

Doctor of Philosophy

Diversified Probabilistic Graphical Models

by

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CERTIFICATE OF AUTHORSHIP/ORIGINALITY

I certify that the work in this thesis has not previously been submitted for a degree nor has it been submitted as part of requirements for a degree except as fully acknowledged within the text.

I also certify that the thesis has been written by me. Any help that I have received in my research work and the preparation of the thesis itself has been acknowledged. In addition, I certify that all information sources and literature used are indicated in the thesis.

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ABSTRACT

Probabilistic graphical models (PGMs) as diverse as Bayesian networks and Markov random fields have provided a fundamental framework to learn and reason using limited and noisy observations. Examples include, but are not limited to, hidden Markov models (HMMs), sequential graphical models, and probabilistic principal component analysis mixture models (PPCA-MM). PGMs have been used in a wide variety of applications such as speech recognition, natural language processing, web searching, and image understanding. However, one potential drawback of using PGMs with traditional learning and inference methods is that the learned parameters or inferred variables are easily trapped within local, clustered optima rather than distributed evenly across the whole space. Taking mixture models as an example, the learned mixing components might overlap. Consequently, the resulting models might show ambiguity when clustering is performed based on these overlapping mixing components. This phenomenon might limit PGM performance.

Although efforts have been made to explore a variety of priors to alleviate this potential drawback and to enhance PGM performance, diverse priors have yet to be fully explored and utilized. Diversity is a concept that encourages counterpart model parameters and variables to repel as much as possible and, in doing so, spread out model components and decrease overlapping. However, how to explicitly encode these priors into a PGM and how to solve the resulting diversified PGMs are two critical problems that must be solved. This thesis proposes a unified framework to constrain PGMs with diverse priors. Three different PGMs -HMMs, time-varying determinantal point processes (TV-DPPs), and PCA-MMs - are elaborated to demonstrate the proposed diversified PGM framework. For each PGM, three basic constituent framework elements are examined: which part of the traditional PGM is diversified, how to formulate the diversity, and how to solve the diversified version, e.g., parameter learning and inference. In addition, experiments are conducted using various application scenarios to verify the effectiveness of the proposed diversified PGMs.

Keywords: Probabilistic graphical models (PGMs), diversity prior, determinantal point processes (DPPs), hidden Markov models (HMMs), time-varying DPPs, probabilistic principal component analysis (PPCA).

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- 3.2Illustration of Sequential Monte Carlo: At time stamp t - 1, 10 particles in red with equal weights are given, i.e. $\{x_{t-1}^{(i)}\}_{i=1}^{10}$. At this stage, two computations will be done - One is computing the incremental weights; the other is computing the particles for the next time stamp. For the incremental weight of each particle at time t-1, according to Eq. (3.7), it is simply the likelihood ratio between time stamps t and t-1. The corresponding relationship is denoted by the dashed line connecting two neighbour distributions and the weight for each particle is illustrated by size of blue solid circle. For the particle's location at next time stamp t, usually, a Markov transition kernel is used to qualify the transition job between two slightly different neighbour distributions. The transition relationship is indicated by solid line with arrow. For the particle's weight at time stamp t, it is gained by multiplying the weight at time stamp t-1 by the incremental weight. To alleviate the degeneracy of the algorithm which is measured by effective sample size (ESS), a re-sampling step is applied when $N_{ESS} < \alpha \cdot N$. High weighted particles will re-birth as several equal weighted particles, while particles with low weights may disappear. To increase the samples' diversity, a move step is followed. Once particle' locations and weights at t are prepared, it will recursively carry out the whole above procedure.

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