An Information-Theoretic Parameter-Free Bayesian Framework for Probing Labeled Dependency Trees from Attention Scores

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Abstract

Figuring out how neural language models comprehend syntax acts as a key to revealing how they understand languages. We systematically analyzed methods of extracting syntax from models, namely *probing*, and found five limitations yet widely exist in previous probing practice. We proposed a method that can directly extract labeled dependency trees from attention scores without training any network, while being able to calculate the mutual information (MI) in a mathematical-rigorous way. Compared with previous approaches, our method has a much *simpler* model, while being able to probe more *complex* dependency trees, providing much more fine-grained information about model *explanation* at the same time. We demonstrated our method's effectiveness by systematically comparing it with a great many competitive baselines, and gained informative conclusions, shedding light on our method's explanation potential. Our code is included in the "software" materials of the openreview system to keep anonymity, and we'll make them publicly available upon publication.

1 Introduction

Recent advancements in Large Language Models (LLMs) have left the world with deep impressions. This process is accompanied by confusion, since LLMs are usually trained on simple next-token-prediction LM tasks, while languages have complex hierarchical structures, known as syntax, and recent brain science has proved that humans implicitly build up syntax structures while reading (Lopopolo et al., 2020; Dotan et al., 2022; Fallon et al., 2024). Thus, figuring out whether language models have learned to implicitly comprehend syntax is the key to revealing the essentials of their intelligence.

There are already methods for extracting syntax (or other linguistics concepts) from model internal states, called *probing* methods. A common practice of probing is to train a supervised classifier network on top of model states (Hewitt and Manning, 2019; Pimentel et al., 2020; Müller-Eberstein et al., 2022) to predict dependency syntax trees, or directly take some model states as evidence for syntax (Htut et al., 2019).

Despite the insights they gave, it is obvious that previous probing methods are **explaining by unexplainibility**: Most of them are introducing external trainable networks to extract syntax, ranging from simple linear mappings (Liu et al., 2019) to deep MLPs (Hewitt and Liang, 2019; Voita and Titov, 2020; Pimentel et al., 2020a) or pseudo attention heads (Pimentel et al., 2022). This is causing a trade-off: Linear mappings are simple and explainable, but have limited expressivity. Deeper networks can fit any co-relationships, but a deep probing network is unexplainable itself, so it's natural to raise the doubt on whether the extracted syntax structures really come from the probed LM, or just the strong probes have learned to unconditionally predict them. Moreover, since modern LLMs have larger hidden dimensions compared with pre-trained models, the trainable networks have to be even larger to fit in the dimensionality, inevitably making them more unexplainable.

If we dive deeper, we might find clues about this bitter tradeoff: previous methods are putting their attention mainly on contextualized hidden states. Since vector-based hidden states have completely different modalities compared with dependency trees, a trainable mapping network is necessary. Using hidden states is also a primary cause for the aforementioned concern of *did the probe learn the*

task: Contextualized embeddings embed abundant semantics, so even though the probed LM knows nothing about syntactics, it's still possible that the deep probing model learns it. As an example, if we see two words, *eat* and *breakfast*, even without any context, there're still good reasons for us to believe that *breakfast* acts as the object of the verb *eat*. This is exactly the case of this concern.

If hidden states are not yet good enough, what's the better choice? Maybe we should *put attention on attention*. Attention is the only component that involves inter-token relationships (MLP and add/norm are applied token-wise), while dependency syntactics is exactly inter-word relationships. Moreover, attention maps are topologically consistent with syntax trees: Attention scores are matrices, while dependency trees can also be described as adjacency matrices. Even the matrix size can be the same (ignoring sub-word tokenization, which is also overridable by simple indexing).

Unfortunately, despite those nice consistencies, to our best knowledge, there're only few works focusing on probing attention (Clark et al., 2019; Htut et al., 2019; Vig and Belinkov, 2019; Ravishankar et al., 2021), only being able to extract inferior or even incomplete dependency trees. There seems to be a contradiction. Why is it? Our opinion is that, due to those consistencies, researchers are **overtrusting attention scores**, which is right another limitation. Since attention scores are softmax-normalized, constituting a probability distribution across tokens, they tend to directly use attention scores as probabilities of a dependency relationship between two words. However, attention scores definitely do not only have this single functionality of syntax, so filtering out highly syntactical attention heads together with transformations on attention scores is necessary.

Based on our analysis, we proposed our method of **I**nformation-theoretic **P**arameter-free **B**ayesian **P**robing (IPBP): Instead of training supervised networks, we chose to directly estimate the multivariate probabilistic distributions between attention scores and dependency relationships. With those distributions, we're able to estimate mutual information (MI) in a mathematically rigorous way, obtaining a good metric for each head's individual importance for each dependency label. We further designed a novel decoding algorithm incorporating the estimated MI and Bayesian posteriors, being able to efficiently reconstruct *labeled* dependency trees, while preventing us from dropping into the trap of directly using attention scores as dependency probabilities. We systematically compared with a series of strong baselines, even with those methods requiring far more complex probe networks, and achieved state-of-the-art head importance estimation and tree-constructing performance. We further derived informative conclusions on the estimated MI and distributions. In a word, our method is addressing the two limitations in an elegant way, while offering vast possibilities for the upcoming conclusion-intensive research thanks to its fine-grained MI and probabilities functions.

