

# Active Inference for Dynamic Bayesian Networks

Caner Komurlu

Illinois Institute of Technology, Chicago Illinois  
ckomurlu@hawk.iit.edu

## Abstract

In supervised learning, many techniques focus on optimizing training phase to increase prediction performance. *Active inference*, a relatively novel paradigm, aims to decrease overall prediction error via selective collection of some labels based on relations among instances. In this research, we use dynamic Bayesian networks to model temporal systems and we apply active inference to dynamically choose variables for observation so as to improve prediction on unobserved variables.

## 1 Introduction

In supervised learning, a mathematical model is trained by tuning its parameters using labeled data in order to automatically predict labels for unseen data. Many studies have focused on training. For example, active learning tries to train a model using fewer labeled data by selecting most informative instances. This helps reducing labeling cost [Settles, 2012].

A relatively new approach, *active inference*, maximizes prediction performance by selective information gathering during prediction [Bilgic and Getoor, 2009]. In this approach, relations between instances are utilized with the intuition that knowing true label of some instances help predicting others.

Dynamic Bayesian network (DBN) is a generative statistical model which asserts probabilities of random variables accounting complex dependencies. Two main properties of DBNs make them powerful: i) factorizing joint probability distributions into conditional probability distributions, ii) dynamically representing random variables in time dimension.

In majority of cases, lack of evidence degrades prediction performance in DBNs over time. As random variables are correlated, observing one contributes to evaluating probabilities on its dependents. In some scenarios, observed variables are specified by definition. Otherwise, selecting variables to observe arises as a problem to tackle. Therefore, active inference can help to detect variables to observe and eventually it can increase prediction performance significantly. This objective revolves around assessment of prediction uncertainty and calculating observation cost. To the best of our knowledge, this is the first time active inference is applied on DBNs.

In the following, Section 2 presents description of these objectives on two practical problems along with results ob-

tained. Section 3 continues with short and long term research plans, followed by a conclusion in Section 4.

## 2 Preliminary Research and Results

In this section, our proposed method, *active inference for dynamic Bayesian networks*, will be described and evaluated on two practical problems: i) detecting optimal time for observation for tissue engineering, and ii) dynamic detection of optimal observation subset on wireless sensor networks.

### 2.1 Active Inference for Tissue Engineering

In tissue engineering domain, experts seek conditions for optimal tissue development. One criterion for optimal development is blood vessel network which should develop in tandem with tissue, also named as vascularization. Though many factors affect the performance of vascularization, few are known. Therefore, this phenomenon is partially observable, hence stochastic. Given an initial configuration, e.g initial blood vessel and tissue cell locations, stress level of tissue cells, we try to estimate probabilities of each atomic locations being occupied by blood vessel in a sequence of time stamps.

We modeled the environment as a grid, of which each cell represents an atomic location. We assumed that the direction of blood vessel progress is from bottom to top. We designed a DBN in which each random variable represents a location whose parents are lower neighbors from previous time. Hence occurrence of blood vessel in a location becomes more likely when parents have blood vessel [Komurlu *et al.*, 2014]. Given initial settings, i.e. each location's value at time slice  $t = 0$ , we compute probabilities at following time slices until the final time slice,  $T - 1$ . Next, we find most probable complete observation of each time slice. Then for each most probable observation, we compute uncertainty of predictions on the final time slice. The objective is to find the earliest time slice,  $t^*$ , on which observation for each location yields an uncertainty at time slice  $T$  less than a given threshold  $\sigma$ .

For three different stress levels of tissue cells, we computed uncertainty at each time slice which can be seen in Figure 1. Note that the uncertainty computation is expensive and we cannot merely generate this uncertainty curve for any given initial setting. Therefore, we tried some search methods to find  $t^*$  in the search space of uncertainty and we made analytical evaluation. The reference article is hidden as it is under revision of a journal.

