. The models are grouped into three types defined by their dependency graphs and the positions of `mm` in the programs (see Fig.11 in the appendix). We ran our meta-algorithm using 600 programs from all three types using

Figure 7: Moments and ESS for `pgm19` from `mulmod`.

importance samples (not HMC samples). Then for 60 test programs from the last model type, we measured ESS and the sum of second moments along the wall-clock time using three approaches: IS (IS-pred; ours) with the predicted posteriors as proposal using 70K samples, IS (IS-prior) with prior as proposal using 100K samples, and HMC with 1M samples after 500 warmups. As the reference sampler, we used IS (IS-ref) with prior as proposal using 5M samples. All the approaches were repeated 10 times.

Table 2 shows the average ESS per unit time over the 60 test programs, by the three approaches. For HMC, “ESS” and “ESS / sec” were computed using 10 Markov chains, averaged over the 60 programs. For IS-{pred, prior}, “ESS” and “ESS / sec” were averaged over the 10 trials and the 60 programs. GM is the geometric mean, and Q1 and Q3 are the first and third quartiles, respectively. We used the geometric mean, since the ESSes had outliers. The results show that IS-pred achieved the highest ESS per unit time in terms of both mean (GM) and the quartiles (Q1 and Q3). Similarly to §5.2, we also observed the improvements in $KL[p'(z)||q(z)]$ for the marginal distributions (see Appendix K).

We provide further analysis for a test program (`pgm19`; see Appendix L). Fig. 7a and 7b show the moments estimated by HMC and IS-{pred, prior}, respectively, in comparison with the same (across the two figures) reference moments by IS-ref. The estimates by IS-pred (red) quickly converged to the reference (green) within 18ms, while those by HMC (orange) did not converge even after 84s. IS-pred (red) and IS-prior (blue) tended to produce better estimates as the elapsed time increased, but each time, IS-pred estimated the moments more precisely with a smaller variance than IS-prior. In the same runs of the three approaches as in the last columns of Fig. 7a and 7b, IS-pred produced over 16K effective samples in 18ms, while HMC generated only 80 effective samples even after 84s. Similarly, IS-prior generated fewer than 1.4K effective samples in the approximately same elapsed time as in IS-pred. In fact, Fig. 7c shows that as the time increases, the gap between the ESSes of IS-pred and IS-prior gets wider, because the former increases at a rate significantly higher than the latter. Note that IS-pred has to scan a program twice at test time, once for computing the proposal and another for IS with the predicted proposal, whereas IS-prior only once; we observed that the advantage to IS-prior quickly disappeared as the number of IS samples grew (see Appendix M). By manual inspection, we found that the programs in `mulmod` often had multimodal posteriors. We refer the reader to Appendix N for more details.

6 Limitations and future work

Ideally, the meta-algorithm should learn an inference algorithm that effectively solves an inference problem for an arbitrary probabilistic program in the target PPL. Here we clarify the limitations of this work and suggest important future work, as stepping stones to the ultimate goal:

- Though we have demonstrated effective generalisation of the learnt inference algorithm to unseen probabilistic programs, the generalisation is only within a controlled class of modeling problems. For example, the learnt inference algorithm does not generalise to programs with different sizes (Yan et al., 2020); for clustering problems, it means that the learnt algorithm for two-cluster settings is

unapplicable to ten-cluster scenarios. Each model class that we studied assumes a fixed number of variables, and the neural networks crucially exploit the assumption.

- Our meta-algorithm does not scale in practice. When applied to large programs, e.g., state-space models with a few hundred time steps, it cannot learn an optimal inference algorithm within a reasonable amount of time. Overcoming this limitation is important for more realistic problem solving.
- Another direction of improvement is to remove the strong independence assumption (via mean field Gaussian) on the approximating distribution in our inference algorithm, and to equip the algorithm with the capability of generating an appropriate form of the approximation distribution with rich dependency structure, by, e.g., incorporating the ideas from Ambrogioni et al. (2021). This direction is closely related to automatic guide generation in Pyro (Bingham et al., 2018).

7 Conclusion

In this paper, we presented a white-box inference algorithm that computes an approximate posterior and a marginal likelihood estimate by analysing the given program sequentially using neural networks, and a meta-algorithm that learns the network parameters over a training set of probabilistic programs. In our experiments, the meta-algorithm learnt an inference algorithm that generalises well to unseen programs within class, and the learnt inference algorithm brought test-time advantages over alternatives in the important application to IS that we studied. A moral of this work is that the description of a probabilistic model itself has useful information, and learning to exploit the information may lead to an efficient inference. We hope that our work encourages further exploration of this direction.

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A Further discussion about the translation of an expressive PPL into our intermediate language

Programs with recursion or while-loops cannot generally be translated into our PPL, since such programs may go into infinite loops while the programs in our language always terminate. Programs with for-loops and general branches can in theory be translated into a less expressive language such as ours. For example, van de Meent et al. (2018) explain a language called FOPPL (Section 2), which has for-loops and branches, and the translation of FOPPL into graphical models (Section 3). We think that these graphical models can be translated into programs in our PPL. Of course, this does not mean that the learnt inference algorithm would interact well with the compilation; the interaction between compilation and inference in the context of meta-learning is something to be explored in future work.

```

one := 1; t := 2; f := 5; ten := 10;
z1 ~  $\mathcal{N}(f, \textit{ten})$ ; // log of the mass of Milky Way
mass1 := z1 × t;
z2 ~  $\mathcal{N}(\textit{mass}_1, f)$ ; // for the first satellite galaxy
obs( $\mathcal{N}(z_2, \textit{one}), \textit{ten}$ ); // x1 = 10 for x1 ~  $\mathcal{N}(z_2, \textit{one})$ 
mass2 := z1 + f;
z3 ~  $\mathcal{N}(\textit{mass}_2, t)$ ; // for the second satellite galaxy
obs( $\mathcal{N}(z_3, \textit{one}), 3$ ) // x2 = 3 for x2 ~  $\mathcal{N}(z_3, \textit{one})$ 

```

Figure 8: Milky Way example compiled to the probabilistic programming language used in the paper.

B Milky Way example in the probabilistic programming language

Fig. 8 shows the compiled version of the Milky way example to the probabilistic programming language of the paper.

C Formal semantics of the probabilistic programming language

In §2, we stated that a program C in our language denotes an unnormalised density p_C that is factorised as follows:

$$p_C(z_{1:n}) = p_C(x_{1:m} = r_{1:m} | z_{1:n}) \times \prod_{i=1}^n p_C(z_i | z_{1:i-1}).$$

Here z_1, \dots, z_n are all the variables assigned by the sampling statements $z_i \sim \mathcal{N}(\dots)$ in C in that order, the program C contains m observe statements with observations r_1, \dots, r_m , and these observed random variables are denoted by x_1, \dots, x_m . The goal of this section is to provide the details of our statement. That is, we describe the formal semantics of our probabilistic programming language, and from it, we derive a map from programs C to unnormalised densities p_C .

To define the formal semantics of programs in our language, we need a type system that tracks information about updated variables and observations, and also formalises the syntactic conditions that we imposed informally in §2. The type system lets us derive the following judgements for programs C and atomic commands A :

$$(S, V, \alpha) \vdash_1 C : (T, W, \beta), \quad (S, V, \alpha) \vdash_2 A : (T, W, \beta),$$

where S and T are sequences of distinct variables, V and W are sets of variables that do not appear in S and T , respectively, and α and β are sequences of reals. The first judgement says that if before running the

program C , the latent variables in S are sampled in that order, the program variables in V are updated by non-sample statements, and the real values in the sequence α are observed in that order, then running C changes these three data to T , W , and β . The second judgement means the same thing except that we consider the execution of A , instead of C . The triples (S, V, α) and (T, W, β) serve as types in this type system.