2 Related Work

Just after the born of deep contextualize embeddings (Peters et al., 2018) and transformer-based pre-trained models (Devlin et al., 2019), researchers have started to investigate whether or not linguistics properties are embedded in these models (Conneau et al., 2018; Liu et al., 2019; Tenny et al., 2019; Hewitt and Manning, 2019). Then arguments began in this area. The frontline of these arguments is about *what probing model can we use to prevent it from learning the task itself*. While early practices and preliminary works suggested strictly-linear probes (Alain and Bengio, 2017; Hewitt and Manning, 2019; Liu et al., 2019), Hewitt and Liang, 2019 proposed control tasks that penalizes models being ability to learn the task itself, and had attempts on several Deep MLPs. Furthermore, Pimentel et al., 2020b admitted this trade-off and took probing as an accuracy-complexity two-goal optimizing problem, and most radically, Pimentel et al. 2020a, insisted that probes should be as deep and complex as possible since they used them as estimations of $\mathcal{V}$-Information (Xu et al., 2020). Apart from disputes, there are also alternative theories proposed by the researchers, like the code-description-length theory by Voita and Titov, 2020 and the architectural bottleneck principle by Pimentel et al., 2022. These theories can be seen as patches under the supervised probing context since they're also addressing the complexity vs. accuracy tradeoff.

Apart from supervised probes, there're also naïve parameter-free probes, mainly based on extracting dependency trees (or partial dependency arcs) from attention scores (Clark et al. 2019, Vig and Belinkov, 2019; Ravishankar et al., 2021), yielding not yet good enough probing performances. If we take a broader view, we'll also find parameter-free explaination methods for more general-purpose

concept in deep learning research (Mu and Andreas, 2020; Antverg and Belinkov, 2021). Together with some supervised works (Radford et al., 2019; Lakretz et al., 2019; Dalvi et al., 2019), these methods, also called neuron analysis methods, were systematically evaluated by a recent work (Fan et al., 2024). In section 4, we'll systematically compare our methods with the principles of these strong baselines.

3 IPBP Methodology

To foster understanding, we'll first break our method into key points in the first section, and then introduce the details.

3.1 Key Aspects Analysis

Given a sentence $X = x_1 x_2 \dots x_n$ and an arbitrary token pair (x_i, x_j), there might or might not be a dependency arc from x_i to x_j. We define $l^{[i][j]}$ as the variable for which kind of dependency exists from x_i to x_j. $l^{[i][j]}$ can be a specific dependency type like `nsubj`, or ϕ when there's no dependency arc from x_i to x_j. If the sentence is sent into a transformer-based LM, there will also be a series of attention scores from x_i to x_j, namely $a_{b,h}^{[i][j]}$, which stands for the attention score of the h-th attention head from the b-th transformer block. If we take the dataset as a series of token pairs, and get the observation $l^{[i][j]}$ and $a_{b,h}^{[i][j]}$ with respect to each token pair, we'll get two co-occurring dataset-wide variables, L and $A_{b,h}$ which stand for the dependency type and head (b, h)'s attention score with respect to *any* token pair.

Therefore, the goal of our probing can be divided into two:

- **MI Estimation**: estimating mutual information (MI) between L and every $A_{b,h}$: $\mathrm{MI}(L; A_{b,h})$.
- **Tree Reconstruction**: A method of deriving a full dependency tree based on attention scores $A_{b,h}$

Specifically, since L is a discrete variable while $A_{b,h}$ is continuous, the joint distribution is a mixture distribution, and the formula of MI is slightly different from its classical discrete or continuous ones, shown as follows:

$$\mathrm{MI}(L; A_{b,h}) = \sum_{l \in \mathcal{L} \cup \{\phi\}} \int f(l,a) \log \frac{f(l,a)}{P(l) f(a)} \mathrm{d}a \quad (1)$$

Where $\mathcal{L}$ stands for the set of all dependency relationships {`nsubj`, `dobj`, …} and $f(l,a)$, $P(l)$, $f(a)$ is short for the density value of joint distribution $f(L, A_{b,h})$ at $L = l, A_{b,h} = a$, density of marginal distribution $f(A_{b,h})$ at $A_{b,h} = a$, and scalar probability $P(L = l)$.

Moreover, the second goal can be regarded as a Bayesian inference process taking $A_{b,h}$ as evidence and L as hypothesis. The posterior distributions ($f(L = l | A_{b,h} = a)$) are required for tree reconstruction. Therefore, the key to achieving these two goals is those probabilistic distributions.

The above formulation intuitively explains our method in a nutshell, in the following sections we'll dive deep into how we can infer these distributions from the dataset.

3.2 Getting the Distributions

Now assume we have a dataset $\mathcal{D}$ consisting of a series of <*sentence, dependency tree*> pairs. We also have a model with $\mathscr{b}$ blocks and $\mathscr{h}$ attention heads within each block. We first initialize a series of attention score sets $\mathcal{A}_{b,h;l}$ where $b \in \{1 \dots \mathscr{b}\}$, $h \in \{1 \dots \mathscr{h}\}$ and $L \in \mathcal{L} \cup \{\phi\}$. $\mathcal{A}_{b,h;l}$ means all possible attention scores of attention head b, h between token pairs having dependency l.

Then we'll iterate over the dataset. For a specific sentence $X \in \mathcal{D}$, we feed $X = x_1 \dots x_n$ into the model, and for any token pair $\langle x_i, x_j \rangle$ ($i, j \in \{1 \dots n\}$), we'll have its dependency relationship $l^{[i][j]}$ and a series of attention scores $a_{b,h}^{[i][j]}$ for any $b \in \{1 \dots \mathscr{b}\}$ and $h \in \{1 \dots \mathscr{h}\}$. We'll add each attention score $a_{b,h}^{[i][j]}$ to the corresponding attention score set $\mathcal{A}_{b,h;l^{[i][j]}}$. After iteration, all attention score sets will have all possible attention scores for any token pair $\langle x_i, x_j \rangle$ in any sentence $X \in \mathcal{D}$.

