# The Dirichlet-tree distribution <br> Tom Minka <br> Justsystem Pittsburgh Research Center <br> July, 1999 (revised Oct, 2004) 


#### Abstract

This note further explores the Dirichlet-tree distribution developed by Dennis (1991). An inclusion relationship is given for different tree structures and an independence relationship is given for probabilities in the tree. The posterior, evidence, and predictive density are derived for multinomial observations.


## 1 Introduction

The Dirichlet distribution has enjoyed considerable popularity as a prior distribution for multinomial parameters. This is mainly because the Dirichlet is conjugate to the multinomial under the conventional parameterization. But the Dirichlet distribution has key limitations:

1. Each variable has its own mean, but they must all share a common variance parameter.
2. Aside from the constraint that they sum to one, the variables are mutually independent (Mosimann, 1962).

This note describes a new distribution which overcomes these limitations while preserving computational simplicity. The new distribution is simply the conjugate distribution to the multinomial under a different parameterization.


Figure 1: (a) A finite stochastic process (b) Notation for a general process
Instead of representing a multinomial sample as the outcome of a $K$-sided die, we can represent it as the outcome of a finite stochastic process, such as the tree shown in figure 1(a). The probability of a leaf is the product of branch probabilities leading to that leaf. To represent an arbitrary tree,
we can use the notation in figure $1(\mathrm{~b})$. Before, the parameters were the leaf probabilities $\left[p_{1} \ldots p_{K}\right]$, so that the probability of a sample $x$ was

$$
\begin{equation*}
p(x \mid \mathbf{p})=\prod_{k=1}^{K} p_{k}^{\delta(x-k)} \tag{1}
\end{equation*}
$$

Under the tree parameterization, it is instead written

$$
\begin{align*}
p(x \mid \mathbf{B}, T) & =\prod_{(\text {nodes } j)} \prod_{(\text {branches } c)} b_{j c}^{\delta_{j c}(x)}  \tag{2}\\
\delta_{j c}(x) & = \begin{cases}1 & \text { if branch } j c \text { leads to } x \\
0 & \text { otherwise }\end{cases} \tag{3}
\end{align*}
$$

The probability of reaching any interior node can also be computed this way. The conjugate prior for this parameterization is no longer a single Dirichlet density but rather a product of Dirichlet densities, one for each node:

$$
\begin{align*}
p(\mathbf{B} \mid \alpha) & =\prod_{\text {(nodes } j \text { ) }} p\left(\mathbf{b}_{j} \mid \alpha\right)  \tag{4}\\
p\left(\mathbf{b}_{j} \mid \alpha\right) & \sim \mathcal{D}\left(\alpha_{j c}\right) \tag{5}
\end{align*}
$$

The "Dirichlet-tree distribution" is the distribution over leaf probabilities $\left[p_{1} \ldots p_{K}\right.$ ] that results from this prior on branch probabilities. The distribution is a function of the tree structure $T$ as well as $\alpha$. The explicit density over $\left[p_{1} \ldots p_{K}\right]$ can be computed by noting that

$$
\begin{equation*}
b_{j c}=\frac{\sum_{k} \delta_{j c}(k) p_{k}}{\sum_{k c^{\prime}} \delta_{j c^{\prime}}(k) p_{k}} \tag{6}
\end{equation*}
$$

i.e. $b_{j c}$ is proportional to the probability mass in its subtree. Combining this with (4) and multiplying by the Jacobian gives (Dennis, 1991)

$$
\begin{align*}
p(\mathbf{p} \mid \alpha, T) & =\prod_{k} p_{k}^{\alpha_{\text {parent }(k)}-1} \prod_{j} \frac{\Gamma\left(\sum_{c} \alpha_{j c}\right)}{\prod_{c} \Gamma\left(\alpha_{j c}\right)}\left(\sum_{k c} \delta_{j c}(k) p_{k}\right)^{\beta_{j}}  \tag{7}\\
\beta_{j} & =\alpha_{\text {parent }(j)}-\sum_{c} \alpha_{j c} \quad \text { (or } 0 \text { if } j \text { is the root node) } \tag{8}
\end{align*}
$$

where $\alpha_{\text {parent }(j)}$ means the $\alpha$ parameter for the branch immediately leading to $j$.
Since the Dirichlet distribution at a node can be arbitrarily broad or sharp, the Dirichlet-tree distribution can give an independent variance to each $p_{k}$. Also, the leaves in a subtree are correlated since they all depend on the ancestors of that subtree.

## 2 Properties

The Dirichlet-tree distribution obviously generalizes the Dirichlet distribution, since we can always have a tree of depth 1. Surprisingly, a depth 1 tree is not necessary: it is possible for any tree
structure to realize a Dirichlet distribution on $\left[p_{1} \ldots p_{K}\right]$, if the $\alpha$ 's are chosen carefully. In particular, if the $\beta_{j}$ in (7) are all zero, then the distribution is Dirichlet. This result is generalized in the following theorem:

Theorem 1 Let $S(T)$ be the set of Dirichlet-tree distributions realizable by tree structure $T$. If $T^{\prime}$ is identical to $T$ except for an additional interior node, then $S\left(T^{\prime}\right)$ is a proper superset of $S(T)$.