The rules for deriving the judgements for C and A follow from the intended meaning just explained. We show these rules below, using the notation $@$ for the concatenation operator for two sequences and also $\text{set}(S)$ for the set of elements in the sequence S :

$$\frac{(R, U, \alpha) \vdash_1 C_1 : (S, V, \beta) \quad (S, V, \beta) \vdash_1 C_2 : (T, W, \gamma)}{(R, U, \alpha) \vdash_1 (C_1; C_2) : (T, W, \gamma)} \quad \frac{(S, V, \alpha) \vdash_2 A : (T, W, \beta)}{(S, V, \alpha) \vdash_1 A : (T, W, \beta)}$$

$$\frac{z \notin \text{set}(S) \cup V \quad v_1, v_2 \in \text{set}(S) \cup V}{(S, V, \alpha) \vdash_2 (z \sim \mathcal{N}(v_1, v_2)) : (S@[z], V, \alpha)} \quad \frac{v_0, v_1 \in \text{set}(S) \cup V}{(S, V, \alpha) \vdash_2 \text{obs}(\mathcal{N}(v_0, v_1), r) : (S, V, \alpha@[r])}$$

$$\frac{v_0 \notin \text{set}(S) \cup V \quad v_1, v_2, v_3, v_4 \in \text{set}(S) \cup V}{(S, V, \alpha) \vdash_2 (v_0 := \text{if } (v_1 > v_2) \text{ } v_3 \text{ else } v_4) : (S, V \cup \{v_0\}, \alpha)}$$

$$\frac{v_0 \notin \text{set}(S) \cup V}{(S, V, \alpha) \vdash_2 (v_0 := r) : (S, V \cup \{v_0\}, \alpha)} \quad \frac{v_0 \notin \text{set}(S) \cup V \quad v_1 \in \text{set}(S) \cup V}{(S, V, \alpha) \vdash_2 (v_0 := v_1) : (S, V \cup \{v_0\}, \alpha)}$$

$$\frac{v_0 \notin \text{set}(S) \cup V \quad v_1, v_2 \in \text{set}(S) \cup V}{(S, V, \alpha) \vdash_2 (v_0 := p(v_1, v_2)) : (S, V \cup \{v_0\}, \alpha)}$$

We now define our semantics, which specifies mappings from judgements for C and A to mathematical entities. First, we interpret each type (S, V, α) as a set, and it is denoted by $\llbracket (S, V, \alpha) \rrbracket$:

$$\begin{aligned} \llbracket (S, V, \alpha) \rrbracket = \{ & (p, f, l) \mid p \text{ is a (normalised) density on } \mathbb{R}^{|S|}, f = (f_v)_{v \in \text{set}(S) \cup V}, \\ & \text{each } f_v \text{ is a measurable map from } \mathbb{R}^{|S|} \text{ to } \mathbb{R}, \\ & l \text{ is a measurable function from } \mathbb{R}^{|S|} \times \mathbb{R}^{|\alpha|} \text{ to } \mathbb{R}_+\}, \end{aligned}$$

where $|S|$ and $|\alpha|$ are the lengths of the sequences S and α , and \mathbb{R}_+ means the set of positive reals. Next, we define the semantics of the judgements $(S, V, \alpha) \vdash_1 C : (T, W, \beta)$ and $(S, V, \alpha) \vdash_2 A : (T, W, \beta)$ that can be derived by the rules from above. The formal semantics of these judgements, denoted by the $\llbracket - \rrbracket$ notation, are maps of the following type:

$$\begin{aligned} \llbracket (S, V, \alpha) \vdash_1 C : (T, W, \beta) \rrbracket : \llbracket (S, V, \alpha) \rrbracket &\rightarrow \llbracket (T, W, \beta) \rrbracket, \\ \llbracket (S, V, \alpha) \vdash_2 A : (T, W, \beta) \rrbracket : \llbracket (S, V, \alpha) \rrbracket &\rightarrow \llbracket (T, W, \beta) \rrbracket. \end{aligned}$$

The semantics is given by induction on the size of the derivation of each judgement, under the assumption that for each procedure name $p \in \mathbb{P}$, we have its interpretation as a measurable map from \mathbb{R}^2 to \mathbb{R} :

$$\llbracket p \rrbracket : \mathbb{R}^2 \rightarrow \mathbb{R}.$$

We spell out the semantics below, first the one for programs and next that for atomic commands.

$$\begin{aligned} \llbracket (S, V, \alpha) \vdash_1 A : (T, W, \beta) \rrbracket(p, f, l) &= \llbracket (S, V, \alpha) \vdash_2 A : (T, W, \beta) \rrbracket(p, f, l), \\ \llbracket (R, U, \alpha) \vdash_1 (C_1; C_2) : (T, W, \gamma) \rrbracket(p, f, l) &= (\llbracket (S, V, \beta) \vdash_2 C_2 : (T, W, \gamma) \rrbracket \\ &\quad \circ \llbracket (R, U, \alpha) \vdash_2 C_1 : (S, V, \beta) \rrbracket)(p, f, l). \end{aligned}$$

Let $\mathcal{N}(a; b, c)$ be the density of the normal distribution with mean b and variance c when $c > 0$ and 1 when $c \leq 0$. For a family of functions $f = (f_v)_{v \in V}$, a variable $w \notin V$, and a function f'_w , we write $f \oplus f'_w$ for the extension of f with a new w -indexed member f'_w .

$$\llbracket (S, V, \alpha) \vdash_2 z \sim \mathcal{N}(v_1, v_2) : (S@[z], V, \alpha) \rrbracket(p, f, l) = (p', f', l')$$

$$\begin{aligned}
& \text{(where } p'(a_{1:|S|+1}) = p(a_{1:|S|}) \times \mathcal{N}(a_{|S|+1}; f_{v_1}(a_{1:|S|}), f_{v_2}(a_{1:|S|})), \\
& \quad f'_v(a_{1:|S|+1}) = f_v(a_{1:|S|}) \text{ for all } v \in V, \quad f'_z(a_{1:|S|+1}) = a_{|S|+1}, \\
& \quad l'(a_{1:|S|+1}, b_{1:|\alpha|}) = l(a_{1:|S|}, b_{1:|\alpha|}), \\
& \llbracket (S, V, \alpha) \vdash_2 \text{obs}(\mathcal{N}(v_0, v_1), r) : (S, V, \alpha @ [r]) \rrbracket (p, f, l) = (p, f, l') \\
& \quad \text{(where } l'(a_{1:|S|}, b_{1:|\alpha|+1}) = l(a_{1:|S|}, b_{1:|\alpha|}) \times \mathcal{N}(b_{|\alpha|+1}; f_{v_1}(a_{1:|S|}), f_{v_2}(a_{1:|S|})), \\
& \llbracket (S, V, \alpha) \vdash_2 (v_0 := \text{if } (v_1 > v_2) v_3 \text{ else } v_4) : (S, V \cup \{v_0\}, \alpha) \rrbracket (p, f, l) = (p, f \oplus f'_{v_0}, l) \\
& \quad \text{(where } f'_{v_0}(a_{1:|S|}) = \text{if } (f_{v_1}(a_{1:|S|}) > f_{v_2}(a_{1:|S|})) \text{ then } f_{v_3}(a_{1:|S|}) \text{ else } f_{v_4}(a_{1:|S|}), \\
& \llbracket (S, V, \alpha) \vdash_2 (v_0 := r) : (S, V \cup \{v_0\}, \alpha) \rrbracket (p, f, l) = (p, f \oplus f'_{v_0}, l) \\
& \quad \text{(where } f'_{v_0}(a_{1:|S|}) = r), \\
& \llbracket (S, V, \alpha) \vdash_2 (v_0 := v_1) : (S, V \cup \{v_0\}, \alpha) \rrbracket (p, f, l) = (p, f \oplus f'_{v_0}, l) \\
& \quad \text{(where } f'_{v_0}(a_{1:|S|}) = f_{v_1}(a_{1:|S|}), \\
& \llbracket (S, V, \alpha) \vdash_2 (v_0 := p'(v_0, v_1)) : (S, V \cup \{v_0\}, \alpha) \rrbracket (p, f, l) = (p, f \oplus f'_{v_0}, l) \\
& \quad \text{(where } f'_{v_0}(a_{1:|S|}) = \llbracket p' \rrbracket (f_{v_0}(a_{1:|S|}), f_{v_1}(a_{1:|S|})).
\end{aligned}$$

Finally, we define p_C for the well-initialised well-typed programs C , i.e., programs C for which we can derived

$$(\llbracket \cdot \rrbracket, \emptyset, \llbracket \cdot \rrbracket) \vdash_1 C : (S, V, \alpha).$$

For such a C , the definition of p_C is given below:

$$p_C(z_{1:|S|}) = p(z_{1:|S|}) \times l(z_{1:|S|}, \alpha)$$

where $(p, _, l) = (\llbracket \llbracket \cdot \rrbracket, \emptyset, \llbracket \cdot \rrbracket \rrbracket \vdash_1 C : (S, V, \alpha)) (p_0, f_0, l_0)$ for the constant-1 functions p_0 and l_0 of appropriate types and the empty family f_0 of functions.

