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Dynamic Staged Trees for Discrete Multivariate Time Series: Forecasting, Model Selection and Causal Analysis

Guy Freeman* and Jim Q. Smith†

Abstract. A new tree-based graphical model — the dynamic staged tree — is proposed for modelling discrete-valued discrete-time multivariate processes which are hypothesised to exhibit symmetries in how some intermediate situations might unfold. We define and implement a one-step-ahead prediction algorithm with the model using multi-process modelling and the power steady model that is robust to short-term variations in the data yet sensitive to underlying system changes. We demonstrate that the whole analysis can be performed in a conjugate way so that the potentially vast model space can be traversed quickly and then results communicated transparently. We also demonstrate how to analyse a general set of causal hypotheses on this model class. Our techniques are illustrated using a simple educational example.

Keywords: Staged trees, graphical models, Bayesian model selection, Dirichlet distribution, Bayes factors, forecasting, discrete time series, causal inference, power steady model, multi-process model, clustering

1 Introduction

In this paper we consider a class of dynamic multivariate graphical models applicable to a wide range of discrete-valued processes, including some from biology, medicine and education. We consider those that have the following characteristics:

1. A description is provided of the possible development histories each unit in the process can take for a given time. These histories could be radically different from one another in terms of length of development, the variables encountered, the state spaces of each stage of development, and so on.

2. There are various symmetry hypotheses for a given cohort of units concerning which intermediate situations in the histories have the same distributions over their immediate developments.

3. The units arrive in discrete time cohorts, assumed here for simplicity to be equally spaced apart. The symmetries in the system are allowed to change from one time point to the next to reflect a changing environment.

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4. The system may, at various times, be subject to local interventions. The model then admits a “causal” extension which provides predictions of the process when subject to such controls.

These types of discrete processes are clearly common but to our knowledge have so far not been systematically studied. We are particularly interested in this paper in making good one-step ahead predictions for such processes. This will also provide, as a beneficial side-effect, the probabilities of the symmetry hypotheses through time, which can be used as an explanatory tool.

One example of a system that exhibits the characteristics above is a programme of study provided by an educational establishment which monitors students’ marks over time. We therefore use this as our running example. (Many real-world systems share these characteristics, as has been discussed in the literature for biological (Smith and Anderson 2008) and medical diagnosis (Thwaites et al. 2009) processes). The system characteristics described above might be manifested in the educational setting as follows:

1. The modules of the course are always taken in a particular order (or consistent with some partial order); there might be a requirement to achieve a threshold mark before being allowed to continue onto the next module; and certain modules might have different prerequisite modules.

2. A student’s performance on a previous module could influence the marks on a later one.

3. New students come in yearly cohorts, and because of any number of possible changes in any number of unobserved confounding factors each cohort could exhibit different symmetries in their possible course mark histories.

4. The administrators will be interested in predicting the effect on the mark distribution by changing the program in some way, such as changing the syllabus or lecturer for a module, changing the prerequisites for a module or removing a module entirely.

A simple graphical representation of the different mark combinations a student in such a programme could achieve is the event tree (Shafer 1996), such as the one given by Figure I for a course with two modules. Event trees can represent any discrete event space and naturally codify a chronological order (or partial order) in their topology, and so we base our own proposed model class on them.

Each root-to-leaf path of the event tree represents a distinct combination of grades. In Figure I, for example, the top-most path represents the student achieving the top grade in both modules and those marks not being missing.

The nodes in the tree that are neither root nor leaf nodes indicate points along the process at which histories become differentiated. For example, the node $v_2$ in Figure I represents the point at which the grade histories with a non-missing first
Figure 1: Event tree for marks for two modules in a course. Marks are discretized into 3 grades, and $A$ and $NA$ indicate whether the mark is recorded or missing. The 10 situations are labelled and the 16 leaf nodes are unlabelled.

mark differentiate into those with a high grade, middling grade or low grade. We call these intermediate nodes situations.

However, the event tree’s semantics are not sufficient for addressing the rest of our requirements by itself, in particular because it does not codify the symmetries in the system that we are interested in modelling. These symmetries can be viewed as equivalent, in many cases, to different situations having the same subsequent development, whether only on the immediate possibilities or on the whole remaining process. In the educational example of Figure 1, for example, it might be considered to be the case that the grade distributions for the second module (where available) are the same no matter what the mark for the first module was. In that case, the corresponding situations $v_8$, $v_9$ and $v_{10}$ are exhibiting the symmetry we are interested in. We call such situations stages.

The symmetries discussed above can be conceptualised as a kind of conditional independence (Dawid 1979; Studen 2005) where the random variable describing the subsequent development for symmetrical situations is held to be independent of the history that led to those situations. There are many graphical models that aim to represent conditional independence relations between the different variables of a system. Bayesian networks (BNs) (Pearl 2000; Cowell et al. 2007) are currently the most prominent of
these models. However, they cannot easily represent in their graphical structure the asymmetry in the potential histories of a unit, such as those illustrated by the example of Figure [II] or symmetry among only a subset of a variable’s sample space. Some enhancements to the canonical Bayesian network have therefore been suggested in order to take in particular the latter consideration, called “context-specific independence”, into account — for example by [Boutilier et al.|1996] or [Geiger and Heckerman|1996] — but these approaches can be unwieldy or abandon any graphical representation of the symmetry.

Using different semantics from BNs, [Smith and Anderson|2008] defined the chain event graph (CEG) as an enhancement of the event tree, where non-leaf nodes in an event tree with the same probability distribution over their outgoing edges are linked by undirected edges, and where subtrees with identical probability distributions of their root-to-leaf paths are merged. Our model class is therefore based on CEGs, but extended into a more general dynamic scenario where probabilities are allowed to change with time.

In this paper we describe the dynamics of this type of tree-structured process by a state space model incorporating a switching mechanism to neighbouring models. The earliest example of this class, to the best of our knowledge, was studied for univariate Gaussian series by [Harrison and Stevens|1976] and called Multi-process Models Class II (see [West and Harrison|1997] for a more recent review). [Frühwirth-Schnatter|2006] reviews switching models for non-Gaussian state spaces, none of which have closed posterior forms. Here, we use a type of multi-process model which allows us to dynamically shift from one symmetry partition to neighbouring ones.

In order to take into account possible drifting on the tree parameters through time caused by unobserved background processes, one could follow the filtering approach of stating a transition probability $P(\theta_t \mid \theta_{t-1}, S)$, where $\theta_t$ represents the parameters on the tree at time $t$ and $S$ is the underlying model. The most common way to achieve this is to use a conventional state space formulation. Unfortunately, this approach immediately requires the inference to be undertaken with approximating numerical methods. This is not ideal in this context for several reasons: Firstly, in the stochastic version of the process we consider, conjugacy is retained by using product Dirichlet priors, and it would be a shame to lose this useful modular property. Secondly, because of the enormity of the model space of our domain of application it is convenient to be able to have Bayes factors calculable in closed form, because this greatly speeds up computation of model goodness. Thirdly, models in this class are easier to interpret when they retain their modular and conjugate forms.

