Scaling Hidden Markov Language Models

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Abstract

Asdf

1 Introduction

Neural network-based generative models have led to progress in difficult tasks such as language modeling and machine translation. However, this progress comes at the cost of interpretability. Neural networks are flexible function approximations, but that flexibility results in opaque models that are difficult to analyze. Rather than post-hoc analysis [**?**], we instead propose to explore the space of models that are defined with interpretability in mind. In particular, we focus on probabilistic graphical models [], which allow for the execution of arbitrary probabilistic queries, i.e. the calculation of (functions of) probability distributions via operations such as conditioning and marginalization [**?**] (simplify). We seek to answer the question of whether we can have models that are both performant and interpretable.

We use language modeling as a benchmark task, as the complex and sometimes long-range phenomena in natural language provides a good testbed. Additionally, language modeling captures import scientific and linguistic questions (more on this later). Language model benchmarks are currently dominated by autoregressive neural models, which makes it difficult to analyze what properties of language the models are actually capturing. For example, it is difficult to analyze the inner representations used in syntax-inspired neural language models [1] (this is my own take on that paper, need to find a better citation). This is an issue if the goal is to learn from models in order to answer scientific questions.

In this work, we investigate scaling probabilistic graphical models which explicitly reason about latent variables. We focus on the one of the simplest latent variable models, the hidden Markov model (HMM). HMMs posit a simple generative process, that first generates a sequence of latent states then the emissions. Additionally, HMMs make very strong conditional independence assumptions. Despite the simplicity and constraints of HMMs, they are still computationally expensive to scale due to the cost of marginalization.

In order to scale HMMs, we propose a series of choices in parameterization that result in computational speedups for inference in HMMs: sparse emission constraint, state dropout, and softmax kernel feature map approximation.

Before diving into these contributions, we first review HMMs.

2 Hidden Markov Models for Language Modeling

Hidden Markov models (HMMs) have a rich history in natural language processing. Speech recognition [4], part of speech tagging [3], and word alignment in machine translation [5]. Have seen some use in neural attention-based models (cite posterior attention). We focus on applying HMMs to the task of language modeling, where the goal is to model the tokens in a sentence $x = (x_1, \dots, x_T)$.

HMMs are cool because ?.

Formally, HMMs have the following generative process: for each $t \in [T]$ timestep, first choose a latent state $z_t \in \mathcal{Z}$, where $|\mathcal{Z}|$ is the number of latent states, then choose a token to emit $x_t \in \mathcal{X}$. This defines the joint distribution:

$$p(x, z) = \prod_t p(x_t \mid z_t) p(z_t \mid z_{t-1}), \tag{1}$$

with transitions $p(z_t \mid z_{t-1})$, emissions $p(x_t \mid z_t)$, and a distinguished start state $z_0 = S$.

Training an HMM requires marginalizing over the unobserved $z = (z_1, \dots, z_T)$ in order to obtain the evidence $p(x) = \sum_z p(x, z)$, accomplished via the forward algorithm. The forward algorithm can

be written as a sequence of matrix-vector multiplications: Let the transition operators $\Lambda_t \in \mathbb{R}^{|\mathcal{Z}|\times|\mathcal{Z}|}$ for $t \in [2, \dots, T]$, with entries

$$[\Lambda_t]_{z_t, z_{t-1}} = p(x_t \mid z_t) p(z_t \mid z_{t-1}),$$

with the first operator given by

$$[\Lambda_1]_{z_1, z_0} = \begin{cases} p(x_1 \mid z_1) p(z_1 \mid z_0 = S) & z_0 = S \\ 0 & \text{otherwise.} \end{cases}$$

The evidence is then given by

$$p(x) = \mathbf{1}^\top \Lambda_1 \Lambda_2 \cdots \Lambda_T \mathbf{1}, \tag{2}$$

where $\mathbf{1} \in \mathbb{R}^{|\mathcal{Z}|\times 1}$ is the column vector of all ones.

On a serial machine, the cost of computing each matrix-vector product in the above equation takes time $O(|\mathcal{Z}|^2)$, resulting in a total running time of $O(T|\mathcal{Z}|^2)$ for the forward algorithm. The quadratic dependence on the number of states precludes scaling to extremely large state spaces.

gradient computation requires $O(T|\mathcal{Z}|^2)$ space

3 Kernel Parameterization

Recent work in latent variable modeling with neural components has shown that the parameterization of conditional distributions can have a large effect on generalization [2]. In particular, the use of neural parameterizations can lead to stronger generalization than full rank parameterizations. We will now make this notion more precise.