D Marginal likelihood computation: derivation and correctness

Let x_n be the random variable (RV) that is observed by the command $\text{obs}(\mathcal{N}(v_0, v_1), r)$ and $x_{1:(n-1)}$ be the $(n-1)$ RVs that are observed before the command. When our algorithm is about to analyse this observe command, we have (an estimate of) $p(x_{1:(n-1)})$ by induction. Then, the marginal likelihood of $x_{1:n}$ can be computed as follows:

$$\begin{aligned}
& p(x_{1:(n-1)}, x_n) \\
&= \iint p(x_{1:(n-1)}, x_n, v_0, v_1) dv_0 dv_1 \\
&= \iint p(x_{1:(n-1)}) p(v_0, v_1 | x_{1:(n-1)}) p(x_n | x_{1:(n-1)}, v_0, v_1) dv_0 dv_1 \\
&\approx p(x_{1:(n-1)}) \iint p_h(v_0, v_1) p(x_n | x_{1:(n-1)}, v_0, v_1) dv_0 dv_1 \\
& \quad // \text{ The filtering distribution } p(v_0, v_1 | x_{1:(n-1)}) \text{ is approximated by } p_h. \\
&= p(x_{1:(n-1)}) \iint p_h(v_0, v_1) p(x_n | v_0, v_1) dv_0 dv_1 \\
& \quad // \text{ The RV } x_n \text{ is conditionally independent of } x_{1:(n-1)} \text{ given } v_0, v_1. \\
&= p(x_{1:(n-1)}) \iint p_h(v_0, v_1) \mathcal{N}(r; v_0, v_1) dv_0 dv_1 \\
& \quad // p(x_{1:(n-1)}) \text{ is } Z \text{ in the description of } \text{INFER}(A_i) \text{ in Section 4, and the neural network} \\
& \quad // \text{ } nn_{\text{intg}, \phi_7} \text{ aims at approximating the integral term accurately.}
\end{aligned}$$

This derivation leads to the equation in the main text.

In a setting of probabilistic programming where observations are allowed to be different in true and false branches, the marginal likelihood may fail to be defined, and such a setting is beyond the scope of our language. Using variables multiple times or having observe commands spread out in the program does not make differences in the derivation above.

E Detailed descriptions for probabilistic models used in the empirical evaluation

We detail the program specifications for the classes in §5 using the probabilistic programming language in §2, and then describe how our program generator generated programs from those classes randomly.

In the program specifications to follow, randomly-generated constants are written in the Greek alphabets (θ), and latent and other program variables in the English alphabets. Also, we often use more intuitive variable names instead of using z_i for latent variables and v_i for the other program variables, to improve readability. When describing random generation of the parameter values, we let $U(a, b)$ denote the uniform distribution whose domain is $(a, b) \subset \mathbb{R}$; we use this only for describing the random program generation process itself, not the generated programs (only normal distributions are used in our programs, with the notation \mathcal{N}).

E.1 Generalisation to new model parameters and observations

This section details the model classes in §5.1.

E.1.1 gauss

The model class is described as follows:

$$\begin{aligned} m_z &:= \theta_1; v_z := \theta'_2; c_1 := \theta_3; c_2 := \theta_4; v_x := \theta'_5; \\ z_1 &\sim \mathcal{N}(m_z, v_z); z_2 := z_1 \times c_1; z_3 := z_2 + c_2; \\ \text{obs}(\mathcal{N}(z_3, v_x), o) \end{aligned}$$

For each program of the class, our random program generator generated the parameter values as follows:

$$\begin{aligned} \theta_1 &\sim U(-5, 5), \theta_2 \sim U(0, 20), \theta'_2 = (\theta_2)^2, \theta_3 \sim U(-3, 3) \\ \theta_4 &\sim U(-10, 10), \theta_5 \sim U(0.5, 10), \theta'_5 = (\theta_5)^2 \end{aligned}$$

and then generated the observation o by running the program forward where the value for z_1 was sampled from $z_1 \sim U(m_z - 2 \times \sqrt{v_z}, m_z + 2 \times \sqrt{v_z})$.

E.1.2 hierl

The model class is described as follows:

$$\begin{aligned} m_g &:= \theta_1; v_g := \theta'_2; v_{t_1} := \theta'_3; v_{t_2} := \theta'_4; v_{x_1} := \theta'_5; \\ v_{x_2} &:= \theta'_6; g \sim \mathcal{N}(m_g, v_g); t_1 \sim \mathcal{N}(g, v_{t_1}); t_2 \sim \mathcal{N}(g, v_{t_2}); \\ \text{obs}(\mathcal{N}(t_1, v_{x_1}), o_1); \text{obs}(\mathcal{N}(t_2, v_{x_2}), o_2) \end{aligned}$$

For each program of the class, our generator generated the parameter values as follows:

$$\begin{aligned} \theta_1 &\sim U(-5, 5), \theta_2 \sim U(0, 50), \theta'_2 = (\theta_2)^2, \theta_3 \sim U(0, 10) \\ \theta'_3 &= (\theta_3)^2, \theta_4 \sim U(0, 10), \theta'_4 = (\theta_4)^2, \theta_5 \sim U(0.5, 10) \\ \theta'_5 &= (\theta_5)^2, \theta_6 \sim U(0.5, 10), \theta'_6 = (\theta_6)^2 \end{aligned}$$

and then generated the observations o_1 and o_2 by running the program (i.e., simulating the model) forward.

E.1.3 hierd

The model class is described as follows:

$$\begin{aligned}
m_{a_0} &:= \theta_1; v_{a_0} := \theta'_2; v_{a_1} := \theta'_3; v_{a_2} := \theta'_4; m_b := \theta_5; \\
v_b &:= \theta'_6; d_1 = \theta_7; d_2 = \theta_8; v_{x_1} := \theta'_9; v_{x_2} := \theta'_{10}; \\
a_0 &\sim \mathcal{N}(m_{a_0}, v_{a_0}); a_1 \sim \mathcal{N}(a_0, v_{a_1}); a_2 \sim \mathcal{N}(a_0, v_{a_2}); \\
b &\sim \mathcal{N}(m_b, v_b); \\
t_1 &:= b \times d_1; t_2 := a_1 + t_1; \text{obs}(\mathcal{N}(t_2, v_{x_1}), o_1); \\
t_3 &:= b \times d_2; t_4 := a_2 + t_3; \text{obs}(\mathcal{N}(t_4, v_{x_2}), o_2)
\end{aligned}$$

For each program of the class, our generator generated the parameter values as follows:

$$\begin{aligned}
\theta_1 &\sim \text{U}(-10, 10), \theta_2 \sim \text{U}(0, 100), \theta'_2 = (\theta_2)^2, \theta_3 \sim \text{U}(0, 10) \\
\theta'_3 &= (\theta_3)^2, \theta_4 \sim \text{U}(0, 10), \theta'_4 = (\theta_4)^2, \theta_5 \sim \text{U}(-5, 5) \\
\theta_6 &\sim \text{U}(0, 10), \theta'_6 = (\theta_6)^2, \theta_7 \sim \text{U}(-5, 5), \theta_8 \sim \text{U}(-5, 5) \\
\theta_9 &\sim \text{U}(0.5, 10), \theta'_9 = (\theta_9)^2, \theta_{10} \sim \text{U}(0.5, 10), \theta'_{10} = (\theta_{10})^2
\end{aligned}$$

and then generated the observations o_1 and o_2 by running the program forward where the values for a_0 , a_1 , a_2 , and b in this specific simulation were sampled as follows:

$$\begin{aligned}
a_0 &\sim \text{U}(m_{a_0} - 2 \times \sqrt{v_{a_0}}, m_{a_0} + 2 \times \sqrt{v_{a_0}}) \\
a_1 &\sim \text{U}(a_0 - 2 \times \sqrt{v_{a_1}}, a_0 + 2 \times \sqrt{v_{a_1}}) \\
a_2 &\sim \text{U}(a_0 - 2 \times \sqrt{v_{a_2}}, a_0 + 2 \times \sqrt{v_{a_2}}) \\
b &\sim \text{U}(m_b - 2 \times \sqrt{v_b}, m_b + 2 \times \sqrt{v_b})
\end{aligned}$$

E.1.4 cluster

The model class is described as follows:

$$\begin{aligned}
m_{g_1} &:= \theta_1; v_{g_1} := \theta'_2; m_{g_2} := \theta_3; v_{g_2} := \theta'_4; v_x := \theta'_5; \\
g_1 &\sim \mathcal{N}(m_{g_1}, v_{g_1}); g_2 \sim \mathcal{N}(m_{g_2}, v_{g_2}); \\
zero &:= 0; hund := 100; \\
t_1 &\sim \mathcal{N}(zero, hund); m_1 := \text{if } (t_1 > zero) g_1 \text{ else } g_2; \\
\text{obs}(\mathcal{N}(m_1, v_x), o_1); \\
t_2 &\sim \mathcal{N}(zero, hund); m_2 := \text{if } (t_2 > zero) g_1 \text{ else } g_2; \\
\text{obs}(\mathcal{N}(m_2, v_x), o_2); \\
t_3 &\sim \mathcal{N}(zero, hund); m_3 := \text{if } (t_3 > zero) g_1 \text{ else } g_2; \\
\text{obs}(\mathcal{N}(m_3, v_x), o_3); \\
t_4 &\sim \mathcal{N}(zero, hund); m_4 := \text{if } (t_4 > zero) g_1 \text{ else } g_2; \\
\text{obs}(\mathcal{N}(m_4, v_x), o_4); \\
t_5 &\sim \mathcal{N}(zero, hund); m_5 := \text{if } (t_5 > zero) g_1 \text{ else } g_2; \\
\text{obs}(\mathcal{N}(m_5, v_x), o_5)
\end{aligned}$$

For each program of the class, our generator generated the parameter values as follows:

$$\begin{aligned}
\theta_1 &\sim \text{U}(-15, 15), \theta_2 \sim \text{U}(0.5, 50), \theta'_2 = (\theta_2)^2 \\
\theta_3 &\sim \text{U}(-15, 15), \theta_4 \sim \text{U}(0.5, 50), \theta'_4 = (\theta_4)^2 \\
\theta_5 &\sim \text{U}(0.5, 10), \theta'_5 = (\theta_5)^2
\end{aligned}$$

and then generated the observations $o_{1:5}$ by running the program forward.