After gaining the attention values, we'll estimate those key probabilities. The most intuitive one might be $P(L = l)$, since we can simply take the empirical probability on the dataset as the approximate value, which is $\hat{P}(L = l) = \frac{|\mathcal{A}_{b,h;l}|}{\sum_{l' \in \mathcal{L} \cup \{\phi\}} |\mathcal{A}_{b,h;l'}|}$, where b, h can be any value and this equation means the proportion of token pairs with dependency l among all possible token pairs from the dataset. The tricky ones are the continuous probabilities. Since we already have abundant attention score samples, we'll use Kernel Density Estimation (KDE) to estimate the continuous ones. Specifically, for every possible $\mathcal{A}_{b,h;l}$, we can regard it as

the observation of the attention variable $A_{b,h}$ under the circumstance of $L = l$. The samples in $\mathcal{A}_{b,h;l}$ follow the conditional density of $f(A_{b,h}|L = l)$. We use the Gaussian kernel and take a specific bandwidth B (See Appendix A). Therefore, the kernel density $\hat{f}(A_{b,h}|L = l)$ can be estimated as:

$$\frac{1}{\left|\mathcal{A}_{b,h;l}\right| \cdot B} \sum_{i=1}^{|\mathcal{A}_{b,h;l}|} \frac{1}{\sqrt{2\pi \cdot \sigma_{\mathcal{A}_{b,h;l}}}} \exp\left(-\frac{x_0 - \mathcal{A}_{b,h;l}^{(i)}}{B}\right)^2 \quad (2)$$

Where $\sigma_{\mathcal{A}_{b,h;l}}$ is the standard deviation of $\mathcal{A}_{b,h;l}$, and $\mathcal{A}_{b,h;l}^{(i)}$ means the i-th value of $\mathcal{A}_{b,h;l}$.

Again, if we take the view of Bayesian inference, with $A_{b,h}$ as evidence and L as hypothesis, then the estimated $\hat{f}(A_{b,h}|L)$s can be seen as the *likelihood* densities. Applying the Bayesian theorem, we'll get the following equation:

$$f(L|A_{b,h}) = \frac{f(A_{b,h}|L)P(L)}{f(A_{b,h})} = \frac{f(A_{b,h}, L)}{f(A_{b,h})} \quad (3)$$

This gives us inspirations: the likelihoods $\hat{f}(A_{b,h}|L)$ act as the catalyst of the whole process of IPBP. We can multiply $\hat{f}(A_{b,h}|L)$ with the $\hat{P}(L)$s, which are already gained by empirical probabilities, to get the joint densities $\hat{f}(A_{b,h}, L)$. By summing over all possible Ls, we'll get the estimated marginal density $\hat{f}(A_{b,h})$. Now that each term in the above equation is available, we can get the estimated posterior probability $\hat{f}(L|A_{b,h})$. By now, the probabilities required for the whole IPBP process, as mentioned in Section 3.1, are all set.

3.3 Estimating MI

With all these distributions, we're able to proceed to our two main goals: MI estimation and Tree Reconstruction. However, maybe we should reexamine the MI formulation in Equation 1: the $\mathrm{MI}(L; A_{b,h})$ in Equation 1 measures how much common information head $\langle b, h\rangle$ has about *every* possible dependency relationship. However, there might not be such an all-around attention head that is responsible for *every* possible dependency type, but more probable that some heads are responsible for *certain* dependency labels. This kind of *specialist head* is also the assumption of preliminary attention-analyze work like (Htut et al., 2019). Even though there do exist such versatile heads, an MI corresponding to all dependency types is still too coarse-grained, and not helpful for discovering the different functionality of every attention head. Therefore, we should tweak Equation 1 to make the MI formulation fit this *specialist* assumption. The new formulation is as follows:

$$\mathrm{MI}_{\mathrm{binary}}(l; A_{b,h}) = \int f(l, a)\log\frac{f(l,a)}{P(l)f(a)}\mathrm{d}a + \int f(\neg l, a)\log\frac{f(\neg l,a)}{P(\neg l)f(a)}\mathrm{d}a \quad (4)$$

In that equation, $f(\neg l, a)$ is short for the density value of $f(\neg l, A_{b,h})$ at $A_{b,h} = a$, where $f(\neg l, A_{b,h})$ stands for the joint density between any L other than l and $A_{b,h}$. In practice, this joint density can be gained by marginalizing $\hat{f}(A_{b,h}, L)$ over all possible Ls except for l. $P(\neg l)$ stands for the possibility of dependency relationships other than l, and can be estimated using $1 - \hat{P}(l)$.

3.4 Getting Highly Syntactical Heads

By now, having posterior distributions $\hat{f}(L|A_{b,h})$ and $\mathrm{MI}_{\mathrm{binary}}$ feasible for estimating the independent importance of each dependency type, the road towards the goal of Tree Reconstruction is clear. The basic idea of our approach is: for every dependency relationship $l \in \mathcal{L} \cup \{\phi\}$, we filter out a series of attention heads highly responsible for l, constituting the head set $\mathcal{H}_l$. We then infer the possibilities of dependency arcs of l based on the posteriors of heads from $\mathcal{H}_l$, and use $\mathrm{MI}_{\mathrm{binary}}$ to balance between posteriors conditioned on each head from $\mathcal{H}_l$, forming the overall possibility for a dependency arc with relation l. Finally, we use a decoding algorithm to build the dependency tree based on these overall possibilities.

To filter out $\mathcal{H}_l$, it's natural to set a threshold on $\mathrm{MI}_{\mathrm{binary}}(l; A_{b,h})$. However, since for different dependency relationships, the magnitudes of $\mathrm{MI}_{\mathrm{binary}}$ also differs, setting a fixed threshold for all possible ls will favor those relationships with larger MI values. Therefore, an adaptive threshold conditioning on the relation l is necessary. Remind that mutual information is upper-bounded by the individual entropies of each random variable, in our case, $H(\mathbf{1}_{\{l\}}(L))$ and $H(A_{b,h})$ (where $\mathbf{1}_{\{l\}}(L)$ is the indicator function meaning whether or not L equals to l). Since $A_{b,h}$ is continuous, and the entropy analogs of continuous variables (variational entropies) are known as inferior analogs, possibly non-positive, making it unable to act as an upper bound, we choose to estimate $H(\mathbf{1}_{\{l\}}(L))$ as:

$$\hat{H}\left(\mathbf{1}_{\{l\}}(L)\right) = \hat{P}(L)\log\hat{P}(L) + \hat{P}(\neg L)\log\hat{P}(\neg L) \quad (5)$$

If we divide the MI with the entropy, the resulting proportions $\frac{\mathrm{MI}_{\mathrm{binary}}(l;A_{b,h})}{\hat{H}(\mathbf{1}_{\{l\}}(L))}, \forall l$ will be in a uniform $[0, 1]$ scale. An overall threshold can be set for this proportion, resulting in $\mathcal{H}_l$ for every l.