Another approach, which we take here, is to set a transition function

$$T : P(\theta_{t-1} \mid x^{t-1}, S) \mapsto P(\theta_t \mid x^{t-1}, S)$$  \hspace{1cm} (1)$$

where $x^{t-1}$ are the observations up to time $t - 1$. Although this approach is more restrictive in its scope, it can be justified through various characterisations ([Smith|1979], [1992]) and we show below that it can have the very attractive property of preserving the conjugate structure of each model in this class, encouraging several different authors to
use such transitions.

Interventions on a graphical model are covered by the causal literature (e.g. Pearl (2000)). Causal analysis on event trees was considered by Shafer (1996) and was defined for chain event graphs by Thwaites et al. (2010). We implement causal interventions on the dynamic model class we present here. By retaining conjugacy and modularity when learning model probability parameters, this causal extension of the model class is particularly straightforward, allowing us to utilise it for modelling a controlled environment.

We proceed to developing our new model, the dynamic staged tree. In Section 2 we formally define the necessary concepts. In Section 3 we develop a multi-process model for the dynamic staged tree that can be used to make one-step ahead predictions. In Section 4 we extend the multi-process model to causal analyses on the dynamic staged tree. We end in Section 5 by applying our analyses to study the results from part of a real educational programme by inferring the changing probabilities of symmetry hypotheses on-line.

2 Concepts and definitions

2.1 Event tree

We begin with some definitions. See Smith and Anderson (2008), Thwaites et al. (2010) and Freeman and Smith (2011) for more details and discussion about these concepts.

Let $T = (V(T), E(T))$ be a directed tree where $V(T)$ is its node set and $E(T)$ its edge set. Let $L(T)$ be the set of leaf nodes and $S(T) = \{v : v \in V(T) \setminus L(T)\}$ be the set of situations of $T$. Let $\lambda(v, v')$ be the path from node $v \in S(T)$ to node $v' \in V(T)$ (if it exists), and let $\Lambda(v, T) = \{\lambda(v, v') : v' \in L(T)\}$, the set of paths from $v$ to a leaf node. Let $X = \Lambda(v_0, T)$, where $v_0$ is the root node of $T$, so that $X$ is the set of root-to-leaf paths of $T$. Each path $X \in X$ is an atomic event, corresponding to a possible unfolding of events through time by using the partial ordering induced by the paths.

Let $X(v)$ denote the set of children of $v \in S(T)$. In an event tree (ET), each situation $v \in S(T)$ has an associated random variable $X(v)$ with sample space $X(v)$, defined conditional on having reached $v$. The distribution of $X(v)$ is determined by the primitive probabilities $\theta(v) = \{\theta(v, v') = p(X(v) = v') : v' \in X(v)\}$.

With random variables on the same path being mutually independent, the joint probability of events on a path can be calculated by multiplying the appropriate primitive probabilities together. Each primitive probability $\theta(v, v')$ is a colour for the directed edge $e = (v, v')$, so that we let $\pi(e) := \theta(v, v')$. 

2.2 Staged trees

Starting with an event tree $T$, define a floret of $v \in S(T)$ as

$$F(v) = (V(F(v)), E(F(v)))$$

where $V(F(v)) = v \cup \mathbb{X}(v)$ and $E(F(v)) = \{e \in E(T) : e = (v, v') : v' \in \mathbb{X}(v)\}$. The floret of a vertex $v$ is thus a sub-tree consisting of $v$, its children, and the edges connecting $v$ and its children, as shown in Figure 2. This represents, as defined in section 2.1, the random variable $X(v)$ and its sample space $\mathbb{X}(v)$.

Two situations $v, v' \in S(T)$ are said to be in the same stage $u$ if and only if $X(v)$ and $X(v')$ have the same distribution under a bijection

$$\psi_u(v, v') : \mathbb{X}(v) \rightarrow \mathbb{X}(v')$$

It follows that one necessary condition for $v$ and $v'$ to be in the same stage is that $|\mathbb{X}(v)| = |\mathbb{X}(v')|$, i.e. $v$ and $v'$ have the same number of children. In particular, $\psi_u(v, v)$ is the identity function for any stage $u$ that contains $v$, including $u = \{v\}$.

The set of stages (or staging) of $T$ is written $U(T)$. It is clear that $U(T)$ is a partition of $S(T)$. The set $S(T)$ itself can be thought of as the trivial staging.

Finally, a staged tree $ST(T, U(T))$ is constructed from $T$ by letting $V(ST) = V(T)$ and $E(ST) = E_d(ST) \cup E_u(ST)$, where $E_d(ST)$ and $E_u(ST)$ are constructed as follows:

- $E_d(ST)$ is identical to $E(T)$ except that two edges $(v, v^*)$, $(v', v'^*)$ are given the same colour if and only if $v^* \mapsto v'^*$ under some $\psi_u(v, v')$ as defined above;
- $E_u(ST)$: for every $v, v' \in S(T)$, an undirected edge between $v$ and $v'$ is drawn if and only if $X(v)$ and $X(v')$ have the same distribution.

It is easily shown (Smith and Anderson 2008) that BNs over finite discrete random variables are an important but small subclass of staged trees.
3 Prediction with dynamic staged trees

Let $T$ be an event tree whose topology is known and fixed in time, but with an uncertain and possibly dynamic probability distribution over its structure. Let the set of situations of $T$, $S(T)$, be denoted by $S = \{v_1, \ldots, v_{|S|}\}$.

At each time point $t = 1, \ldots, \tau$, we wish to predict $x_t(v, v')$ for all $v' \in \mathcal{X}(v)$ for all $v \in S$, where $x_t(v, v')$ is the number of times $X(v) = v'$ at time $t$. Let $\mathbf{x}_t = (x_t(v))_{v \in S}$ where $x_t(v) = (x_t(v, v'))_{v' \in \mathcal{X}(v)}$. Then at every time $t$ we need to construct a probability distribution over the possible values of $\mathbf{x}_t$ conditional on all previous observations $\mathbf{x}^{t-1} = (x_1, \ldots, x_{t-1})$. The marginal joint distribution $P(\mathbf{x}^t)$ over time of the full data set can be represented as a product of the one-step ahead predictive probabilities $P(\mathbf{x}_t \mid \mathbf{x}^{t-1})$. Bayes factors associated with different models can then be expressed as a function of these quantities. It is interesting to note that this factorisation corresponds to the prequential likelihood described by [Dawid, 1984] used for comparing probabilistic forecasting systems.

The probability distribution of $\mathbf{x}_t \mid \mathbf{x}^{t-1}$ can be written parametrically as a function of $\theta_t$, the values of $\theta(v)$ for all $v \in S$ at time $t$, so that

$$P(\mathbf{x}_t \mid \mathbf{x}^{t-1}) = \int_{\Theta_t} P(\mathbf{x}_t \mid \theta_t, \mathbf{x}^{t-1}) P(\theta_t \mid \mathbf{x}^{t-1}) d\theta_t. \quad (4)$$

$\theta_t$ is unknown in the general case. One way to specify the distribution of $\theta_t$ is to assume the process can be described by a dynamic staged tree. We define a dynamic staged tree to be an event tree where at each time point $t = 1, \ldots, \tau$ (where $\tau$ can be finite or infinite) an independent sampling over $\mathcal{X}$ occurs but with a possibly different staging $U_t(T)$.