E.1.5 milky and milkyo

The model class `milky` is described as follows:

$$\begin{aligned}
m_{mass} &:= \theta_1; v_{mass} := \theta'_2; c_1 := \theta_3; v_{g_1} := \theta'_4; c_2 := \theta_5; \\
v_{g_2} &:= \theta'_6; v_{x_1} := \theta'_7; v_{x_2} := \theta'_8; \\
mass &\sim \mathcal{N}(m_{mass}, v_{mass}); \\
mass_1 &:= mass \times c_1; g_1 \sim \mathcal{N}(mass_1, v_{g_1}); \\
mass_2 &:= mass + c_2; g_2 \sim \mathcal{N}(mass_2, v_{g_2}); \\
\text{obs} &(\mathcal{N}(g_1, v_{x_1}), o_1); \text{obs}(\mathcal{N}(g_2, v_{x_2}), o_2)
\end{aligned}$$

For each program of `milky`, our generator generated the parameter values as follows:

$$\begin{aligned}
\theta_1 &\sim \text{U}(-10, 10), \theta_2 \sim \text{U}(0, 30), \theta'_2 = (\theta_2)^2, \theta_3 \sim \text{U}(-2, 2) \\
\theta_4 &\sim \text{U}(0, 10), \theta'_4 = (\theta_4)^2, \theta_5 \sim \text{U}(-5, 5), \theta_6 \sim \text{U}(0, 10) \\
\theta'_6 &= (\theta_6)^2, \theta_7 \sim \text{U}(0.5, 10), \theta'_7 = (\theta_7)^2, \theta_8 \sim \text{U}(0.5, 10) \\
\theta'_8 &= (\theta_8)^2
\end{aligned}$$

and then generated the observations o_1 and o_2 by running the program forward.

Everything remained the same for the `milkyo` class, except that the two `obs` commands were extended to `obs($\mathcal{N}(g_1, v_{x_1}), [o_1, o_2, o_3, o_4, o_5]$)` and `obs($\mathcal{N}(g_2, v_{x_2}), [o_6, o_7, o_8, o_9, o_{10}]$)`, respectively, and all the observations were generated similarly by running the extended model forward.

E.1.6 rb

The model class `rb` is described as follows:

$$\begin{aligned}
m_{z_1} &:= \theta_1; v_{z_1} := \theta'_2; m_{z_2} := \theta_3; v_{z_2} := \theta'_4; v_x := \theta'_5; \\
z_1 &\sim \mathcal{N}(m_{z_1}, v_{z_1}); z_2 \sim \mathcal{N}(m_{z_2}, v_{z_2}); r := \text{Rosenbrock}(z_1, z_2); \\
\text{obs} &(\mathcal{N}(r, v_x), o)
\end{aligned}$$

where $\text{Rosenbrock}(z_1, z_2) = 0.05 \times (z_1 - 1)^2 + 0.005 \times (z_2 - z_1^2)^2$. For each program of the class, our generator generated the parameter values as follows:

$$\begin{aligned}
\theta_1 &\sim \text{U}(-8, 8), \theta_2 \sim \text{U}(0, 5), \theta'_2 = (\theta_2)^2, \theta_3 \sim \text{U}(-8, 8) \\
\theta_4 &\sim \text{U}(0, 5), \theta'_4 = (\theta_4)^2, \theta_5 \sim \text{U}(0.5, 10), \theta'_5 = (\theta_5)^2
\end{aligned}$$

and then generated the observation o by running the program forward where the values for z_1 and z_2 in this specific simulation were sampled as follows:

$$\begin{aligned}
z_1 &\sim \text{U}(m_{z_1} - 1.5 \times \sqrt{v_{z_1}}, m_{z_1} + 1.5 \times \sqrt{v_{z_1}}) \\
z_2 &\sim \text{U}(m_{z_2} - 1.5 \times \sqrt{v_{z_2}}, m_{z_2} + 1.5 \times \sqrt{v_{z_2}})
\end{aligned}$$

E.2 Generalisation to new model structures

This section details the model classes, and different types in each model class in §5.2. For readability, we present canonicalised dependency graphs where variables are named in the breadth-first order. In the experiments reported in this section, we used a minor extension of our probabilistic programming language with procedures taking one parameter.

E.2.1 ext1

Fig. 9 shows the dependency graphs for all model types in `ext1`. The variables z_0, z_1, \dots and x_1, x_2, \dots represent latent and observed variables, respectively, and observed variables are colored in gray. The red node in each graph represents the position of the `nl` variable.

Our program generator in this case generates programs from the whole model class `ext1`; it generates programs of all twelve different types in `ext1`. We explain this generation process for the model type (1,1) in Fig. 9, while pointing out that the similar process is applied to the other eleven types. To generate programs of the model type (1,1), we use the following program template:

$$\begin{aligned} m_{z_0} &:= \theta_1; v_{z_0} := \theta'_2; v_{z_2} := \theta'_3; v_{z_3} := \theta'_4; v_{x_1} := \theta'_5; \\ z_0 &\sim \mathcal{N}(m_{z_0}, v_{z_0}); z_1 := \text{nl}(z_0); z_2 \sim \mathcal{N}(z_1, v_{z_2}); z_3 \sim \mathcal{N}(z_2, v_{z_3}); \\ \text{obs}(\mathcal{N}(z_3, v_{x_1}), o_1) \end{aligned}$$

where $\text{nl}(z) = 50/\pi \times \arctan(z/10)$. The generation involves randomly sampling the parameters of this template, converting the template into a program in our language, and creating synthetic observations. Specifically, our generator generates the parameter values as follows:

$$\begin{aligned} \theta_1 &\sim \text{U}(-5, 5), \theta_2 \sim \text{U}(0, 20), \theta'_2 = (\theta_2)^2, \theta_3 \sim \text{U}(0, 20), \theta'_3 = (\theta_3)^2 \\ \theta_4 &\sim \text{U}(0, 20), \theta'_4 = (\theta_4)^2, \theta_5 \sim \text{U}(0.5, 10), \theta'_5 = (\theta_5)^2 \end{aligned}$$

and generates the observation o_1 by running the program forward where the values for $z_{0:3}$ in this specific simulation were sampled (and fixed to specific values) as follows:

$$\begin{aligned} z_0 &\sim \text{U}(m_{z_0} - 2 \times \sqrt{v_{z_0}}, m_{z_0} + 2 \times \sqrt{v_{z_0}}) \\ z_1 &= \text{nl}(z_0) \\ z_2 &\sim \text{U}(z_1 - 2 \times \sqrt{v_{z_2}}, z_1 + 2 \times \sqrt{v_{z_2}}) \\ z_3 &\sim \text{U}(z_2 - 2 \times \sqrt{v_{z_3}}, z_2 + 2 \times \sqrt{v_{z_3}}). \end{aligned}$$

The generator uses different templates for the other eleven model types in `ext1`, while sharing the similar process for generation of the parameters and observations.