3.5 Tree-Reconstruction Algorithm

After getting $\mathcal{H}_l$ s, another problem occurs: As mentioned before, previous probing practices mainly aim at building unlabeled trees. Even those supervised dependency parsing methods (Dozat and Manning, 2017; Tian et al., 2022) are training separate networks for predicting arcs, and then predicting labels for those predicted arcs. Therefore, these methods are operating on a simple probability space with only probabilities on *the existence of dependency arcs*. What's more, in their methods, there's only one network responsible for predicting probabilities, our method, on the other hand, has a bunch of posterior probabilities. Therefore, it's necessary that we design a decoding algorithm that not only *balances each posterior* but also constitutes *a uniform probability space*.

We first make an assumption that the overall possibility of dependency arcs is independently conditioned on each head in $\mathcal{H}_l$ (otherwise the problem might be too complex). An ideal resort for balancing each posterior is to treat the prediction of dependency arcs as a *voting problem*: for dependency l, each head $\langle b_i, h_i\rangle \in \mathcal{H}_l$ can be seen as a participant with weight $\mathrm{MI}_{\mathrm{binary}}\left(l; A_{b_i,h_j}\right)$. The probability of a dependency arc of l can be seen as the probability of a series of heads with total weights larger than a proportion (like half or two-thirds) voting *the arc belongs to* l. However, due to the non-discrete weights, the problem cannot be efficiently dynamically programmed, resulting in a search space of $\mathcal{O}(2^{|\mathcal{H}_l|})$, which will be rather inefficient during inference. Instead, we relax this voting problem to an easy-computing while rational form: We take the geometric mean of the posteriors. Specifically, let $\mathrm{GP}_{\mathcal{H}_l}(x_i, x_j; l)$ be the geometrically-averaged probability of an arc of l between tokens x_i and x_j conditioned on heads in $\mathcal{H}_l$. In logarithm space, the geometric mean is:

$$\log \mathrm{GP}_{\mathcal{H}_l}(x_i, x_j; l) = \frac{\sum_{\langle b_k,h_k\rangle} \mathrm{MI}_{\mathrm{binary}}(l;A_{b_k,h_k})\cdot\hat{f}(L=l;A_{b_k,h_k})}{\sum_{\langle b_m,h_m\rangle\in\mathcal{H}_l} \mathrm{MI}_{\mathrm{binary}}(l;A_{b_m,h_m})} \quad (6)$$

This is approximately equivalent to the Logarithmic Opinion Pooling technique widely adopted in Bayesian inference, thus acting as a reasonable approximation when the number of experts (in our case, heads in $\mathcal{H}_l$) is relatively large. However, the problem of Logarithmic Pooling is that, if we sum over all probabilities of each possible dependency relationship (in our $\mathrm{MI}_{\mathrm{binary}}$ case, l and $\neg l$), it is not guaranteed to be 1, recalling the second problem of *uniform probability space*. To resolve this, we build a larger multivariate probability space of $\{0,1\}^{|\mathcal{L}|+1}$. We take the voting process of the dependency between x_i and x_j as $|\mathcal{L}| + 1$ independent votes, the ℓ-th voting votes for the existence of the ℓ -th dependency from $|\mathcal{L}|$, using the $\mathrm{GP}_{\mathcal{H}_l}(x_i, x_j; l)$ in Equation 6 as the probability of *existence*, and $1 - \mathrm{GP}_{\mathcal{H}_l}(x_i, x_j; l)$ as the probability of *non-existence*. The overall probability $P(x_i, x_j; l)$, meaning the probability of an arc of l between tokens x_i and x_j conditioned on *all* highly responsible heads $\mathcal{H}_1 \cup \ldots \cup \mathcal{H}_{|\mathcal{L}|} \cup \mathcal{H}_\phi$, is calculated as follows:

$$P\left(x_i, x_j; l\right) = \mathrm{GP}_{\mathcal{H}_l}(x_i, x_j; l) \cdot \prod_{l'\in|\mathcal{L}|+\{\phi\}-\{l\}}\left[1 - \mathrm{GP}_{\mathcal{H}_l}(x_i, x_j; l)\right] \quad (7)$$

While the probability of not arc between x_i and x_j is $1 - \sum_{l\in|\mathcal{L}|} P\left(x_i, x_j; l\right)$, thus resulting in a valid probability space. By now, the two problems introduced by *multi-head* and *multi-label* are all solved. We're just one step towards building the tree, that is, the decoding algorithm utilizing the overall probabilities. Specifically, following previous supervised dependency parsing works, we're using the Eisner dynamic programming algorithm (Eisner, 1996) as the decoding algorithm. Readers might refer to Appendix A for implementation details of our methods, like hyperparameters and our GPU-optimized KDE and integral methods.

3.6 The Novelty of IPBP

Since MI estimation is a small hot topic in statistics, in case of re-inventing wheels, we've done research on related methods. We found two methods sharing (minor) principles with our method: The first one (Moon et al., 1995) is a method estimating MI between two observations within a time series using KDE. They're doing three individual KDEs, with one multivariate one. While it's a known issue that KDE quickly becomes inferior when variables become more than one, known as the *dimensionality curse*, their method is inevitably introducing errors

(and also unapplicable to our attention-dependency mixed-joint distribution setting). We, instead, dexterously circumvented the curse and made the least number of estimations possible (limited to 1) by exploiting mixed-joint distribution and Bayesian theorems.