rewrite for more generality

Consider the emission distribution $p(x \mid z; \theta)$, with discrete random variables x, z and model parameters θ.[1][2] The full rank parameterization defines the model parameters $\theta \in \mathbb{R}^{|\mathcal{Z}|\times|\mathcal{X}|}$, giving a one-to-one correspondence between the distribution parameters and model parameters:

$$p(x = j \mid z = i) = [\theta]_{i,j}.$$

A kernel parameterization instead defines model parameters $\theta = \{U, V\}$, with $U \in \mathbb{R}^{|\mathcal{Z}|\times d}, V \in \mathbb{R}^{|\mathcal{X}|\times d}$,[3] and obtains the distributional parameters via

$$p(x = j \mid z = i) = \frac{K(u_i, v_j)}{\sum_{j'} K(u_i, v_{j'})},$$

where u_i and v_j are the ith and jth rows of U and V, and $K : \mathbb{R}^d \times \mathbb{R}^d \to \mathbb{R}$ is a kernel. There are a variety of valid kernels K one could employ; however, we focus on the exponential kernel, given by $K(u, v) = \exp(u^\top v)$, as it is widely used in practice as part of the softmax function.

(turn into footnote, or experimental detail) Finally, a neural parameterization takes a further step in abstraction by parameterizing the embedding matrices U, V as functions of neural networks, allowing for more parameter sharing. For the embedding matrix $U \in \mathbb{R}^{|\mathcal{Z}|\times d}$, this entails having another underlying embedding matrix $U' \in \mathbb{R}^{|\mathcal{Z}|\times d'}$ generating the rows $u_i = f_U(u'_i)$, where the same f_U is used for all $i \in \mathcal{Z}$. In this work, we use the following parameterization for f: (resnet thing).

Pros and cons of each, implications? Probably make this a subsection too.

Add summarizing paragraph about the different levels of abstraction, with a picture. Talk about going up and down the stack? This way when we talk about each technique we can point to exactly what abstraction it operates in.

4 Kernels

In the previous section, we discussed three types of parameterizations: full rank, kernel, and neural. Both the kernel and neural parameterizations hinged on the use of a kernel to model the interaction between the values of random variables. In this section, we will discuss some background on kernels, as well as their use in obtaining an efficient inference algorithm for HMMs.[4]

Kernel methods have a rich history in machine learning, having been applied to nonparametric regression, support vector machines, gaussian processes, among other models, to a large degree of success. In those settings, kernels are intuitively used to measure the similarity between two data points. A key distinction is that in our setting, we use kernels to parameterize conditional distributions. However, similar scaling issues arise in both settings.

[1] The dimension of the model parameters θ will vary depending on the choice of parameterization.

[2] The following parameterizations extend to the transition distribution as well, and any conditional distribution.

[3] Typically U and V are referred to as embedding matrices, as the embeddings of the elements of $\mathcal{Z}$ and $\mathcal{X}$ in $\mathbb{R}^d$ are given by the rows of U and V.

[4] As we will see, the inference procedure extends to more complex models as well.

4.1 Exponential Kernels and Softmax

Kernels, and in particular exponential kernels, are widely used to parameterize conditional distributions in neural networks.

A kernel is a function $K : \mathcal{X} \times \mathcal{X} \to \mathbb{R}$ which intuitively measures the similarity of two elements of the domain $\mathcal{X}$. A kernel is positive definite symmetrics (PDS) if.

Why is softmax / exponential kernel special? Maximum entropy distributions, exponential family. Slides on information projections Power series representation?

5 Kernel Approximations

Taylor approximation, generating functions
limitations?

5.1 Bochner Theorem

5.2 Nystrom

6 Kernelized Inference

7 Generalization: Kernelized Belief Propagation

8 Spectral Methods?

References

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A Gradient Estimator Implementation

In the log-semiring, addition is given by $\bigoplus = \text{L}\Sigma\text{E}$ and multiplication by $\bigotimes = +$. Consider the linear chain CRF, $\bigotimes_t \psi(x_{t-1}, x_t)$, with $\psi(x_{t-1}, t) = f(x_t) \oplus g(x_{t-1}, x_t)$. We would like to compute the gradient of the log partition function, $A = \bigoplus_x \bigotimes_t \psi(x_{t-1}, x_t)$. Recall the gradient identities

$$\begin{aligned} \nabla_a a \bigoplus b &= \frac{\exp(a)}{\exp(a \bigoplus b)} \\ \nabla_a a \bigotimes b &= 1. \end{aligned} \tag{3}$$

We then have

$$\begin{aligned} &\nabla_{\psi(x_a, x_b)} \bigoplus_t \psi(x_{t-1}, x_t) \\ &\qquad = \nabla_{\psi(x_a, x_b)} \bigoplus_t \psi(x_{t-1}, x_t) \end{aligned} \tag{4}$$