E.2.2 ext2

Fig. 10 shows the dependency graphs for all five model types in `ext2`. Programs of these five types are randomly generated by our program generator. As in the `ext1` case, we explain the generator only for one model type, which corresponds to the first dependency graph in Fig. 10. To generate programs of this type, we use the following program template:

$$\begin{aligned} m_{z_0} &:= \theta_1; v_{z_0} := \theta'_2; v_{z_1} := \theta'_3; v_{z_3} := \theta'_4; v_{z_4} := \theta'_5; v_{z_5} := \theta'_6; v_{z_6} := \theta'_7; \\ v_{x_1} &:= \theta'_8; v_{x_2} := \theta'_9; v_{x_3} := \theta'_{10}; v_{x_4} := \theta'_{11}; \\ z_0 &\sim \mathcal{N}(m_{z_0}, v_{z_0}); z_1 \sim \mathcal{N}(z_0, v_{z_1}); z_2 := \text{nl}(z_0); z_3 \sim \mathcal{N}(z_0, v_{z_3}); \\ z_4 &\sim \mathcal{N}(z_1, v_{z_4}); z_5 \sim \mathcal{N}(z_1, v_{z_5}); z_6 \sim \mathcal{N}(z_2, v_{z_6}); \\ \text{obs}(\mathcal{N}(z_4, v_{x_1}), o_1); \text{obs}(\mathcal{N}(z_5, v_{x_2}), o_2); \text{obs}(\mathcal{N}(z_6, v_{x_3}), o_3); \text{obs}(\mathcal{N}(z_3, v_{x_4}), o_4) \end{aligned}$$

In order to generate a program of this model type and observations, our generator instantiates the parameters of the template as follows:

$$\begin{aligned} \theta_1 &\sim \text{U}(-5, 5), \theta_2 \sim \text{U}(0, 10), \theta'_2 = (\theta_2)^2, \theta_3 \sim \text{U}(0, 10), \theta'_3 = (\theta_3)^2, \theta_4 \sim \text{U}(0, 10), \theta'_4 = (\theta_4)^2 \\ \theta_5 &\sim \text{U}(0, 10), \theta'_5 = (\theta_5)^2, \theta_6 \sim \text{U}(0, 10), \theta'_6 = (\theta_6)^2, \theta_7 \sim \text{U}(0, 10), \theta'_7 = (\theta_7)^2 \\ \theta_8 &\sim \text{U}(0, 10), \theta'_8 = (\theta_8)^2, \theta_9 \sim \text{U}(0, 10), \theta'_9 = (\theta_9)^2, \theta_{10} \sim \text{U}(0, 10), \theta'_{10} = (\theta_{10})^2 \\ \theta_{11} &\sim \text{U}(0, 10), \theta'_{11} = (\theta_{11})^2. \end{aligned}$$

Then, it generates the observations $o_{1:4}$ by running the program forward where the values for $z_{0:6}$ in this specific simulation were sampled (and fixed to specific values) as follows:

$$\begin{aligned} z_0 &\sim \text{U}(m_{z_0} - 2 \times \sqrt{v_{z_0}}, m_{z_0} + 2 \times \sqrt{v_{z_0}}) \\ z_1 &\sim \text{U}(z_0 - 2 \times \sqrt{v_{z_1}}, z_0 + 2 \times \sqrt{v_{z_1}}) \end{aligned}$$

$$\begin{aligned}
z_2 &= \text{nl}(z_0) \\
z_3 &\sim \text{U}(z_0 - 2 \times \sqrt{v_{z_3}}, z_0 + 2 \times \sqrt{v_{z_3}}) \\
z_4 &\sim \text{U}(z_1 - 2 \times \sqrt{v_{z_4}}, z_1 + 2 \times \sqrt{v_{z_4}}) \\
z_5 &\sim \text{U}(z_1 - 2 \times \sqrt{v_{z_5}}, z_1 + 2 \times \sqrt{v_{z_5}}) \\
z_6 &\sim \text{U}(z_2 - 2 \times \sqrt{v_{z_6}}, z_2 + 2 \times \sqrt{v_{z_6}}).
\end{aligned}$$

The generator uses different templates for the other four model types in `ext2`, while sharing the similar process for generation of the parameters and observations.

E.3 Application to IS: test-time efficiency in comparison with alternatives

This section details the `mulmod` class in §5.3, which has three different model types. Fig. 11 shows the dependency graphs for all the model types. The red node in each graph represents the position of the `mm` variable. We used all the three types in training, applied the learnt inference algorithm to programs in the third model type, and compared the results with those returned by HMC.

We similarly explain the generator only using the model type corresponding to the first dependency graph in Fig. 11. To generate programs of this type, we use the following program template:

$$\begin{aligned}
m_{z_0} &:= \theta_1; v_{z_0} := \theta'_2; v_{z_1} := \theta'_3; v_{x_1} := \theta'_4; \\
z_0 &\sim \mathcal{N}(m_{z_0}, v_{z_0}); z_1 \sim \mathcal{N}(z_0, v_{z_1}); z_2 := \text{mm}(z_1); \text{obs}(\mathcal{N}(z_2, v_{x_1}), o_1)
\end{aligned}$$

where $\text{mm}(x) = 100 \times x^3 / (10 + x^4)$. For each program in this model type, our generator instantiates the parameter values as follows:

$$\begin{aligned}
\theta_1 &\sim \text{U}(-5, 5), \theta_2 \sim \text{U}(0, 20), \theta'_2 = (\theta_2)^2, \theta_3 \sim \text{U}(0, 20), \theta'_3 = (\theta_3)^2 \\
\theta_4 &\sim \text{U}(0.5, 10), \theta'_4 = (\theta_4)^2
\end{aligned}$$

and synthesises the observation o_1 by running the program forward where the values for $z_{0:2}$ in this specific simulation were sampled (and fixed to specific values) as follows:

$$\begin{aligned}
z_0 &\sim \text{U}(m_{z_0} - 2 \times \sqrt{v_{z_0}}, m_{z_0} + 2 \times \sqrt{v_{z_0}}) \\
z_1 &\sim \text{U}(z_0 - 2 \times \sqrt{v_{z_1}}, z_0 + 2 \times \sqrt{v_{z_1}}) \\
z_2 &= \text{mm}(z_1).
\end{aligned}$$

The generator uses different templates for the other two model types in `mulmod`, while sharing the similar process for instantiation of the parameters and observations.

F Detailed evaluation setup

In our evaluation, the dimension s of the internal state h was 10 (i.e., $h \in \mathbb{R}^{10}$). We used the same neural network architecture for all the neural network components of our inference algorithm `INFER`. Each neural network had three linear layers and used the `tanh` activation. The hidden dimension was 10 for each layer in all the neural networks except for `nmde` where the hidden dimensions were 50. The hyper-parameter in our optimisation objective (§4) was set to $\lambda = 2$ in the evaluation. We did not use GPUs.

Before running our inference algorithm, we canonicalise the names of variables in a given program based on its dependency (i.e., data-flow) graph. Although not perfect, this preprocessing removes a superficial difference between programs caused by different variable names, and enables us to avoid unnecessary complexity caused by variable-renaming symmetries at training and inference times.

G Losses for `hierd`, `cluster`, `milkyo`, and `rb`

Fig. 12 shows the average training and test losses under three random seeds for `hierd`, `cluster`, `milkyo`, and `rb`. The later part of Fig. 12a, 12c and 12d shows cases where the test loss surges. This was when the loss of only

a few programs in the test set (of 50 programs) became large. Even in this situation, the losses of the rest remained small. We give analyses for `cluster` and `rb` separately in §H.

H Multimodal posteriors: cluster and rb

The `cluster` and `rb` classes in §5.1 posed another challenge: the models often had multimodal posteriors, and it was significantly harder for our meta-algorithm to learn an optimal inference algorithm. To make the evaluation partially feasible for `rb`, we changed two parts of our meta-algorithm slightly, as well as increasing the size of the test set from 50 to 100. First, we used importance samples instead of samples by HMC, which often failed to converge, to learn an inference algorithm. Second, our random program generator placed some restriction on the programs it generated (e.g., by using tight boundaries on some model parameters), guided by the analysis of the geometry of the Rosenbrock function (Pagani et al., 2019). Consequently, HMC (with 500K samples after 50K warmups) failed to converge for only one fifth of the test programs.

Fig. 13 shows the similar comparison plots between reference and predicted marginal posteriors for 10 test programs of the `rb` type, after 52.4K epochs. Our inference algorithm computed the posteriors precisely for most of the programs except two (`pgm75` and `pgm79`) with significant multimodality. The latent variable `pgm75_z0` had at least two modes at around -10 (visible in the figure) and around 10 (hidden in the figure)⁵. Our inference algorithm showed a mode-seeking behavior for this latent variable. Similarly, the variable `pgm79_z0` had at least two modes in the similar domain region (one shown and one hidden), but this time our inference algorithm showed a mode-covering behavior.

The multimodality issue raises two questions. First, how can our meta-algorithm generate samples from the posterior more effectively so that it can optimise the inference algorithm for classes of models with multimodal posteriors? For example, our current results for `cluster` suffer from the fact that the samples used in the training are often biased (i.e., only from a single mode of the posterior). One possible direction would be to use multiple Markov chains simultaneously and apply ideas from the mixing-time research. Second, how can our white-box inference algorithm catch more information from the program description and find non-trivial properties that may be useful for computing the posterior distributions having multiple modes? We leave the answers for future work.