Another one is also focusing on mixed joint distribution (Gao et al., 2017). However, they use a kNN-like algorithm to estimate point-wise mutual information (PMI) and average it over the dataset. Their method didn't provide any valid probability distributions, thus offering no possibility of tree reconstruction, and also providing less chance for dataset-level or visualization-based explanations.

4 Experiments

4.1 Baselines

In this section, we're going to systematically compare our method with a series of probing as well as neuron analysis baselines. Corresponding to the two sub-tasks introduced in Section 3.1, we first introduce a series of head-selection baselines, where we replace the estimated MI with other criteria, and keep the tree-construction algorithm unchanged. We'll also compare the tree-construction algorithm with common practices of previous attention-based methods. This is better for illustrating the individual contributions of each corresponding submodule.

For head-selection baselines, we'll start from several strong neuron analysis methods evaluated by a recent paper (Fan et al., 2024):

Probeless (Antverg and Belinkov, 2021): This is a parameter-free method, which gets the correlation scores by calculating mean values with respect to different concepts alongside the dataset. In our situation, we use the following instead of $\mathrm{MI}_{\mathrm{binary}}$:

$$\mathrm{PL}(l; A_{b,h}) = \sum_{l' \in \mathcal{L}+\{\phi\}-\{l\}} \left| \bar{\mathcal{A}}_{b,h;l} - \bar{\mathcal{A}}_{b,h;l'} \right| \quad (8)$$

Where $\bar{\mathcal{A}}_{\cdot,\cdot;\cdot}$ denotes the mean value of a specific attention score set. Note that despite its simplicity, this method is evaluated as the method that is most consistent with others by Fan et al., 2024, thus most robust.

IoU (Mu and Andreas, 2020): This method uses Jaccard Similarity as a correlation criterion. In our implementation, we use the following form:

$$\mathrm{IoU}(l; A_{b,h}) = \frac{\left|\mathcal{A}_{b,h,l} \cap [\tau, +\infty)\right|}{\left|\mathcal{A}_{b,h,l}\right| + \sum_{l' \in \mathcal{L}+\{\phi\}-\{l\}} \left|\mathcal{A}_{b,h,l'} \cap [\tau, +\infty)\right|} \quad (9)$$

Where τ is a threshold serving as selecting a salient score. Following the original authors, we set it to the top 99.5% value among values in $\mathcal{A}_{b,h,l}$.

The Linear Feedforward Family: This method refers to a series of methods performing correlation ranking by training a supervised linear network W_θ. Specifically, the equation below gives a uniform formulation of these methods:

$$W_\theta = \underset{W_\theta \in \Theta}{\operatorname{argmin}} \left[\sum_{X \in \mathcal{D}} \sum_{x_i, x_j \in X \times X} \log P_\theta \left(l = l^{[i][j]} \middle| a^{[i][j]}_{1,1\ldots\mathscr{b},\mathscr{h}} \right) + \right.$$
$$\left. + \lambda_1 \|\theta\|_1 + \lambda_2 \|\theta\|_2 \right] \quad (10)$$

Where W_θ is a matrix of shape $\mathscr{b}\mathscr{h} \times (|\mathcal{L}| + 1)$, and $a^{[i][j]}_{1,1\ldots\mathscr{b},\mathscr{h}}$ denotes the concatenation of attention scores between x_i and x_j for all attention heads, and $P_\theta \left(l^{[i][j]} \middle| a^{[i][j]}_{1,1\ldots\mathscr{b},\mathscr{h}} \right)$ stands for the probability of the ground-truth label estimated by the network. When $\lambda_1 = 1, \lambda_2 = 0$, this equation becomes Lasso (Radford et al., 2019), when $\lambda_1 = 0, \lambda_2 = 1$, it becomes Ridge (Lakretz et al., 2019), and $\lambda_1, \lambda_2 = 1$ corresponds to ElasticNet (Dalvi et al., 2019). We use ElasticNet as a representative. After gaining the trained W_θ, we use the weight entry mapping attention score of head $\langle b, h \rangle$, to the probability of relation l as the correlation value, $\mathrm{LFF}(l; b, h)$.

$\mathcal{V}$-Information: Xu et al., 2020 proposed to use a trainable network as an approximation of conditional probabilities, and use the mean logarithm probabilities as approximations of conditional entropies based on the law of large number. This is the state-of-the-art entropy estimation algorithm, used by previous methods also taking information-theoretic perspectives (Pimentel et al., 2020a, Pimentel et al., 2022). Specifically, in our case, we use $\max H_\mathcal{V}(l|A_{\cdot,\cdot}) - H_\mathcal{V}(l|A_{b,h})$ as replacement of $MI_{\mathrm{binary}}(l; A_{b,h})$, and as the equation shows:

$$H_\mathcal{V}(l|A_{b,h}) = \frac{1}{\sum_{X \in \mathcal{D}} |X|^2} \sum_{X \in \mathcal{D}} \sum_{\langle x_i, x_j \rangle \in X} \left[\mathbf{1}_{\{l\}}(L^{[i][j]}) \log \mathrm{MLP}_{b,h;l}\left(a^{[i][j]}_{b,h}\right) + \right.$$
$$\left. \mathbf{1}_{\mathcal{L}+\{\phi\}-\{l\}}(L^{[i][j]}) \log\left(1 - \mathrm{MLP}_{b,h;l}\left(a^{[i][j]}_{b,h,}\right)\right)\right] \quad (11)$$

Where $\mathrm{MLP}_{b,h;l}(\cdot)$ are deep MLPs individually trained using head $\langle b, h \rangle$ to predict label l.

Under each head-selection setting, for fair comparison, we set a limit of the total number of syntactical heads $\sum_{l \in \mathcal{L} \cup \{\phi\}} |\mathcal{H}_l|$ of 2000.

