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Project 1

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Abstract

This paper describes the mixtures-of-trees model, a probabilistic model for discrete multidimensional domains. Mixtures-of-trees generalize the probabilistic trees of [1] in a different and complementary direction to that of Bayesian networks. We present efficient algorithms for learning mixtures-of-trees models in maximum likelihood and Bayesian frameworks. We also discuss additional efficiencies that can be obtained when data are “sparse,” and we present data structures and algorithms that exploit such sparseness. Experimental results demonstrate the performance of the model for both density estimation and classification. We also discuss the sense in which tree-based classifiers perform an implicit form of feature selection, and demonstrate a resulting insensitivity to irrelevant attributes.

Keywords: Bayesian Networks, Mixture Models, Chow-Liu Trees

1. Introduction

Probabilistic inference has become a core technology in AI, largely due to developments in graph-theoretic methods for the representation and manipulation of complex probability distributions [3]. Whether in their guise as [2] directed graphs (Bayesian networks) or as undirected graphs (Markov random fields), *probabilistic graphical models* have a number of virtues as representations of uncertainty and as inference engines. Graphical models allow a separation between qualitative, structural aspects of uncertain knowledge and the quantitative, parametric aspects of uncertainty...

Remainder omitted in this sample. See <http://www.jmlr.org/papers/> for full paper.

2. Ordinary Least Squares

2.1 Discussion on Scaling

Unscaled sample design matrix fitting one-dimensional polynomial of degree 5

```
[[1.      0.      0.      0.      0.      0.      ]
 [1.      0.25    0.0625  0.01562 0.00391 0.00098]
 [1.      0.5     0.25    0.125   0.0625  0.03125]
 [1.      0.75    0.5625  0.42188 0.31641 0.2373 ]
 [1.      1.      1.      1.      1.      1.      ]]
```

Scaled sample design matrix fitting one-dimensional polynomial of degree 5

```
[[ 0.      -1.41421 -1.0171  -0.83189 -0.728  -0.66226]
 [ 0.      -0.70711 -0.84758 -0.7903  -0.71772 -0.65971]
 [ 0.      0.      -0.33903 -0.49913 -0.56348 -0.58075]
 [ 0.      0.70711  0.50855  0.29116  0.10488 -0.0433 ]
 [ 0.      1.41421  1.69516  1.83016  1.90431  1.94603]]
```

Unscaled β after fitting two-dimensional polynomial of degree 5

```
[ 0.4097   7.55124   3.79304 -32.85696 -14.83669 -8.81645 45.45889
 43.33221 20.70625 -7.63623 -21.24552 -51.81866 -7.53731 -29.60175
 28.57282  0.73824 18.29253 10.60883 -5.52465 16.60259 -16.13743]
```

Scaled β after fitting two-dimensional polynomial of degree 5

```
[ 2.21481  1.1093  -9.93941 -3.32124 -2.6612  13.11848  8.70306  4.14949
-2.19979 -5.79259 -9.48697 -1.28166 -5.40093  7.77825  0.19055  3.09519
1.60557 -0.83485  2.797   -4.1589 ]
```

First x_{unscaled} and x_{scaled} is not that different, the original data was close to zero-centered and not that spread out, which means that initially by just looking at the data scaling is not that necessary. When we add the polynomial terms we can now see that some of the entries of X_{unscaled} get really small as an example $0.1^5 = 0.00001$ this makes the columns of X_{unscaled} live in their own order of magnitude and scaling should be considered to bring them back to the same ish order of magnitude. The act of not scaling results in β spanning from -50 to 48 while scaling gives a smaller span from -10 to 13 (We would imagine that keeping everything in the same order of magnitude is something that the computer likes, perhaps reducing floatingpoint error). Now this alone may not justify why we should scale this dataset, as scaled and unscaled OLS yields the same MSE. However when doing ridge regression the cost function is directly dependent on the magnitude of β_i . This explains why the MSE for unscaled data is much higher than for scaled data in the ridge case. This leads us to conclude that we should scale the data, making it easier to tweak λ and giving us nicer numbers to work with.

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Appendix A.

In this appendix we prove the following theorem from Section 6.2:

Theorem *Let u, v, w be discrete variables such that v, w do not co-occur with u (i.e., $u \neq 0 \Rightarrow v = w = 0$ in a given dataset \mathcal{D}). Let N_{v0}, N_{w0} be the number of data points for which $v = 0, w = 0$ respectively, and let I_{uv}, I_{uw} be the respective empirical mutual information values based on the sample \mathcal{D} . Then*

$$N_{v0} > N_{w0} \Rightarrow I_{uv} \leq I_{uw}$$

with equality only if u is identically 0. ■

Proof. We use the notation:

$$P_v(i) = \frac{N_v^i}{N}, \quad i \neq 0; \quad P_{v0} \equiv P_v(0) = 1 - \sum_{i \neq 0} P_v(i).$$

These values represent the (empirical) probabilities of v taking value $i \neq 0$ and 0 respectively. Entropies will be denoted by H . We aim to show that $\frac{\partial I_{uv}}{\partial P_{v0}} < 0 \dots$

Remainder omitted in this sample. See <http://www.jmlr.org/papers/> for full paper.

Bibliography

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