I Training and test losses for the other cases in ext1

Fig. 14 shows the average training and test losses in the `ext1` experiment runs (under three different random seeds) for generalisation to the last (4th) dependency graph and to all three positions of the `nl` variable.

J Training and test losses for the other cases in ext2

Fig. 15 shows the average training and test losses in the `ext2` experiment runs (under three different random seeds) for generalisation to the 4th and 5th dependency graphs.

K Comparisons in terms of KL

For accuracy, it would be ideal to report $\text{KL}[p||q]$, where p is the fully joint target posterior and q is the predicted distribution. It is, however, hard to compute this quantity since often we cannot compute the density of p . One (less convincing) alternative is to compute $\text{KL}[p'(z)||q(z)]$ for a latent variable z where $p'(z)$ is the best Gaussian approximation (i.e., the best approximation using the mean and standard deviation) for the true marginal posterior $p(z)$, and average the results over all the latent variables of interest. We computed $\text{KL}[p'(z)||q(z)]$ for the test programs from `ext1` and `ext2` in §5.2, and for the three from `mulmod` that are reported in §5.3.

⁵The blue reference plots were drawn using an HMC chain, but the HMC chain got stuck in the mode around -10 for this variable.

For `ext1` and `ext2`, we measured the average $\text{KL}[p'(z)||q(z)]$ over all the latent variables z in the test programs. For instance, if there were 90 test programs and each program had three latent variables, we averaged $90 \times 3 = 270$ KL measurements. In an experiment run for `ext1` (which tested generalisation to an unseen dependency graph), the average KL was around 1.32. In an experiment run for `ext2`, the estimation was around 0.95. When we replaced q with a normal distribution that is highly flat (with mean 0 and standard deviation 10K), the estimation was 7.11 and 7.57, respectively. The results were similar in all the other experiment runs that were reported in §5.2.

For three randomly picked test programs (`pgm19`, `pgm30`, and `pgm38`) from `mulmod` in §5.3, p' was the best Gaussian approximation whose mean and standard deviation were estimated by the reference importance sampler (IS-ref), and q was either the predicted marginal posterior by the learnt inference algorithm or the best Gaussian approximation whose mean and standard deviation were estimated by HMC. The average KL was around 1.19 when q was the predicted posterior, while the estimation was 40.9 when q was the best Gaussian approximation by HMC. The results demonstrate that the predicted posteriors were more accurate on average than HMC at least in terms of p' .

L Program in §5.3

Fig. 16 shows the program that is reported in §5.3, written in our probabilistic programming language.

M Discussion of the cost of IS-pred vs. IS-prior

Our approach (IS-pred) must scan the given program “twice” at test time, once for computing the proposal using the learnt neural networks and another for running the importance sampler with the predicted proposal, while IS-prior only needs to scan the program once. Although it may seem that IS-prior has a huge advantage in terms of saving the wall-clock time, our observation is that the effect easily disappears as the sample size increases. In fact, going through the neural networks in our approach (i.e., the first scanning of the program) does not depend on the sample size, and so its time cost remains constant given the program; the time cost was 0.6ms for the reported test program (`pgm19`) in §5.3.

N Manual inspection into `mulmod` (§5.3)

Our manual inspection revealed that the programs in `mulmod` often have multimodal posteriors. Fig. 17a shows the posteriors for `{pgm19, pgm30, pgm38}` in the test set, computed by our learnt inference algorithm (without IS), HMC (200K samples after 10K warmups), and IS-ref. The variable `z0` in the three programs had multimodal posteriors. For `pgm19`, the learnt inference algorithm took only 0.6ms to compute the posteriors, while HMC took 120s on average to generate a chain. The predictions (red) from the learnt inference algorithm for `z0` describe the reference posteriors (green) better than those (blue) by HMC in terms of mean, variance, and mode covering.⁶ The contour plots in Fig. 17b to 17d visualise three HMC chains for `pgm19`. Here, HMC failed to converge, and Fig. 17b explains the poor estimate (blue) in the first column of Fig. 17a.

O Broader impact

Our work partially shows that an inference algorithm itself can be learnt from given inference tasks and can generalise to unseen probabilistic models to some degree. If an inference algorithm can reason about the specifics of the problem at hand (customisation) for distinct cases (universality) automatically, practitioners will benefit directly from it even without much knowledge about probabilities and programming languages. This paper shows potential to develop such an automated reasoning over probabilistic models, and eventually to lower the “entry barriers” to probabilistic programming.

⁶Our inference algorithm in a multimodal-posterior case leads to a good approximation in the following sense: the approximating distribution q covers the regions of the modes well, and also approximates the mean and variance of the target distribution accurately. Note that such a q is useful when it is used as the proposal of IS.

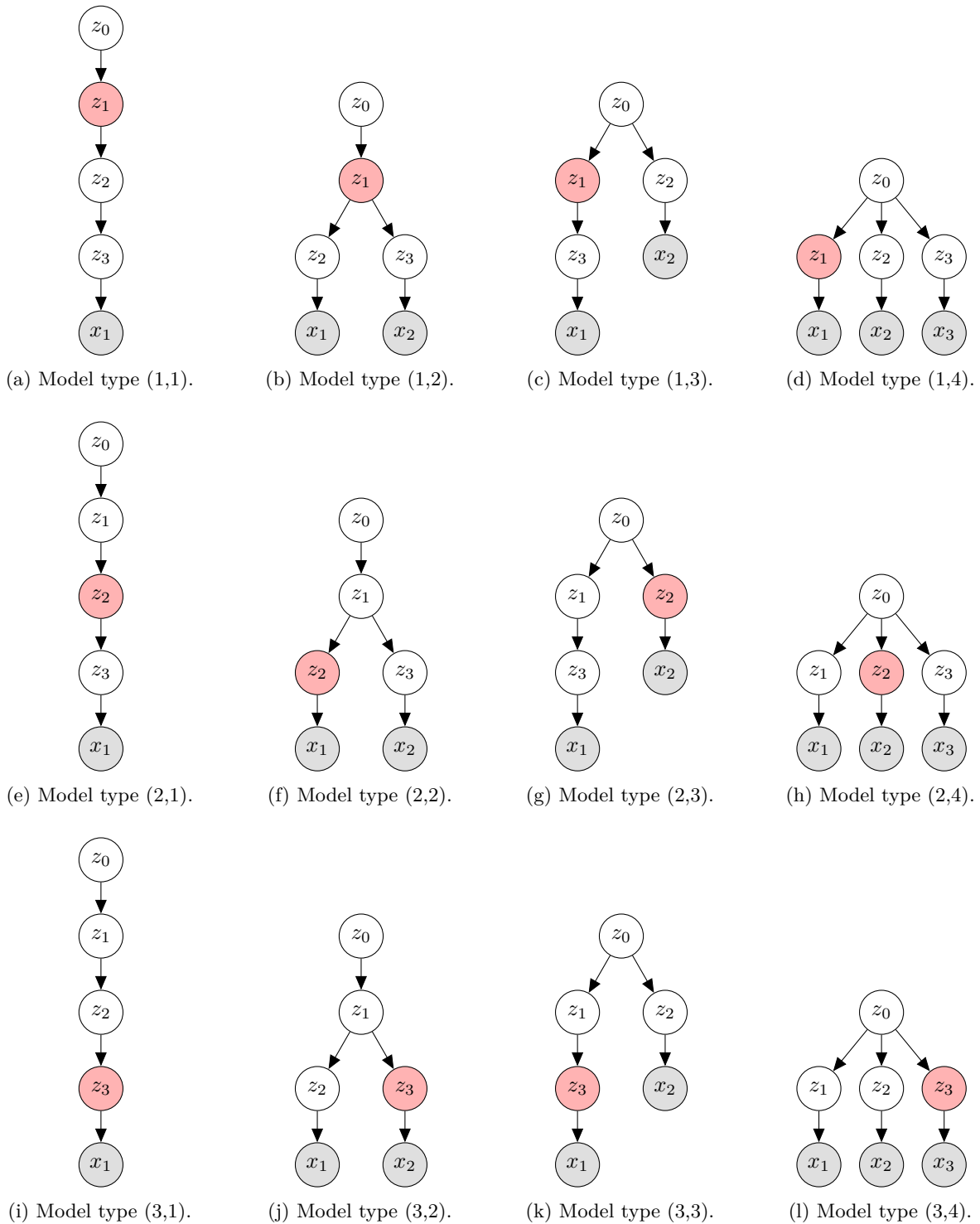


Figure 9: Canonicalised dependency graphs for all 12 model types in ext1. The rows are for different positions of the nl variable, and the columns are for different dependency graphs: the model type (i,j) means one of the 12 model types in ext1 that corresponds to the i -th position of the nl variable and j -th dependency graph.