For the tree-construction alternative, we use **Raw attention score**: Under this setting, we're still using the estimated MI as head importance criteria,

while for a specific head $\langle b, h\rangle \in \mathcal{H}_l$, we use the attention score $a_{b,h}^{[\cdot][\cdot]}$ instead of the posterior $\hat{f}(A_{b,h}, L)$ in the reconstruction algorithm. This simple intuitive is the common underlying principle of previous works focusing on attention (Clark et al., 2019; Vig and Belinkov, 2019; Ravishankar et al., 2021). We found that due to the absence of our estimated posteriors, if $\sum_{l\in\mathcal{L}\cup\{\phi\}}|\mathcal{H}_l|$ reaches 2000, the scores of all heads will be rather noisy. Therefore, we choose to select top-k heads based on MI for each label. We did a grid search and found the top-8 settings have ideal performance.

4.2 Model, Dataset and Metrics

We're using open_llama_7b[1] as our probed model. open_llama_7b is a decoder-based LLM consisting of 32 layers and 32 attention heads within each layer. Compared with pre-trained language models like BERT (Devlin et al., 2019), open_llama_7b might consist of attention heads with rather varied functionalities, offering more insights under the contemporary LLM research context.

Specifically, open_llama_7b is a decoder-based model having triangular-masked attention scores. In implementation, we cache the Key Values of each attention head and use them to re-calculate the unmasked attention scores. While our reconstruction is inevitably introducing "useless" attention scores, we think that it is still necessary for two reasons: 1. Making compromises to the decoder structure will hinder our method from applying to non-decoder models (Chung et al., 2024; Zeng et al., 2024), thus less universal. 2. As sentences become longer, the softmax-normalized scores will be *diluted*. This is more serious for triangular attention since it has rows of varying lengths. While softmax is not bijective, using cached QK to reconstruct the unnormalized scores is inevitable.

Following previous supervised dependency parsing works (Tian et al., 2022), we use Universal Dependencies (UD) 2.9 (Zeman et al., 2021), as dataset, with 39832 sentences in the training set and 1700 sentences in the validation set. UD 2.9 is an English treebank covering texts from multiple sources like literature, news articles, spoken languages, *etc.*, with diverse morphological and grammatical features. We also use labeled attachment scores (LAS), and unlabeled attachment scores (UAS) as metrics.

[1] https://github.com/openlm-research/open_llama

4.3 IPBP Structural Alternatives

Apart from comparing with previous methods, we're also curious about our model's designs. Therefore, we propose two alternative structures:

Positive MI: We noticed that the attention score samples exhibit a long-tail characteristic: most samples come from $\mathcal{A}_\phi$, since most pairs of words don't have dependency arc in between. $\mathcal{A}_\phi$ might be noisy, consisting of various non-syntactic inter-token relationships, and MI estimations based on samples in $\mathcal{A}_\phi$ might be affected by this long tail noisy distribution. Other score sets $\mathcal{A}_1, \ldots \mathcal{A}_{|\mathcal{L}|}$ are having approximately the same magnitudes and their corresponding token pairs are guaranteed to have any dependency relationship. Therefore, we also calculated a more syntactical MI, namely $\mathrm{MI}_{\mathrm{pos}}$, with the following formulation:

$$\mathrm{MI}_{\mathrm{pos}}(L; A_{b,h}) = \sum_{l\in\mathcal{L}} \int f_{\mathrm{pos}}(l,a)\log\frac{f_{\mathrm{pos}}(l,a)}{P_{\mathrm{pos}}(l)f_{\mathrm{pos}}(a)}\mathrm{d}a \tag{12}$$

In that equation, $P_{\mathrm{pos}}(\cdot)$, $f_{\mathrm{pos}}(\cdot,\cdot)$ actually stand for conditional possibilities when $l \neq \phi$, estimated by $\hat{P}_{\mathrm{pos}}(L=l) = \frac{\mathcal{A}_{b,h;l}}{\sum_{l'\in\mathcal{L}}\mathcal{A}_{b,h;l'}}$ and $\hat{f}_{\mathrm{pos}}(L, A_{b,h}) = \hat{f}(A_{b,h}|L)\hat{P}_{\mathrm{pos}}(L)$. During implementation, we'll use a balance factor α and calculate the mixed MI $\mathrm{MI}_{\mathrm{mix}}(\cdot;\cdot) = \alpha\mathrm{MI}_{\mathrm{binary}}(\cdot;\cdot) + (1-\alpha)\mathrm{MI}_{\mathrm{pos}}(\cdot;\cdot)$

Arc First: Unlike previous methods, we're directly obtaining labeled dependency trees, bypassing the process of dependency arc predicting. We're curious about whether it's a good choice. Under this setting, instead of estimating $\hat{f}(A_{b,h}|L=l)$, we'll directly estimate the unlabeled likelihoods $\hat{f}(A_{b,h}|L\in\mathcal{L})$ and $\hat{f}(A_{b,h}|L=\phi)$, and calculate the corresponding multivariate probabilities together with corresponding MI values. We'll compare UAS to check the quality of reconstructed unlabeled trees.

Transposed: Sometimes, we're unsure whether the attended token acts as a dependency head, or a dependant. So we'll let $l^{[i][j]}$ correspond to $a_{b,h}^{[j][i]}$, and repeat the whole IPBP process in this setting.

4.4 Result and Analysis

Results are shown in Table 1. We can see that our method is overperforming all competitive baselines, including the state-of-the-art conditional entropy

Method	UAS	LAS
Probeless	34.8	20.9
IoU	38.3	26.6
ElasticNet	41.9	31.3
V-Information	41.3	20.9
Raw Score	32.3	16.6
IPBP	49.1	30.6
IPBP (transposed)	42.6	28.0
IPBP + MI_{pos}	**49.9**	**34.8**
IPBP (arc only)	36.5	N/A

Table 1: Results of our IPBP and different baselines.

estimation method, $\mathcal{V}$-Information, which shares principles with our method while requiring a much more computational budget. In fact, our implemented $\mathcal{V}$-Information MLP is optimized using several tricks (see Appendix B), while during its training, we still find that the trained MLPs are relatively good at detecting arcs while having poor performances on labeling. This aligns with its low LAS in tree-construction results. An insight can be drawn that supervised methods may still fall behind statistical ones, especially when the data is long-tailed or low-dimensional. Moreover, even though the head-selection settings for the raw-score method is specifically tuned, and the method is already selecting attention heads based on our estimated MI, it still has a great performance gap with our posterior-based method, further justifying the necessity of our posterior-based algorithm.