If $v, v' \in S(T)$ are in the same stage $u$ in a partition $U$ at time $t$ then we assume that

$$\theta_t(v) = \theta_t(v') \triangleq \theta_t(u). \quad (5)$$

With these assumptions, the distribution of $\theta_t$ under a staging $U_t$ can be written as the product of the distribution of each stage’s parameters:

$$P(\theta_t \mid U_t, \mathbf{x}^{t-1}) = \prod_{u \in U_t} P(\theta_t(u) \mid U_t, \mathbf{x}^{t-1}). \quad (6)$$

Therefore equation (4) can be written as

$$P(\mathbf{x}_t \mid \mathbf{x}^{t-1}) = \sum_{U_t \in \mathcal{U}} \int_{\Theta_t} P(\mathbf{x}_t \mid \theta_t, U_t, \mathbf{x}^{t-1}) P(\theta_t \mid U_t, \mathbf{x}^{t-1}) P(U_t \mid \mathbf{x}^{t-1}) d\theta_t$$

$$= \sum_{U_t \in \mathcal{U}} \int_{\Theta_t} \left( P(\mathbf{x}_t \mid \theta_t, U_t, \mathbf{x}^{t-1}) P(U_t \mid \mathbf{x}^{t-1}) \prod_{u \in U_t} P(\theta_t(u) \mid U_t, \mathbf{x}^{t-1}) \right) d\theta_t. \quad (7)$$

So to carry out a one-step ahead forecast on the system three probability distributions must be specified: the sampling distribution $P(\mathbf{x}_t \mid \mathbf{x}^{t-1}, \theta_t, U_t)$, the stage
parameter distributions \(P(\theta_t(u) \mid U_t, x^{t-1})\), and the staging distributions \(P(U_t \mid x^{t-1})\). We show below how this can be achieved for each term in turn.

### 3.1 The sampling distributions

Under complete sampling the distribution of \(X(v)\) for any situation \(v \in S\) is conditionally independent of any other quantity given \(\theta(v)\). In particular, this means that the distributions of \(X(v)\) and \(X(v')\) for two situations \(v, v' \in S, v \neq v'\), are assumed to be independent conditional on \(\theta(v), \theta(v')\).

This does not necessarily apply to \(x_t(v)\), because the distribution of the number of samples \(N_t(v)\) from \(X(v)\) at time \(t\) is unknown in the general case. We assume here, however, that for all situations \(v\) bar the root node \(v_0\) that \(N_t(v)\) equals the value of \(x_t(v^*, v)\), where \(v^*\) is the situation such that \(v \in \mathcal{X}(v^*)\), i.e. where \(v^*\) is the parent node of \(v\). We discuss the setting of \(N_t(v_0)\) shortly.

Assuming that \(x_t \perp \perp x^{t-1} \mid \theta_t\), a standard state space model assumption (where \(\perp \perp\) signifies independence), and that the components of \(x_t(v)\) for each \(v\) are independently sampled conditional on \(\theta_t(v)\), we can therefore write

\[
P(x_t \mid \theta_t, U_t, x^{t-1}) = \sum_{N_t(v_0)} P(x_t \mid N_t(v_0), \theta_t, U_t, x^{t-1}) P(N_t(v_0) \mid \theta_t, U_t, x^{t-1})
\]

where \(I_A\) is the indicator variable for an event \(A\) and \(x_t(v^*, v_0)\) is understood to mean \(N_t(v_0)\).

The modelling of the distribution of \(N_t(v_0)\) depends on the details of the system under consideration. One common scenario is when \(N_t(v_0)\) is believed to be independent of all other system parameters apart from, at most, values of \(N_s(v_0)\) for \(s < t\). One approach in this case is to model \(N_t(v_0)\) as a Poisson variable with parameter \(\lambda\), where \(\lambda\) can either be constant or itself given a conjugate prior of Gamma\((\alpha_\lambda, \beta_\lambda)\) at time 1.

When \(N_t(v_0)\) is known, equation (11) becomes

\[
P(x_t \mid \theta_t, U_t, x^{t-1}) = \prod_{v \in S} \left[ I_{\{\sum x_t(v,v')=x_t(v^*,v)\}} \prod_{v' \in \mathcal{X}(v)} \theta_t(v, v')^{x_t(v,v')} \right]
\]

where \(x_t(v^*, v_0)\) should again be read as \(N_t(v_0)\).
3.2 The stage parameter distributions

As with every aspect of the model, the specification of the probability distribution over the floret parameters for each possible stage should be tailored to the scenario at hand. In many cases, however, it is possible to characterise the distribution from some common qualitative modelling assumptions.

Consider first the trivial staging \( U_t = S \). It is shown in Freeman and Smith (2011) that if it is assumed that the relative rates of the root-to-leaf paths are independent, then the additional assumption of mutual independence of the floret distributions implies that each non-trivial floret’s distribution must be Dirichlet. Therefore, denoting its set of hyperparameters as \( \alpha_t(v) = (\alpha_t(v, v'))_{v' \in \mathcal{X}(v)} \), the density of \( \theta_t(v) \mid U_t = S, x^{t-1} \) for a non-trivial floret \( v \in S \), where \( \theta_t(v, v') \) is the value of \( \theta(v, v') \) at time \( t \), is

\[
f_{\theta_t(v)}(\theta_t(v) \mid U_t = S, x^{t-1}) = \Gamma \left( \sum_{v' \in \mathcal{X}(v)} \alpha_t(v, v') \right) \prod_{v' \in \mathcal{X}(v)} \frac{\theta_t(v, v')^{\alpha_t(v, v')-1}}{\Gamma(\alpha_t(v, v'))} \tag{13}
\]

for \( \sum_{v' \in \mathcal{X}(v)} \theta_t(v, v') = 1 \) and \( \alpha_t(v, v') > 0 \) for all \( v' \in \mathcal{X}(v) \), and 0 otherwise.

Now consider a staging \( U \) that is not a trivial partition of \( S \). In Freeman and Smith (2011) we show that requiring margin equivalency to hold for its stages \( u \in U \) characterises the prior on the floret distributions. A stage \( u \) has margin equivalency when

\[
P(X(u) \mid \theta, U) = P(X(u) \mid \theta, S) \tag{14}
\]

where \( X(u) \) is the random variable with sample space \( \bigcup_{v' \in v_u} \{v' \cup \{v_u, \psi(v_u, v(v'))\}\} \), where \( v_u \) is any representative situation in \( u \), i.e. the sample space is the set of edge equivalence classes under a stage. This property is analogous to that of parameter modularity for Bayesian networks (Heckerman 1999). With the distribution for florets in \( S \) given above, this implies that the prior probability of \( \theta_t(u) \mid U_t = U, x^{t-1} \) has a Dirichlet distribution too, with hyperparameters that are sums of the corresponding hyperparameters under \( S \) of the constituent florets:

\[
f_{\theta_t(u)}(\theta_t(u) \mid U_t = U, x^{t-1}) = \Gamma \left( \sum_{v' \in \mathcal{X}(v_u)} \tilde{\alpha}_t(u, v') \right) \prod_{v' \in \mathcal{X}(v_u)} \frac{\theta_t(u, v')^{\tilde{\alpha}_t(u, v')-1}}{\Gamma(\tilde{\alpha}_t(u, v'))} \tag{15}
\]

where \( v_u \) is any situation in \( u \), \( \theta_t(u, v') \) are the elements of the vector \( \theta_t(u) \) and \( \tilde{\alpha}_t(u, v') = \sum_{v \in v_u} \alpha_t(v, \psi(v_u, v(v'))). \) Informally, equation (15) says that the hyperparameter vector for all of the floret distributions of the situations in stage \( u \) is equal to the sum of the hyperparameter vectors of the floret distributions under \( S \).