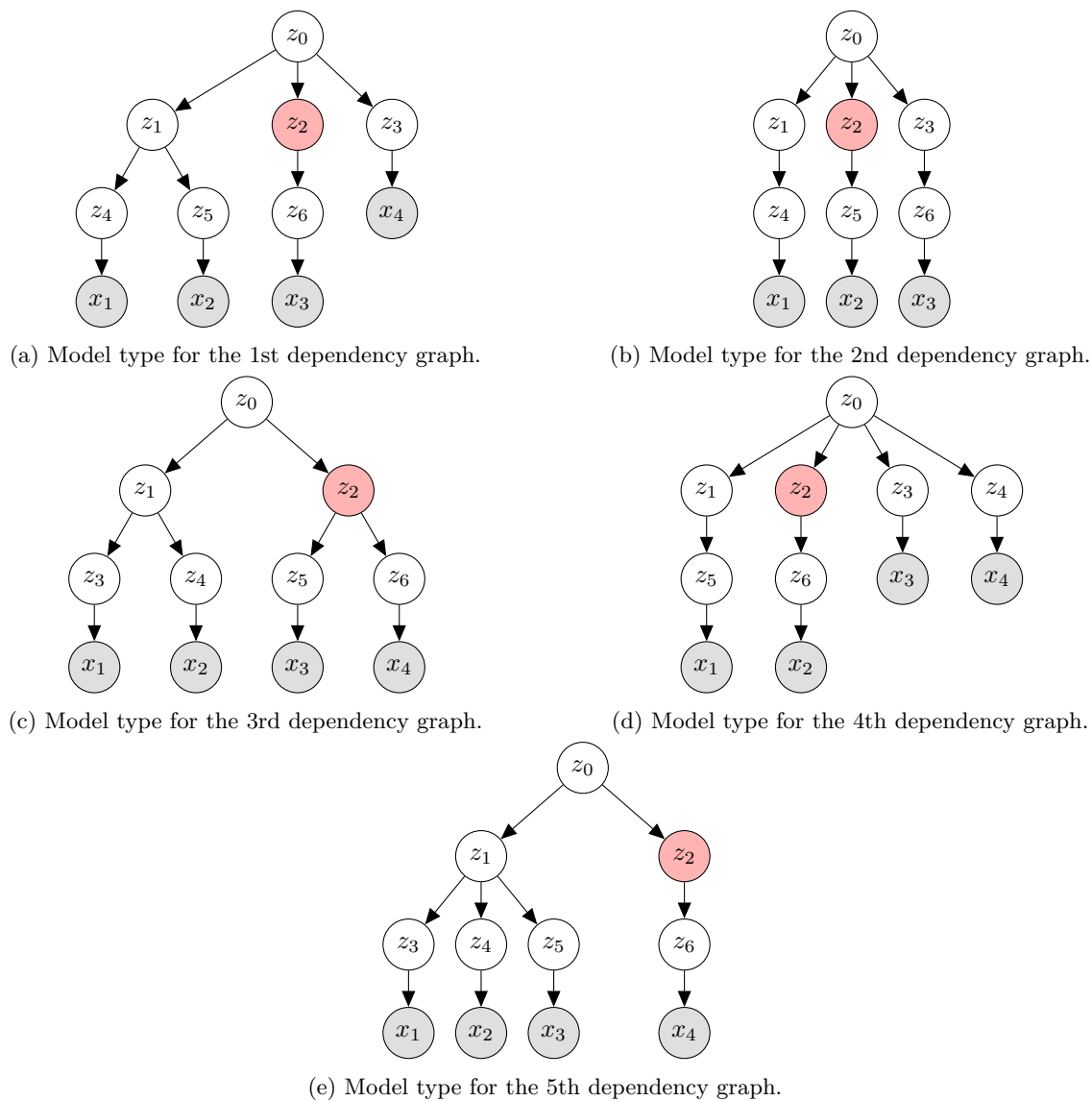


Figure 10: Canonicalised dependency graphs for all five model types in ext2.

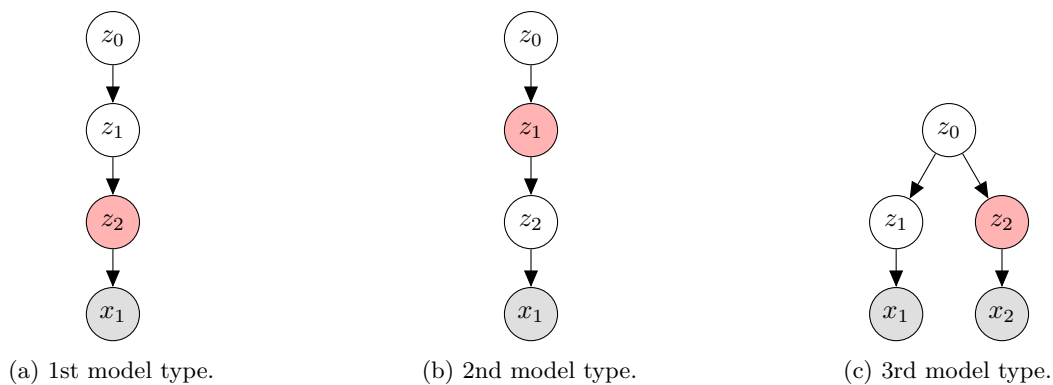


Figure 11: Canonicalised dependency graphs for all three model types in the mulmod class.

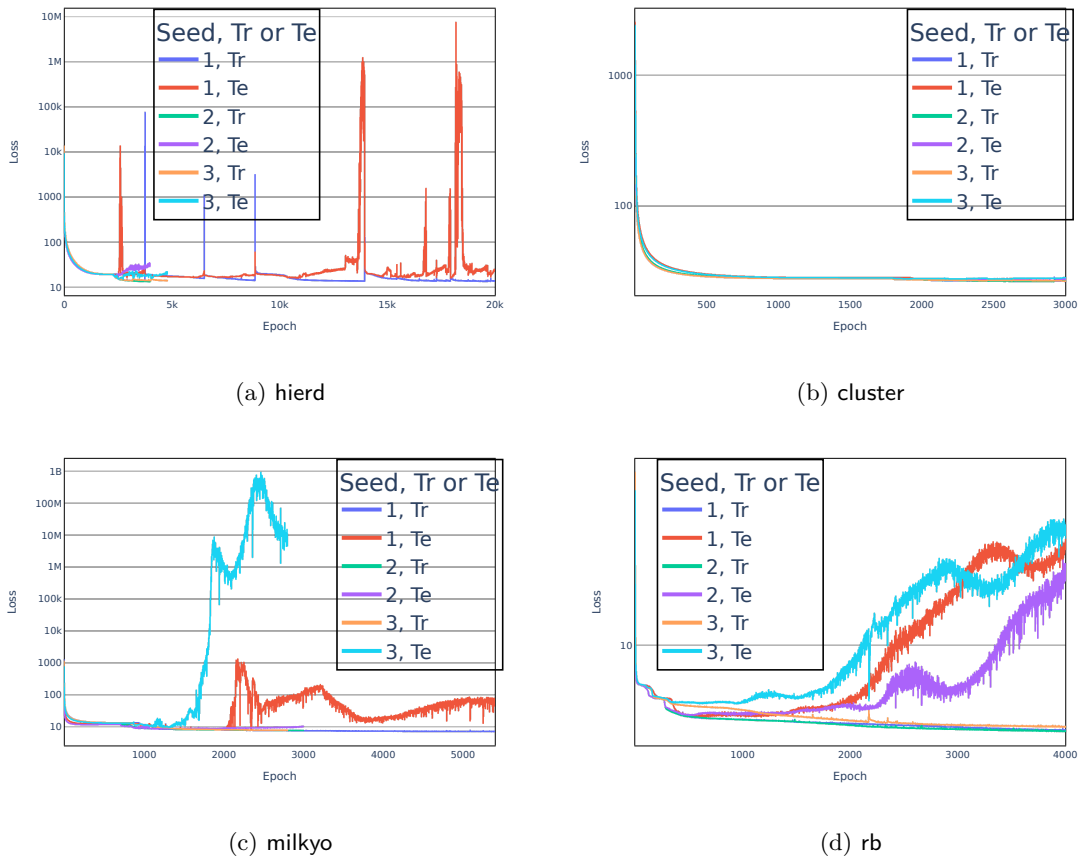


Figure 12: Losses for hierd, cluster, milkyo, and rb. The y -axes are log-scaled. The surges in later epochs of Fig. 12a, 12c and 12d were due to only a single or a few test programs out of 50.