For structure alternatives, we notice that incorporating MI_{pos} will give performance benefits, shedding light on potential improvements to our methods. The transposed setting will still capture a relatively smaller portion of dependencies. Last but not least, by comparing with our arc-based baseline, we'll find we're actually at a triangular balance, we probed for more *accurate*, also *labeled* trees, while choosing a more *straightforward* method, with no need for individual arc probing.

4.5 Further Analysis

Like previous probing works, we'll also do fine-grained analysis of our reconstructed trees and estimated MI values. Instead of listing up MI and doing trivial analyses, we decide to provide two intriguing and informative conclusions, giving inspirations to upcoming works.

The first conclusion is that **decoder models adaptively capture look-back/ahead dependencies**: Since the masked decoder attention can only *look back*, there are good reasons that dependencies that also *look back* (pointing to front words) can be well captured. What makes it more intriguing is that dependencies *looking ahead* might also be captured in a look-back manner. We draw this conclusion by comparing the top-10 most well-reconstructed labels between original IPBP and the transposed alternative. We find that there're more look-ahead dependencies (5 of 10) under the transposed setting compared with the original setting (3 of 10).

The second conclusion is that **model layers correspond to tree layers** to some extent: lower layers are for local/phrasal dependencies, while higher layers are for global/sentence-wide dependencies. The conclusion is consistent with our intuitions and worth hypothesizing, but it was never justified by previous works, which either lack good MI-like criteria or focus on unlabeled trees. Thanks to the fine-grained MI, we can calculate the MI-weighted layer indices for each label, where smaller weighted indices indicate dependencies having more lower-layer heads responsible for it. Among these top-10 labels, we calculated the Pearson correlation coefficient ρ between the weighted layer index and average depth (maximum distance to leaf nodes) of each dependency label, getting a result of 0.69 with p=0.03 for a null hypothesis of no correlation.

5 Conclusion

We proposed a method that can estimate MI and reconstruct labeled dependency trees without introducing any trainable networks. Indeed, our method is achieving an "impossible triangle": it has *simpler* architectures requiring negligible computation budgets, while producing more *complicated* and *high-quality* trees, and also *transparent for explanation*, meaning that researchers can get fine-grained head-level MI estimation, and a bunch of intuitive probability functions, without worrying about *did my network furtively learnt the task?* Through comparing with a series of competitive baselines, we ensured its effectiveness, and then made two informative conclusions based on our estimated MI and reconstructed trees. The number of conclusions is limited due to content limit, and since our method is providing an analytical backbone, we strongly appeal to future research for fine-grained analysis on those estimated MI values and distributions.

Limitations

Despite its efficiency, our method still has several shortages: The most important one is that, to prevent the problem from being too complicated and bounded by the curse of dimensionality, our method does not consider the multivariate case, taking an assumption that all attention heads are independent. Moreover, as mentioned in Section 4.2, the introduction of "useless" attention scores is also noticeable, meaning that the density estimations might contain noises. Lastly, our method is only applicable to discrete-continuous mixtures, where all probed concepts are discrete labels, but not applicable to multivariate continuous joint distributions.

Ethical Considerations

Since our method is an explanation method, readers might exploit our method to perform syntactical attacks, like getting poorly captured dependency labels and designing specific prompts to confuse models. For models that are put into use in production environments, this might cause unexpectable effects.

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A Implementation Details of IPBP

In this section, we'll briefly introduce the implementation details, like the hyperparameters and key algorithms we use to implement IPBP.

As shown by the source code, we use the PyTorch framework to implement the whole IPBP process. We're not relying on off-the-shelf packages that have KDE functionalities like SciPy and Scikit-Learn, since their KDE implementation is CPU-based and thus too inefficient under our experiment settings.

Specifically, we take samples in $\mathcal{A}_{b,h;l}$ as a whole long tensor $\boldsymbol{a}_{b,h,l} \in \mathbb{R}^{|\mathcal{A}_{b,h;l}|}$. We calculate the minimum and maximum values of $\mathcal{A}_{b,h;l}$, and build a tensor of real numbers $\boldsymbol{X} = \{x_1, x_2, \ldots x_{n_x}\}$, ensuring that $x_1 < \min \mathcal{A}_{b,h;l}$, $x_{n_x} > \max \mathcal{A}_{b,h;l}$, and $x_1 < x_2 < \cdots < x_{n_x}$. These discrete x values serve as the points to calculate densities. Next, we calculate the mutual differences between each point in $\boldsymbol{X}$ and each element in $\boldsymbol{a}_{b,h,l}$, by repeating $\boldsymbol{X}^T$ for n_x times, getting a matrix $\overbrace{[\boldsymbol{X}^T, \ldots \boldsymbol{X}^T]}^{|\mathcal{A}_{b,h;l}| \text{ times}}$ of shape $n_x \times |\mathcal{A}_{b,h;l}|$, and repeating $\boldsymbol{a}_{b,h,l}$ for n_x times, also getting a matrix $\overbrace{[\boldsymbol{a}_{b,h,l}^T, \ldots \boldsymbol{a}_{b,h,l}^T]}^{n_x \text{ times}}{}^T$ of shape $n_x \times |\mathcal{A}_{b,h;l}|$. The absolute differences of the two matrices $|[\boldsymbol{X}^T, \ldots \boldsymbol{X}^T] - [\boldsymbol{a}_{b,h,l}^T, \ldots \boldsymbol{a}_{b,h,l}^T]|$, are the mutual differences, let's say $D(\boldsymbol{X}^T, \boldsymbol{a}_{b,h,l}) \in \mathbb{R}^{n_x \times |\mathcal{A}_{b,h;l}|}$. Then we calculate the standard deviation of $\mathcal{A}_{b,h;l}$, i.e., $\sigma_{\mathcal{A}_{b,h;l}}$, and take a rule-of-thumb value $\left(\frac{1}{\sum_{i=1}^{|\mathcal{A}_{b,h;l}|} w_i^2}\right)^{-\frac{1}{5}}$ for the bandwidth B, with all weight w_is equal to 1. We then calculate element-wise, following the following equation:

$$\frac{1}{B\sqrt{2\pi \cdot \sigma_{\mathcal{A}_{b,h;l}}}} \exp\left\{-\left(\frac{D(\boldsymbol{X}^T, \boldsymbol{a}_{b,h,l})}{B}\right)^2\right\} \quad (13)$$

In order to get the kernel values, which are also in shape $n_x \times |\mathcal{A}_{b,h;l}|$, we then calculate the row-wise mean of the kernel values to get the final kernel density values in shape n_x. Since all operations of this process are element-wise matrix operations, this is easily parallel-optimizable by PyTorch. As a result, the computation time for extracting attention score sets ($\mathcal{A}_{\cdot,\cdot;\cdot}$) and performing all kernel density estimations is within 1 hour using a single RTX 4090 GPU. Since attention score allocating and distribution estimations are only required to be done once, our method is extremely time-saving compared to most of the supervised probing methods.

For inferring on estimated probabilities (like inferring on posteriors $\hat{f}(l|A_{b,h})$ in Section 3.5, we take the estimated posteriors as a set of n_x discrete points, and an attention score in range $[x_i, x_{i+1}]$ will get its corresponding posterior value by interpolating between x_i and x_{i+1}. The interpolation-based inferring, together with other processes mentioned in Section 3.5, like head selection and score-weighted averaging, are all parallel-optimized, resulting in being able to run inference within 5 minutes on all baseline settings on a 4090 GPU.

For calculating integrals, specifically, the MI values like $\mathrm{MI}_{\mathrm{binary}}$, $\mathrm{MI}_{\mathrm{pos}}$, we use the trapezoid method to estimate the integral value: as mentioned in the section before, the kernel densities are described by n_x points, we take the n_x points as the integral limits and for every interval between x_i and x_{i+1}, we calculate the trapezoid areas and add them up to get the integral values.

During tree reconstruction, we empirically set the total number of heads, i.e., $\sum_{l \in \mathcal{L} \cup \{\phi\}} |\mathcal{H}_l|$ to a fixed value (2000), and use binary search.

We're using Universal Dependencies 2.9 (Zeman et al., 2021) as our dataset. That dataset is publicly available, using CC BY-SA 4.0 license[2] allowing free redistributions upon notifications. The Universal Dependencies (UD) dataset is designed to provide a standardized framework of grammatical identifications for NLP researchers, so we're following its intended usage.

B Implementation Details of Baselines

For the $\mathcal{V}$-Information MLPs, we found that training on all datasets will result in a network always predicting ϕ for all possible attention scores, due to the long-tail essential discussed in Section 4.3. This will result in many infinite $\mathcal{V}$-Information values, since there will be many estimated probabilities (for label in $\mathcal{L}$ other than ϕ) rather close to zero. Therefore, we apply a sample balancing technique, truncating $\mathcal{A}_{\cdot,\cdot;\phi}$ to make their numbers of samples

[2] https://creativecommons.org/licenses/by-nc-sa/4.0/deed.en

the same as the total number of samples in other score sets $\mathcal{A}_{\cdot,\cdot;1}, \mathcal{A}_{\cdot,\cdot;2}, \ldots \mathcal{A}_{\cdot,\cdot;|\mathcal{L}|}$. What's more, we also did a search on several network sizes, and found that if MLP$(\cdot)$ is $W_2(\text{act}(W_1(\cdot)))$, where W_1 in shape 1×2 and a W_2 in shape 2×4 achieves better fitting. This aligns with (Pimentel et al., 2020a) to some extent. We also use PyTorch to implement the baselines. Specifically, for ElasticNet that requires additional training, we use AdamW optimizer, $1e-5$ for both λ_1 and λ_2, and use a constant learning rate of $1e-3$, training for 12 epochs. For the $\mathcal{V}$-Information MLP, since we need $\mathcal{b}\mathcal{h} \times (|\mathcal{L}|+1)$ individual networks for predicting the alternatives of binary MI, we initialize $\mathcal{b}\mathcal{h} \times (|\mathcal{L}|+1)$ sets of matrices, each constituting the weights of a specific network $W_1, W_2, \ldots W_{\text{n\_layers}}$, with W_1 having a dimension of 1 and $W_{\text{n\_layers}}$ having a dimension of $|\mathcal{L}|+1$. During training and inferencing, we concatenate all attention scores $a_{b,h}^{[i][j]}$ for any $b \in \{1 \ldots \mathcal{b}\}$ and $h \in \{1 \ldots \mathcal{h}\}$ into a tensor of shape $\mathcal{b}\mathcal{h}$, and use `torch.bmm` to map each element of that tensor to $\mathcal{b}\mathcal{h} \times (|\mathcal{L}|+1)$ probabilities (standing for the probabilities of each label conditioned on each attention head's attention score, estimated by the variational family). Using `torch.bmm` will avoid training $\mathcal{b}\mathcal{h} \times (|\mathcal{L}|+1)$ networks separately, which is a disaster on computation loads, and can exploit GPU's parallel processing abilities. We use leaky_relu between hidden layers and use sigmoid to form the final probabilities. We use $1e-2$ as learning rate with exponential decay (0.8 at each epoch), together with an additional warmup epoch at the beginning. The hyperparameters differ for $\mathcal{V}$-Information since otherwise the variational family network will be more poorly trained. We also trained for 12 epochs.