With margin equivalency and independence between the floret distributions under \( S \), the floret distributions under different stagings for stages composed of the same situations will always be the same. Therefore the probability distributions for a stage’s parameters (13) and (15) depend only the composition of the stage and not on the rest of the staging. This property is useful since it allows us to discuss the characteristics of stage clusters of variable groups without reference to the partition in which they
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appear. This makes individual models much simpler to explain. It also reduces the computational complexity in calculating (13) and (15).

As every \( \theta_t(u) \) is conditionally independent of all other quantities given \( \alpha_t(u) = ( \alpha_t(v) )_{v \in u} \), setting \( P(\theta_t \mid U_t, x^{t-1}) \) only requires the setting of \( \alpha_t(v) \) for each \( v \in S \) for every \( t \). This task can be simplified further by assuming

\[
f_{t+1,v}(\theta) = T(f_{t,v}^*(\theta))
\]

for some function \( T \) for all \( t > 1 \), where \( f_{t,v}(\theta) \) is the density of \( \theta_t(v) \mid x^{t-1}, U_t = S \) as given in equation (13), and \( f_{t,v}^*(\theta) \) is the density of \( \theta_t(v) \mid x^t, U_t = S \), so that for every \( v \in S \) only \( \alpha_1(v) \) needs to be set.

The simplest choice of \( T \) is the identity functional, so that \( f_{t+1,v}(\theta) = f_{t,v}^*(\theta) \) for \( t > 1 \). With \( f_{t,v}(\theta) \) as given in equation (13) and \( P(x_t(v) \mid \theta_t(v)) \propto \prod_{v' \in X(v)} \theta_t(v,v')^{x_t(v,v')} \) as given by equation (11), Bayes’ theorem requires

\[
f_{t,v}(\theta_t(v) \mid x^t) = \Gamma \left( \sum_{v' \in X(v)} \alpha^*_t(v,v') \right) \prod_{v' \in X(v)} \frac{\theta_t(v,v')^{\alpha^*_t(v,v')-1}}{\Gamma(\alpha^*_t(v,v'))}
\]

where \( \alpha^*_t(v,v') = \alpha_t(v,v') + x_t(v,v') \), and so

\[
\alpha_{t+1}(v) = \alpha_t(v) + x_t(v).
\]

As equation (18) is true for all \( t > 1 \), \( \alpha_t(v) \) can be written as a function of only \( \alpha_1(v) \) and \( x^{t-1}(v) \),

\[
\alpha_t(v) = \alpha_1(v) + \sum_{\tau=1}^{t-1} x_\tau(v)
\]

for all \( v \in S \).

Letting \( T \) be the identity functional reflects a modelling assumption that the underlying probabilities associated with each stage do not evolve from year to year for a given staged tree. Sometimes this will be too strong an assumption to make. In this case, a weaker set of assumptions are needed which will represent the fact that there is an “information drift” between the time points. This will also guard against spurious jumps in the model probabilities from expected model drift.

One way to characterise \( T \) to meet this need is provided by the power steady model (Smith, 1979, 1981, 1992). It was shown by Smith (1979) that if, loosely speaking, it is assumed that the Bayes decision under a step loss function would stay the same over time if no more information was gathered about the system but that the expected loss of the decision increases due to increasing uncertainty, then it is required that

\[
f_{t+1,v}(\theta) \propto (f_{t,v}^*(\theta))^k
\]

for some \( 0 < k \leq 1 \).
This transition function has a number of appealing properties. Firstly, when applied to the joint distribution \( P(\theta_t \mid U_t, x_t^{t-1}) \) of the \( \theta \) under any staging \( U_t \), the floret independence structure is kept intact.

Secondly, it can be shown that use of the steady model guards against misspecified priors, making predictions more robust. Let the local de Robertis measure \( \text{DR}_A \) be defined as follows (Smith and Daneshkhah 2010):

\[
d_{L}(f,g) = \sup_{\theta,\phi \in A} \{(\log f(\theta) - \log g(\theta)) - (\log f(\phi) - \log g(\phi))\}
\]

for any \( A \in \Theta \). Smith and Daneshkhah (2010) show that the local de Robertis measure is a separation measure whose separations do not change under Bayesian updating. It therefore represents artifacts of the model that cannot be changed by observation. It can easily be shown that, where \( f^* \propto f^k \) and similarly for \( g^* \),

\[
d_{L}(f^*,g^*) = k(d_{L}(f,g)).
\]

Thus using the steady model brings distributions closer together when \( 0 < k \leq 1 \). In this sense steady models tend to be robust against initial prior misspecification, if we see \( f \) as the prior used in the analysis and \( g \) as the “true” prior. See Smith and Rigat (2008) for further details.

A similar result can be shown for Kullback-Leibler (KL) distances (Kullback and Leibler 1951): recall that for two densities \( f \) and \( g \) the KL distance is given by

\[
d_{KL}(f;g) = - \int (\log f(\theta) - \log g(\theta))g(\theta)d\theta
\]

and that the entropy \( H \) of a density is given by

\[
H(f) = - \int f(\theta) \log f(\theta)d\theta.
\]

Let \( f_1, f_2 \) be any two densities such that \( H(f_1) = H(f_2) \). Then

\[
d_{KL}(p_{t+1}; f_1) - d_{KL}(p_{t+1}; f_2) = k(d_{KL}(p_t; f_1) - d_{KL}(p_t; f_2))
\]

where \( p_t \) is the density of the stage parameters at time \( t \), both after observing \( x^t \), so that \( p_{t+1} \propto (p_t)^k \). Equation (20) says that the distance between the density of the stage parameters under any staging and two arbitrary densities with the same entropy decreases by a fixed proportion at each time step, again indicating a robustness to prior mis-specification.

Thirdly, with \( \alpha_t^*(v) = \alpha_t(v) + x_t(v) \), equation (20) implies that \( \theta_{t+1}(v) \) is still distributed Dirichlet if \( \theta_t(v) \) is Dirichlet but with the hyperparameters of the distribution now given by the values

\[
\alpha_{t+1}(v, v') = k\alpha_t(v, v') + kx_t(v, v') - k + 1.
\]
Dynamic staged trees

Solving this recurrence relation for a constant $k$ yields

$$\alpha_t(v, v') = k^{t-1}(\alpha_1(v, v') - 1) + \sum_{\tau=1}^{t-1} k^{t-\tau} x_\tau(v, v') + 1$$

which heuristically can be seen as weighting recent observations more heavily for the setting of the latest prior.

