Global Sensitivity Analysis of MAP inference in Selective Sum-Product Networks

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1 Introduction

Sum-Product Networks (SPN) are deep probabilistic models that have exhibited state-of-the-art performance in several machine learning tasks [1, 2, 3, 4, 5, 6, 7, 8]. As with many other probabilistic models, performing Maximum-A-Posteriori (MAP) inference is NP-hard in SPNs [9, 10]. A notable exception is selective SPNs [11, 9], that allows MAP inference in linear time. Due to the high number of parameters, SPNs learned from data can produce unreliable and overconfident inference. This effect can be partially detected by performing a Sensitivity Analysis (SA) of the model predictions to changes in the parameters [12]. In this work, we develop efficient algorithms for global quantitative analysis of MAP inference in selective SPNs. In particular, we devise a polynomial-time procedure to decide whether a given MAP configuration is *robust* with respect to changes in the model parameters. Experiments with real-world datasets show that this approach can discriminate easy- and hard-to-classify instances, often more accurately than criteria based on the probabilities induced by the model.

2 MAP Inference in Sum-Product Networks

An SPN S is a rooted weighted acyclic directed graph with indicator variables as leaves, and sum and product as internal nodes. We assume that SPN are *complete*, *decomposable* and *normalized* [1]. To ensure linear-time MAP inference, we also assume that SPNs are *selective* [11] (i.e., that at most one child sub-network of any sum node evaluates to nonzero at any realization). For example, the evaluation of the selective SPN in Figure 1(a) at $X_0 = 0$, $X_1 = 0$ and $X_2 = 1$ is $S(0, 0, 1) = 0.4(0 \times 0.3 \times 0.6) + 0.6(0.8 \times 0.9 \times 1) = 0.432$.



Figure 1: (a) Selective SPN. (b) CSPNs obtained by 0.1-contamination of the SPN in (a).

An SPN S induces a probability measure \mathbb{P}_{s} over the domain of its scope. Thus, given n SPN S with scope **X**, **E** and *evidence* $\mathbf{e} \in val(\mathbf{E})$, we define the set of **Maximum-A-Posteriori** (MAP) inference instantiations as:

$$\mathbf{x}^* \in \arg\max_{\mathbf{x} \in val(\mathbf{X})} \mathbb{P}(\mathbf{x}|\mathbf{e}) = \arg\max_{\mathbf{x} \in val(\mathbf{X})} \mathbf{S}(\mathbf{x}, \mathbf{e}) \,. \tag{1}$$

The problem is solvable in linear in the size of the network for selective SPNs by the simple Max-Product algorithm, that consists in replacing sum operations with maximizations and then evaluating the corresponding Max-Product network [11, 9]. The corresponding MAP instantiation is obtained by backtracking the solutions of the maximizations from the root toward the leaves. The highlighted subnetwork in Figure 1(a) contains the arcs selected by Max-Product.

3 Global Sensitivity Analysis of MAP Inferences in SPNs

In order to investigate the sensitivity of an inference w.r.t.] perturbations in the model parameters, we consider a set of SPNs { $S_w : w \in C$ }, where each S_w is parameterized by weights w and share the same network structure. We also assume that C is given by the Cartesian product of sets C_i , one for each sum node *i*. Two common approaches to obtaining C_i is ϵ -contamination:

$$\mathcal{C}_i = \left\{ (1 - \epsilon) \mathbf{w}_i + \epsilon \mathbf{v} : v_j \ge 0, \sum_j v_j = 1 \right\},\tag{2}$$

where $\epsilon \in (0, 1)$, \mathbf{w}_i are the weights associated to sum node *i* and *j* ranges over the children of *i*, and the **Imprecise Dirichlet Model** (IDM) [13]:

$$C_{i} = \{ \mathbf{w}_{i} : w_{ij} = \frac{N_{ij} + s \cdot v_{i}}{s + \sum_{j} N_{ij}}, v_{j} \ge 0, \sum_{j} v_{j} = 1 \},$$
(3)

where N_{ij} denote the number of times that the edge $i \rightarrow j$ is nonzero over the dataset, and s is the prior strength s of a Multinomial-Dirichlet model.