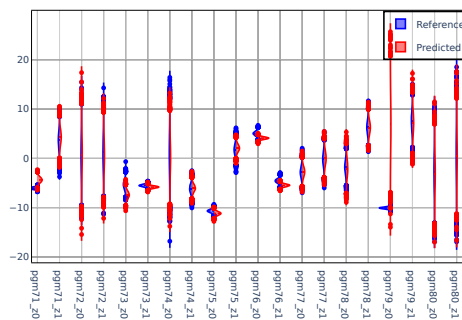


Figure 13: Comparisons of reference and predicted marginal posteriors for 10 programs in the rb test set.

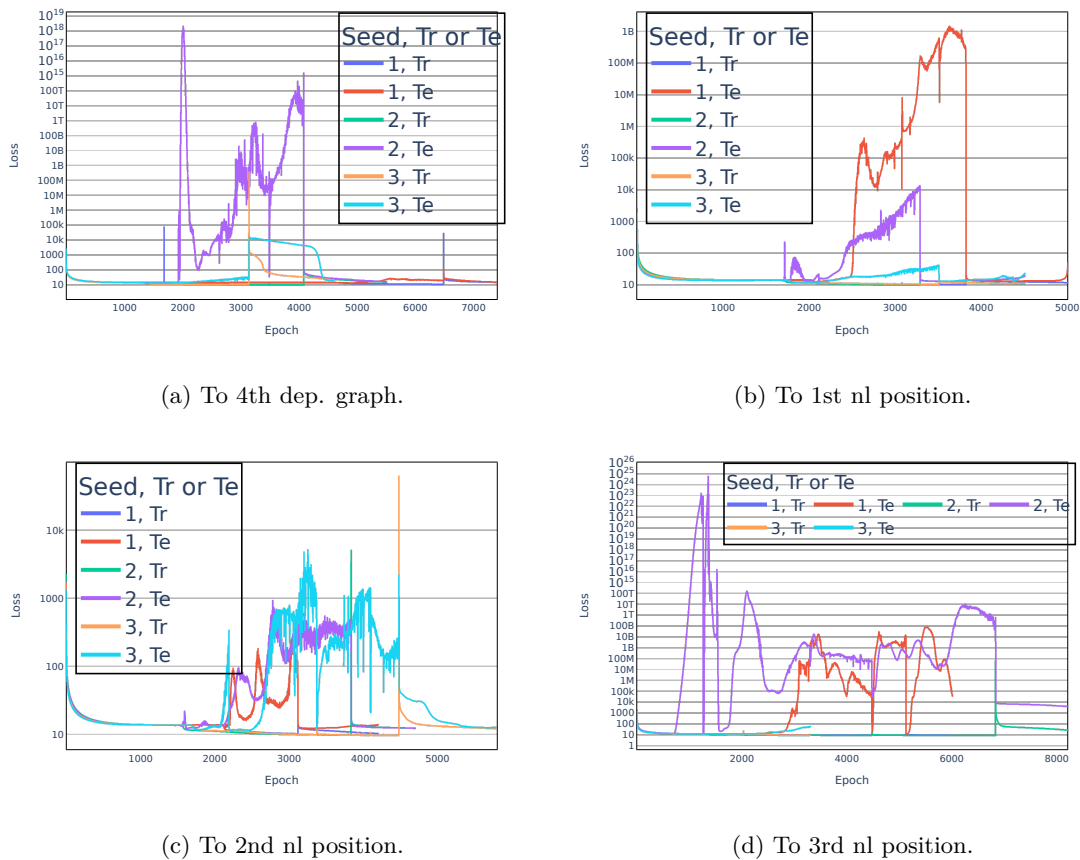


Figure 14: Average training and test losses for generalisation to the last (4th) dependency graph and to all three positions of the nl variable in ext1. The y-axes are log-scaled.

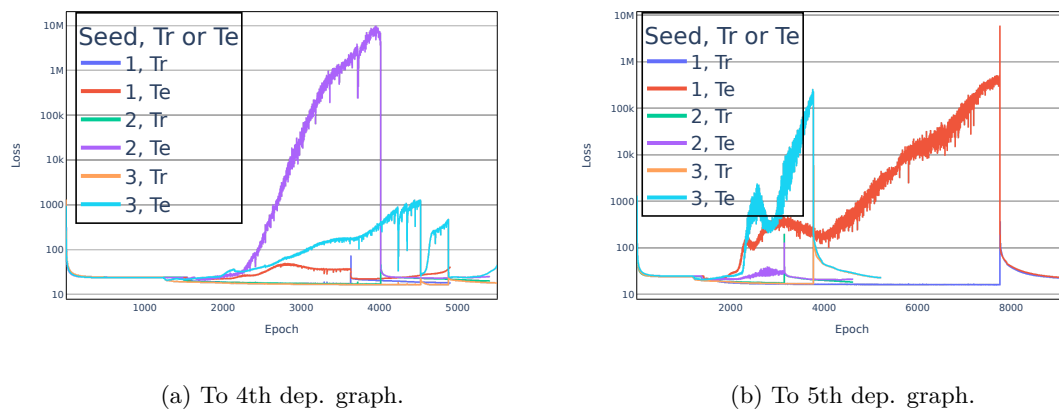


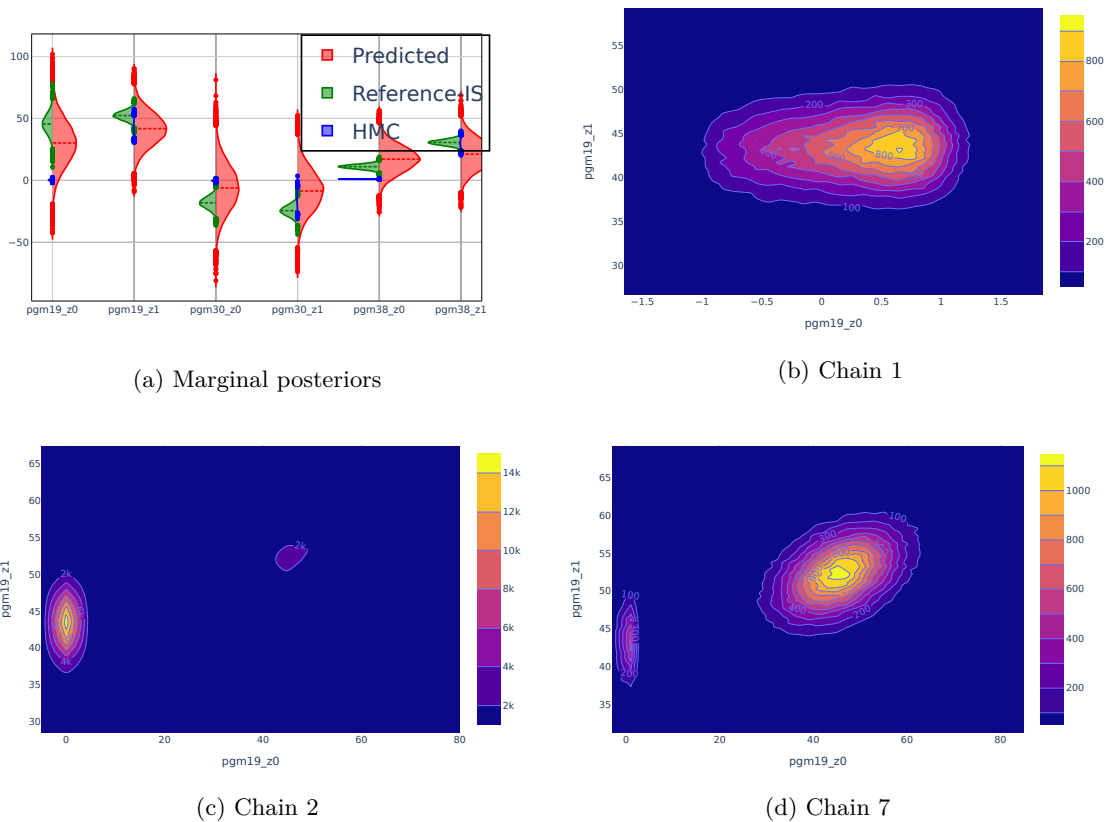
Figure 15: Average training and test losses for generalisation to the 4th and 5th dependency graphs in ext2. The y-axes are log-scaled.

```

a := 3.93; b := 348.16; c := 57.5; d := 14.04; e := 40.34;
z1 ~ N(a, b); z2 ~ N(z1, c); z3 := mm(z1);
obs(N(z2, d), 53.97); obs(N(z3, e), 0.12)

```

Figure 16: The program that is reported in §5.3, written in our probabilistic programming language.

Figure 17: Marginal posteriors, and contours of three HMC chains for `pgm_19` from `mulmod` in §5.3. The x-axis is for `z0` and the y-axis is `z1`.