Each situation can have its own $k$, $k(v)$, and it might be desired that this $k(v)$ be different for different $t$, for example when an external intervention in the system occurs.

We note that the use of the power steady model has a long history with Dirichlet distributions (e.g. in Smith (1979); Queen et al. (1994); Cowell et al. (2007)) and more generally (e.g. Ibrahim and Chen (2000); Rigat and Smith (2009)), and has also been used in Bayesian forecasting under the alternative name of exponential forgetting (Raftery et al. 2010). Here we use the power steady model as a justifiable and conjugate method for making inference about tree models whose floret probabilities evolve.

### 3.3 The staging distributions

We have allowed for drift over time in the values of probabilities associated with the conditional independence structure implicit in a staged tree model. However, we also want to allow for the possibility that the tree stagings themselves — and not just their parameters — evolve in time. It is unfeasible and usually unnecessary to model all possible changes of this partition space; in most applications it is appropriate to assume that changes in stage structure will be small in number and occur locally.

We therefore propose a dynamic model for the staged trees analogous to the Class 2 Multi-process Models used for dynamic linear models (DLMs) (Harrison and Stevens 1976; West and Harrison 1997). This was developed for the case where “no single [model] adequately describes what might happen to the process in the next time interval” (West and Harrison 1997). We describe the C2MPM here as it applies to the staged tree setting.

Let $U$ be the set of all possible stagings of $T$, and for each $U \in U$ and $t > 1$ let $\pi_t(U) = P(U_t = U \mid x^{t-1})$. There is obviously a large class of possible specifications for $\pi_t(U)$, but we note the three “practically important possibilities” mentioned by (West and Harrison 1997):

1. Fixed model selection probabilities, such that

$$\pi_t(U) = \pi(U) \quad \text{for all } t \geq 1.$$  

Here, one needs to only specify one prior over $U$. This prior remains fixed through time and is not changed by observations.

2. First-order Markov probabilities, where fixed transition probabilities between the models

$$\pi(U \mid U') = P(U_t = U \mid U_{t-1} = U')$$
are specified a priori, so that
\[ \pi_t(U) = \sum_{U' \in \mathcal{U}} \pi(U' | U') P(U_{t-1} = U' | x^{t-1}). \] (30)

Some initial prior distribution over the staged tree space, \( \pi_1(C) \), would need to be set.

3. Higher-order Markov probabilities, where the probabilities of the stagings at time \( t \) additionally depend on the stagings at \( t - 2, t - 3 \), etc.

While the first possible modelling strategy, of fixed staging probabilities, is much the simpler one (and the option used by West and Harrison for exposition), the second and third strategies are often going to be more accurate reflections of experts’ beliefs. We show here how to implement first-order Markov transitions between stagings.

A common assumption will be that \( \pi(U' | U) \) is larger the “closer” \( U \) is to \( U' \) in some sense, so that the underlying process is unlikely to change too much over a short period of time. If \( \pi(U' | U) = 0 \) for some \( U \in \mathcal{U} \), this has the advantage of reducing the number of terms in equation (30).

We therefore require a metric on \( \mathcal{U} \) and then let \( \pi(U' | U) \) be a function of this metric. Any intuitive metric on general sets of partitions can be used, e.g. that of Meila (2007). A simple metric that we use here can be derived from the Hasse diagram of the lattice of partitions of \( S \) under the relation “finer than” (see Stanley (1997) for a detailed overview of such lattice terminology). The Hasse diagram for \( |S| = 4 \) is shown in Figure 3.

![Hasse diagram](image)

Figure 3: The Hasse diagram of the lattice of partitions of \( S \) when \( |S| = 4 \)
Dynamic staged trees

The length of the shortest path between two partitions on the Hasse diagram is a metric on the partition space of \( S \), and we call it \( \ell \) here. A distance of \( \ell = 1 \) represents the division of a stage or the merging of two stages. One possible way to set \( \pi(U \mid U') \) based on this metric is

\[
\pi(U \mid U') = \begin{cases} 
\rho & \text{if } U = U' \\
[B_r(U')]^{-1}(1 - \rho) & \text{if } 0 < \ell(U, U') \leq \epsilon, \\
0 & \text{otherwise}
\end{cases}
\]

where \( B_r(U') = \{ U \in U : \ell(U, U') \leq \epsilon, U \neq U', U \in U \} \) is an \( \epsilon \)-ball of stagings around \( U' \) under the \( \ell \) metric. This represents a belief that the underlying symmetry process changes only locally and slowly. If more radical changes in the symmetry process are taking place due to external intervention in the system then the methodology in Section 4.1 can be deployed.

The other term in (29), \( P(U_{t-1} = U' \mid x^{t-1}) \), can be calculated for each \( U_{t-1} \) using Bayes’ theorem:

\[
P(U_{t-1} = U' \mid x^{t-1}) \propto P(x_{t-1} \mid U_{t-1} = U')P(U_{t-1} = U' \mid x^{t-2})
\]

\[
= \sum_{U' \in U} P(x_{t-1} \mid U_{t-1} = U')P(U_{t-1} = U' \mid x^{t-2})
\]

The \( P(U_{t-1} = U' \mid x^{t-2}) \) terms on the right-hand side of (33) will already be available at time \( t - 1 \). The term \( P(x_{t-1} \mid U_{t-1} = U') \), meanwhile, can be calculated as follows, using equations (11) and (15) at time \( t - 1 \):

\[
P(x_{t-1} \mid U_{t-1} = U') = \int_{\Theta_{t-1}} P(x_{t-1} \mid \theta_{t-1}, U_{t-1} = U')P(\theta_{t-1} \mid U_{t-1} = U')d\theta_{t-1}
\]

\[
\propto \int_{\Theta_{t-1}} \prod_{u \in U'} \left( \frac{\Gamma \left( \sum_{v' \in X(v_u)} \tilde{\alpha}_{t-1}(u, v') \right)}{\prod_{v' \in X(v_u)} \theta_{t-1}(u, v')^{\tilde{\alpha}_{t-1}(u, v') - 1}} \right) d\theta_{t-1}
\]

\[
= \prod_{u \in U'} \left( \frac{\Gamma \left( \sum_{v' \in X(v_u)} \tilde{\alpha}_{t-1}(u, v') \right)}{\prod_{v' \in X(v_u)} \Gamma(\tilde{\alpha}_{t-1}(u, v'))} \right)
\]

where \( v_u \) is any situation in \( u \), \( \tilde{\alpha}_{t-1}(u, v') = \tilde{x}_{t-1}(u, v') + \tilde{\alpha}_{t-1}(u, v') \), where \( \tilde{x}_{t-1}(u, v') = \sum_{v \in U} x_{t-1}(v, \psi_u(v_u, v)(v')) \) and \( \tilde{\alpha}_{t-1} \) is as defined in equation (15). Note that the outer product is over the stages \( u \in U' \) due to \( |U'| \) being the dimension of \( \Theta_{t-1} \), whereas in equation (11) the relevant index is that of the situation being sampled from.