In this work, we perform global SA by verifying whether a MAP inference \mathbf{x}^* is the single maximizer of each SPN in a set $\{S_{\mathbf{w}} : \mathbf{w} \in C\}$ given evidence \mathbf{e} , that is, whether:

$$\max_{\mathbf{x} \neq \mathbf{x}^*} \max_{\mathbf{w} \in \mathcal{C}} \left(\frac{\mathbf{S}_{\mathbf{w}}(\mathbf{x}, \mathbf{e})}{\mathbf{S}_{\mathbf{w}}(\mathbf{x}^*, \mathbf{e})} \right) < 1.$$
(4)

The following algorithm computes the left-hand side of the inequality below:

$$V^{i} = \begin{cases} 1 & \text{if } i \text{ is consistent,} \\ 0 & \text{if } i \text{ is inconsistent,} \\ \prod_{j} V^{j} & \text{if } i \text{ is a product node,} \\ \max\left\{\max_{j \in ch(i), j \neq k} \left(\max_{\mathbf{w}_{i} \in \mathcal{C}_{i}} \frac{w_{ij} \max_{\mathbf{x}} \mathbf{S}_{\mathbf{w}_{k}}^{j}(\mathbf{x}, \mathbf{e})}{w_{ik} \mathbf{S}_{\mathbf{w}_{k}}^{k}(\mathbf{x}^{*}, \mathbf{e})}\right), V^{k} \right\} \text{ if } i \text{ is a sum node,} \end{cases}$$
(5)

where k is the active child of i for S at MAP configuration and the evidence $(\mathbf{x}^*, \mathbf{e})$.

4 Experiments and Results

We perform SA of MAP inferences on 8 benchmark multilabel classification domains. We assess the ability to distinguish between reliable and unreliable classifications by measuring both exact match (i.e., whether all labels are correct) and accuracy (which rewards relevant predictions while discounting for irrelevant predictions), on instances deemed robust or non-robust by each method. We compare our approach against a baseline procedure that considers a classification robust if the difference between the most probable and the second most probable configurations exceeds a given threshold p. We select the best values $\epsilon *$, s * and p * for resp. ϵ -contamination, IDM and difference of probability criteria using a validation dataset, an then perform a SA for each multilabel prediction in the test dataset. The results are in the table below.

During		Accuracy			Exact Match			During		Accuracy			Exact Match		
Dataset		Robust	$\neg Robust$	ΔAcc	Robust	$\neg Robust$	ΔEM	Dataset		Robust	$\neg Robust$	ΔAcc	Robust	$\neg Robust$	ΔEM
Arts	$\epsilon *$	0.88	0.196	0.634	0.833	0.143	0.69	Health	$\epsilon *$	0.667	0.557	0.11	0.5	0.416	0.084
	s*	0.107	0.351	-0.244	0.089	0.247	-0.158		s*	0.637	0.482	0.155	0.537	0.304	0.233
	p*	0.81	0.159	0.651	0.75	0.107	0.643		p*	0.655	0.552	0.103	0.552	0.409	0.143
Business	$\epsilon *$	0.751	0.582	0.169	0.617	0.598	0.019	Human	$\epsilon *$	0.203	-	-	0.146	-	-
	s*	0.781	0.641	0.14	0.642	0.469	0.173		s*	0.203	-	-	0.146	-	-
	p*	0.762	0.581	0.181	0.62	0.392	0.228		p*	0.211	0.198	0.013	0.155	0.14	0.015
Emotions	$\epsilon *$	0.595	0.41	0.185	0.238	0.163	0.075	Plant	$\epsilon *$	0.331	0.217	0.114	0.324	0.213	0.111
	s*	0.686	0.413	0.273	0.308	0.16	0.148		s*	0.367	0.162	0.205	0.362	0.157	0.205
	p*	0.574	0.391	0.183	0.176	0.176	0		p*	0.345	0.212	0.132	0.338	0.208	0.13
Flags	$\epsilon *$	0.917	0.468	0.449	0.5	0.118	0.382	Scene	$\epsilon *$	0.857	0.277	0.58	0.857	0.212	0.645
	s*	0.917	0.468	0.449	0.5	0.1	0.4		s*	0.929	0.293	0.636	0.929	0.293	0.636
	p*	0.917	0.468	0.449	0.5	0.118	0.382		p*	0.923	0.276	0.647	0.923	0.211	0.712

One sees that the SA obtained with IDM achieve the highest accuracy and exact math in 5 of the 8 domains, followed by the ϵ -contamination (3 out of 8). Probability difference outperforms the other methods w.r.t. accuracy only for Human, where the our methods consider all instances robust. Yet the difference in accuracy (or exact match scores) for robust and non-robust is very small, showing that the two sets perform indeed very similar (but perhaps it would be more sensible to classify all instances as non-robust). The IDM-based SA perform particularly poorly in the Arts domain, where the accuracy of the robust portion is significantly inferior to the accuracy in the non-robust portion.

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