The number of terms in (30) can be reduced further by setting the values of \( P(U_{t-1} = U' \mid x^{t-1}) \) below a threshold \( q \) as zero and normalising the remaining probabilities to...
ensure they still sum to 1. A similar approach advocated by [Madigan and Raftery (1994)] as “Occam’s window” is to discard models $U'$ that are not in the set

$$U^*_t = \left\{ U_t \in U : \frac{P(U_t \mid x^t)}{\max_U P(U \mid x^t)} \leq q \right\}$$

(37)

for some $0 < q < 1$, i.e., to only keep models where the Bayes factor between them and the most probable model a posteriori are above a certain threshold. This has the advantage of guaranteeing that at least one model will be kept.

### 3.4 One-step-ahead prediction

The terms in equation (7) can now be defined, using the foregoing, as follows:

$$P(x_t \mid \theta_t, U_t, x^{t-1}) = \sum_{N_t(v_0)} P(N_t(v_0) \mid \theta_t, U_t, x^{t-1}) \prod_{v \in S} \prod_{v' \in \mathcal{X}(v)} \pi(U_t(v'), x_t(v', v'))$$

(38)

$$P(\theta_t \mid U_t, x^{t-1}) = \Gamma \left( \sum_{v' \in \mathcal{X}(v_u)} \tilde{\alpha}_t(u, v') \right) \prod_{v' \in \mathcal{X}(v_u)} \frac{\theta_t(u, v') \tilde{\alpha}_t(u, v')^{-1}}{\Gamma(\tilde{\alpha}_t(u, v'))}$$

(39)

$$P(U_t \mid x^{t-1}) = \sum_{U_t \in U} \pi(U_t \mid U_{t-1}) P(U_{t-1} \mid x^{t-1})$$

(40)

where $A$ is the event $\forall v \in u \setminus v_0, \sum_{v'} x_t(v, v') = x_t(v^*, v)$. If it is assumed that the distribution of $N_t(v_0)$ depends only on $x^{t-1}$ then (7) can be further simplified to the closed-form solution

$$P(x_t \mid x^{t-1}) = \sum_{U_t \in U} \left( \sum_{U_{t-1} \in U} \pi(U_t \mid U_{t-1}) P(U_{t-1} \mid x^{t-1}) \right) \left( \sum_{N_t(v_0)} P(N_t(v_0) \mid x^{t-1}) \right)$$

$$\times \prod_{v \in x_t} \left[ \frac{\Gamma \left( \sum_{v' \in \mathcal{X}(v_u)} \tilde{\alpha}_t(u, v') \right)}{\Gamma \left( \sum_{v' \in \mathcal{X}(v_u)} \tilde{\alpha}_t^*(u, v') \right)} \prod_{v' \in \mathcal{X}(v_u)} \frac{\Gamma(\tilde{\alpha}_t^*(u, v'))}{\Gamma(\tilde{\alpha}_t(u, v'))} \right] \cdot I_A$$

(41)

If $N_t(v_0)$ is always known, then (41) can be simplified further to become

$$P(x_t \mid x^{t-1}) = \sum_{U_t \in U} \left( \sum_{U_{t-1} \in U} \pi(U_t \mid U_{t-1}) P(U_{t-1} \mid x^{t-1}) \right)$$

$$\times \prod_{v \in x_t} \left[ \frac{\Gamma \left( \sum_{v' \in \mathcal{X}(v_u)} \tilde{\alpha}_t(u, v') \right)}{\Gamma \left( \sum_{v' \in \mathcal{X}(v_u)} \tilde{\alpha}_t^*(u, v') \right)} \prod_{v' \in \mathcal{X}(v_u)} \frac{\Gamma(\tilde{\alpha}_t^*(u, v'))}{\Gamma(\tilde{\alpha}_t(u, v'))} \right] \cdot I_A$$

(42)
4 Causal intervention

With many forecasting systems there is also an attendant need to consider the effects of external intervention in the system, including by the forecasters themselves (West and Harrison 1989). This ensures that all relevant information is taken into account, increasing the accuracy of future forecasts.

The predicted effect of an intervention depends both on the nature of that intervention and the context in which it applies. Many interventions act only on certain local features of a model while leaving the other features of the model unchanged. We note that these types of interventions have now been extensively studied on non-dynamic BNs (Pearl 2000; Spirtes et al. 2001), which are called causal Bayesian networks (CBNs) in this context. Dynamic extensions of CBNs also exist (Queen and Smith 1993; Eichler and Didelez 2007; Queen and Albers 2009). A Bayesian way of learning causal Bayesian networks is given by Cooper and Yoo (1999).

We believe that tree-based graphical models are very useful in general for undertaking causal analysis. Due to the multiple representations of each variable in the graph — one for each possible path-history on parent variables — much more refined interventions in the system can be represented (Shafer 1996). How causal hypotheses can be represented within the framework of static CEGs has been investigated by Thwaites and Smith (2006) and Thwaites et al. (2010).

We now show how causal analysis affects the one-step ahead forecast on a dynamic staged tree given by equations (41) and (42) for two different types of intervention: one on the possible stagings on a tree $T$ and one on the structure of the tree $T$ itself.

4.1 Intervention on the staging distribution

Suppose that at time $t$ it is hypothesised that an intervention will cause particular situations to be moved into their own stage $u^\dagger$. For example, in our educational example, the exams for the second module might be tailored so that performance in the first module is no longer a predictor in how well students should perform in it. The one-step ahead forecasts then change by reflecting the updated probability distribution over stagings.

Recall that $\pi^*_t(U)$ denotes $P(U_{t-1} = U \mid x^{t-1})$, the posterior probability of the staging $U$ in the unintervened system. After the intervention described above, which we denote with $I_t$, the probability of $U$ being the staging at time $t$ will be $\pi^u_t(U) = P(U_t = U \mid x^{t-1}, I_t)$. The decision needs to be made for how to relate the distribution of $\pi^u_t(U)$ to that of $\pi^*_t(U)$ over $U$. One approach is to let

$$\pi^u_t(U^\dagger) = \sum_{U \in U^{-\dagger}} \pi^*_t(U)$$

(43)

where $U^{-\dagger}$ is the set of stagings that would be turned into $U^\dagger$ under the intervention described above of having a set of situations $u$ put into the same stage.
A remaining issue is how the distribution of $\theta_t \mid U_t$ is affected. In the absence of further information, a good default is to use the steady model as in the idle system but with a lower value for the steady parameter $k$. This indicates that past data might not be considered as relevant in helping to make predictions as it would have been under the idle system. We note that this is analogous to setting a higher variance on evolution parameters in dynamic linear models when forecasting after an external intervention (Section 1.2.2 of [West and Harrison](1997)).

### 4.2 Intervention on $T$

Recalling the event tree pictured in Figure 1, consider the case where at time $t$ the course directors decide to eliminate the first module on the tree from course. This means that the marks that students would have got for this module are unknown from that time onwards, and therefore all of the data at time $t$ for this module will be concentrated on the second (“NA”) edge of the $v_1$ floret.

This type of intervention is analogous to the $do$ operator introduced for CBNs by ([Pearl](2000)), where a random variable is forced to take a particular value with probability 1. The difference with CBNs is that staged trees allow a richer set of interventions on their structure, including letting an intervention take place at specific times and situations, and not merely changing the value of a variable under all circumstances.

We assume that the probability distributions on any unmanipulated florets remain unchanged, just as manipulations on CBNs are local ([Pearl](2000)). We will also assume that once an intervention is made, it endures thereon. We now describe how the learning framework outlined previously can be adapted to prediction after an intervention of this type occurs.

Without loss of generality, say that at time $t$ an intervention $I_t(v, v')$ at situation $v \in S$ occurs so that $\theta_t(v, v')$ is equal to 1 for a specific $v' \in \mathcal{X}(v)$ and to 0 for all other $v^* \in \mathcal{X}(v)$. By the definition of the event tree, along with the causal assumptions, all other floret distributions are technically unchanged. However, notice that the probability of reaching any node in any $\Lambda(v^*, T)$, the sub-tree with $v^*$ as the root node, is zero. It follows that the tree $T$ is equivalent to the reduced tree $T'$ where all $\Lambda(v^*, T)$ are deleted and only the edge $(v, v')$ remains in the floret $\mathcal{F}(v)$, and so the process can henceforth be considered to take place on this reduced tree $T'$.

The one-step ahead forecasts can now be calculated as before with only a few modifications that are due to the set of situations $S$ changing; call this new set $S^\dagger$. Firstly, the distribution over $U^\dagger$, the new set of possible stagings, must be set. There are several possible choices here. In the absence of any other information, a good default is to let

$$P(U_t = U^\dagger \mid x^{t-1}, I_t(v, v')) = P(U_{t-1} = U \mid x^{t-1}), \quad (44)$$

where $U^\dagger$ is the staging formed from $U$ by replacing each stage $u \in U$ with a new stage $u^\dagger := u \setminus \{v^\dagger\}_{v^\dagger \in S_t \setminus S^\dagger}$, and by splitting the stage $u^\dagger \in U^\dagger$ that contains the intervention node $v$ into $u^\dagger \setminus v$ and $v$. 

Secondly, the distributions of the stage parameters \( \theta_t(u) \) for any \( u \) need to be reconsidered. Under the causal assumptions considered here, interventions have only local effects, so a sensible default model is to let \( f_{\theta_t(u)}(\theta_t(u) \mid U_t, x^{t-1}, I_t(v, v')) \) be calculated as before, i.e. as given in Equation (15).

Assuming that all of the other system characteristics, e.g. the steady model and the multinomial sampling, are intact post-intervention, the one-step ahead forecast (41) is adjusted slightly to become

\[
P(x_t \mid x^{t-1}, I_t(v, v')) = \sum_{U_t^1 \in U^1} \left( P(x_t \mid U_t^1) \sum_{U_{t-1} \in U} \pi^{1}(U_t^1 \mid U_{t-1}) P(U_{t-1} \mid x^{t-1}) \right)
\]

where

\[
P(x_t \mid U_t^1) = \sum_{N_t(v_0)} \left( P(N_t(v_0) \mid x^{t-1}) \prod_{u \in U_t^1} \frac{\Gamma \left( \sum_{v' \in X(v_u)} \bar{\alpha}_t(u, v') \right)}{\Gamma \left( \sum_{v' \in X(v_u)} \bar{\alpha}_t^*(u, v') \right)} \prod_{v' \in X(v_u)} \frac{\Gamma(\bar{\alpha}_t^*(u, v'))}{\Gamma(\bar{\alpha}_t(u, v'))} \right) \cdot I_A
\]

and where \( \pi^{1}(U_t^1 \mid U_{t-1}) = \pi(U_t \mid U_{t-1}) \) by the argument above.

5 A simple educational example

In this section we illustrate how to carry out one-step ahead predictions with dynamic staged trees using 12 years’ worth of exam marks for two first-year undergraduate modules. The underlying event tree is that shown in Figure 1.

We made the following assumptions:

1. \( N_t(v_0) \) was known for all values of \( t \).

2. The distribution over the root-to-leaf paths at time \( t = 1 \) under \( U_1 = S \) was Dirichlet with all path hyperparameters equal to 1.

3. For the transitions between stagings we used the \( \ell \) metric with \( \epsilon = 1 \), i.e. only transitions between models with one local change were considered.

We present here the posterior probabilities \( P(U_t \mid x^t) \) for the stagings after \( t = 1 \) for each time \( t \) for different hyperparameter values, when analysed with and without an external intervention. Recall that the same analysis can also produce predictions \( P(x_t \mid x^{t-1}) \) if desired.

In a fuller analysis this application could be run over a distribution of the hyperparameters \( k \) (the steady model parameter), \( \rho \) (the probability of the underlying model...
G. Freeman and J. Q. Smith

Table 1: All stagings with positive probabilities at each time $t$ for $k = 0.9$, $\rho = 0.9$, $q = 0.2$ with $P(U_t = \{v_1, v_2, \{v_3, \ldots, v_6\}, \{v_7, \ldots, v_{10}\}) = 1$

<table>
<thead>
<tr>
<th>Time</th>
<th>$U_t$</th>
<th>$P(U_t \mid x^t)$</th>
</tr>
</thead>
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<td>1</td>
</tr>
<tr>
<td>2</td>
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<tr>
<td></td>
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</tr>
<tr>
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<td>0.766</td>
</tr>
<tr>
<td></td>
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<td>0.233</td>
</tr>
<tr>
<td>4</td>
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</tr>
<tr>
<td></td>
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<td>0.322</td>
</tr>
<tr>
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<td>0.328</td>
</tr>
<tr>
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</tr>
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<td>1</td>
</tr>
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<tr>
<td></td>
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</tr>
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<td>1, 2, 3, {4,5,6}, {7,10}, 8, 9</td>
<td>1</td>
</tr>
</tbody>
</table>

5.1 Analysis of the series without intervention

In Table 1 we present $P(U_t \mid x^t)$ for $t = 1 \ldots 12$ for the model where $U_t = \{v_1, v_2, \{v_3, \ldots, v_6\}, \{v_7, \ldots, v_{10}\})$ with probability 1 and $k = 0.9$, $\rho = 0.9$ and $q = 0.2$. The latter two parameter values ensure that few new models will be kept in the analysis, as the high value of $\rho$ gives a low prior probability on transitions between stagings and this value of $q$ makes the Occam’s window set of equation (37) small. This speeds up the computation of the forecasts at the expense of possibly worse predictions through fewer stagings being included in the model averaging.

An alternative way of presenting this information is to plot how $P(v_i, v_j \in u)$, the probability that situations $v_i, v_j$ are in the same stage, changes for increasing values of $t$. Figure 4 shows this for the information in Table 1.

To illustrate how the level of detail in the staging distribution changes as a function of the hyperparameters, we carried out the analysis again with radically different values: we set $k = 0.5$ (so that floret distributions are flattened more quickly and therefore past
Figure 4: Plots of probabilities that each pair of situations are in the same stage for different values of \( t \), for the case when \( k = 0.9, \rho = 0.9, q = 0.2 \) with \( P(U_1 = \{v_1, v_2, \{v_3, \ldots, v_6\}, \{v_7, \ldots, v_{10}\}\}) = 1 \), using the values in Table 1.

Observations more heavily discounted), \( \rho = 0.25 \) (so that the probability of moving between stagings is more likely), and \( q = 0.05 \) (so that stagings with poorer Bayes factors relative to the most likely are kept in the analysis) with \( P(U_1 = \{v_1, v_2, \{v_3, \ldots, v_6\}, \{v_7, \ldots, v_{10}\}\}) = 1 \) still assumed for consistency. The resulting matrix plot of probabilities of situations being in the same stage against time is as shown in Figure 5.

It can be seen from the latter figure that the analysis with the new hyperparameter values gives much the same qualitative description of the system as the more conservative hyperparameters at greater computational expense, with the pay-off of greater detail.

Some interesting characteristics of the system can be discerned from this initial exploratory analysis of this system. With regard to the situations concerning whether marks are available for a module or not, \( \theta(v_3) \) — the probability distribution for the second module's marks being available given that the mark in the first module is itself missing — does not appear to be related to the others at any time point. Until \( t = 7 \), \( v_4, v_5 \) and \( v_6 \), the situations representing the probability of marks being missing in the second module after gaining a high, medium or low mark respectively in the first module, had high but falling probabilities of being in the same stage, implying that independence of the second module's marks being missing from skill in the first module kept decreasing from a high point. At \( t = 8 \), in contrast, these probabilities are much
Figure 5: Plots of probabilities that each pair of situations are in the same stage for different values of $t$, for the case when $k = 0.5$, $\rho = 0.25$, $q = 0.05$ with $P(U_1 = \{v_1, v_2, \{v_3, \ldots, v_6\}, \{v_7, \ldots, v_{10}\}\}) = 1$

lower, although the probability distributions of marks being missing after gaining a medium or low mark in the first module are deemed to become slightly more likely to be the same after that, with students performing well in the first module continuing to have a very different probability distribution for the missingness of their second module marks. We investigate a possible causal hypothesis that might explain what might have changed at $t = 8$ in the next section.

Another notable finding is that $v_7$ and $v_{10}$ — the situations concerning marks in the second module after getting a poor grade or having a missing mark in the first module, respectively — are always strongly related. It therefore appears that the second module marks of students who did poorly in the first module should be used to predict the second module performance of students whose first module marks are missing.

While the less conservative hyperparameter values in the latter analysis allow for more exploration of the space of possible stagings, it is inevitable that it will also take longer to run the algorithm. Implemented in R, our program to produce the posterior probabilities of the stagings at each time point took 5 minutes in the former analysis but just over 2 hours in the latter. The trade-off is transparent to the analyst and can be tuned as desired.
It is worth noting again that these detailed homogeneities would not have been as easily identifiable if the model class was restricted to Bayesian networks.

5.2 Analysis of the series after intervention

We also carried out an analysis with the latter parameters after a hypothesised causal intervention: we assumed that at $t = 8$ the situations for the grades $(v_2, v_7, v_8, v_9, v_{10})$ were put into the same stage. This could have happened, for example, because the modules were re-defined to be very similar in difficulty for students with different skills. The resulting matrix of probabilities of situations being in the same stage through time is shown in Figure 6.

![Figure 6: Plots of probabilities that each pair of situations are in the same stage for different values of $t$, for the case when $k = 0.5$, $\rho = 0.25$, $q = 0.05$ with $P(U_1 = \{v_1, v_2, \{v_3, \ldots, v_6\}, \{v_7, \ldots, v_{10}\}\}) = 1$, and situations $v_2, v_7, v_8, v_9, v_{10}$ caused to be in the same stage at $t = 8$](image-url)

It can be seen that the probabilities are not too different from those in Figure 5, but there are increased probabilities of $v_8, v_9$ and $v_{10}$ being in the same stage even for $t > 8$, which indicates slightly higher probabilities of dependence between the second module’s grades for students who performed differently in the first module under the causal hypothesis considered here.
6 Discussion

We have presented in this paper a new discrete time series modelling class, the dynamic staged tree, that is intuitive to use and suitable for carrying out causal analysis.

Obviously the class of models we define here can be usefully refined. In many potential applications we would like to allow for multiple possible trees at any time point. If the general class of event trees $T$ is required, then $P(x_t \mid x_{t-1})$ can still be calculated as outlined in this paper but with the additional step of marginalising over the $T \in T$ such that $P(T \mid x_{t-1}) > 0$, assuming the number of such $T$ is tractable. If all that is required is the subclass of $T$ which consists of trees that are merely different partitionings of the same set of root-to-leaf path events, then assuming that the same root-to-leaf path events on different trees have the same probability, the floret distributions on all trees can be characterised as Dirichlet by the method used here. The method of assigning probabilities over the tree space in either case, or how those probabilities change over time, would still need to be resolved. We plan to explore this class in a later paper.

Another way of enlarging the model space is to allow for uncertainty in $\psi_u(v, v')$, i.e. which edges of two florets are coloured identically when their root nodes are in the same stage. The type of hypothesis this could capture includes a belief in the stability in values between different random variables. In our educational example, this would translate into believing that the probability of getting the same grade in the second module as in the first one is the same for all grades.

We note that the number of possible $\psi_u(v, v')$ for any pair $v, v'$ is $|X(v)|!$. Therefore to make the model search tractable in general either the number of possibilities must be restricted using contextual information or a local neighbourhood switching function, as with the staging space model in this paper.

Various classes of discrete multivariate time series are of course well studied. Possibly the closest class to the one considered here is the model used in event history analysis. Event history data relates to the time at which events of interest occur, rather than the reverse representation of what events occur at specific time points. Formally, an event history can be identified as a marked point process, a set $\{(T_s, E_s) : s = 1, \ldots, S\}$ of pairs of times $T_s$ when events $E_s$ occurred, where the times are random variables while the events of interest are fixed beforehand, although their order might be uncertain a priori (Arjas 1989). Two graphical models developed for event history analysis are local independence graphs (Didelez 2008) and graphical duration graphs (Gottard 2007). While there is an overlap between event history data and the problem outlined here, it is clear that the two address quite separate concerns. In event history analyses the number of events under consideration is typically small, with the focus of analysis being the time of events, usually allowed to occur within a continuous time domain. Here, in contrast, we wished to model a complex discrete distribution over a discrete time domain. We are currently investigating the link between staged tree models of the type described here and event history models which explicitly acknowledge this extra source of variation.

Finally, it appears that the dynamic staged tree process can be extended to model
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processes defined on continuous as well as discrete variables. Converting the leaf nodes on a tree into continuous sample spaces is trivial. When other variables are continuous then analogous conjugate models can be defined which describe hierarchical clustering models. We will report on these developments in a later paper.

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