Proof $T^{\prime}$ adds a new $\alpha$ to the tree and a new $\beta_{j}$ to (7). By setting $\beta_{j}=0$, we can realize anything in $S(T)$. If $\beta_{j} \neq 0$, then we can get distributions not in $S(T)$.


The marginal distribution of any $p_{k}$ is difficult to compute, since it is a multiplicative convolution of Dirichlet densities. But the moments of $p_{k}$ are easy. The first two moments are (Dennis, 1991):

$$
\begin{align*}
& E\left[p_{k}\right]=\prod_{j c} E\left[b_{j c}\right]^{\delta_{j c}(k)}=\prod_{j c}\left(\frac{\alpha_{j c}}{\sum_{c^{\prime}} \alpha_{j c^{\prime}}}\right)^{\delta_{j c}(k)}  \tag{9}\\
& E\left[p_{k}^{2}\right]=\prod_{j c} E\left[b_{j c}^{2}\right]^{\delta_{j c}(k)}=E\left[p_{k}\right] \prod_{j c}\left(\frac{1+\alpha_{j c}}{1+\sum_{c^{\prime}} \alpha_{j c^{\prime}}}\right)^{\delta_{j c}(k)} \tag{10}
\end{align*}
$$

The most probable $\mathbf{p}$, given by maximizing (7), corresponds to setting the branch probabilities to

$$
\begin{equation*}
b_{j c} \propto \alpha_{j c}-\sum_{k} \delta_{j c}(k) \tag{11}
\end{equation*}
$$

The subtracted term is simply the total number of leaves under branch $j c$. In the Dirichlet case, this reduces to the usual formula $p_{k} \propto \alpha_{k}-1$.

A special case of the Dirichlet-tree distribution was developed by Connor \& Mosimann (1969). It restricts the tree structure to be a binary cascade, as shown to the right.


Connor \& Mosimann showed that the cascade-tree distribution satisfies a certain independence property. The following theorem generalizes this property to arbitrary trees.

Theorem 2 Let $a$ be a node in the tree such that nodes $b$ and $c$ are descendants of $a$ while node $d$ is not. Then the ratio $p_{b} / p_{c}$ is independent of $p_{d}$. That is, relative probabilities in a subtree are independent of anything outside the subtree.

Proof The probability of a node is the product of branch probabilities leading to it. The ratio $p_{b} / p_{c}$ only involves branch probabilities below $a$, so the theorem follows.


## 3 Bayesian inference

Since the Dirichlet-tree distribution is a conjugate distribution, it is straightforward to compute the posterior distribution given data $D=\left\{x_{1} \ldots x_{N}\right\}$ :

$$
\begin{align*}
p(\mathbf{B} \mid D, \alpha, T) & \propto p(\mathbf{B} \mid \alpha) \prod_{i} p\left(x_{i} \mid \mathbf{B}, T\right)  \tag{12}\\
& \propto \prod_{j c} b_{j c}^{\alpha_{j c}-1} b_{j c}^{\sum_{i}} \delta_{j c}\left(x_{i}\right)  \tag{13}\\
& \sim \prod_{j} \mathcal{D}\left(\alpha_{j c}+\sum_{i} \delta_{j c}\left(x_{i}\right)\right) \tag{14}
\end{align*}
$$

We simply add 1 to $\alpha_{j c}$ for every data point reachable from branch $j c$. Each observation $x$ is equivalent to a boolean matrix of decision outcomes, given by $\delta_{j c}(x)$. So from the perspective of node $j$, it is just as if its children were leaves. The same result holds for the evidence:

$$
\begin{align*}
p(D \mid \alpha, T) & =\int_{\mathbf{B}} p(\mathbf{B} \mid \alpha) \prod_{i} p\left(x_{i} \mid \mathbf{B}, T\right)  \tag{15}\\
& =\prod_{j}\left(\frac{\Gamma\left(\sum_{c} \alpha_{j c}\right)}{\Gamma\left(\sum_{c} \alpha_{j c}+\sum_{c} n_{j c}\right)} \prod_{c} \frac{\Gamma\left(\alpha_{j c}+n_{j c}\right)}{\Gamma\left(\alpha_{j c}\right)}\right)  \tag{16}\\
n_{j c} & =\sum_{i} \delta_{j c}\left(x_{i}\right) \tag{17}
\end{align*}
$$

The predictive density for a new sample $D^{\prime}=\left\{y_{1} \ldots y_{M}\right\}$ is therefore

$$
\begin{align*}
p\left(D^{\prime} \mid D, \alpha, T\right) & =p\left(D^{\prime}, D \mid \alpha, T\right) / p(D \mid \alpha, T)  \tag{18}\\
& =\prod_{j}\left(\frac{\Gamma\left(\sum_{c} \alpha_{j c}+\sum_{c} n_{j c}\right)}{\Gamma\left(\sum_{c} \alpha_{j c}+\sum_{c} n_{j c}+\sum_{c} m_{j c}\right)} \prod_{c} \frac{\Gamma\left(\alpha_{j c}+n_{j c}+m_{j c}\right)}{\Gamma\left(\alpha_{j c}+n_{j c}\right)}\right)  \tag{19}\\
m_{j c} & =\sum_{i} \delta_{j c}\left(y_{i}\right) \tag{20}
\end{align*}
$$

These results also appear in Dennis (1996).


Figure 2: Greedy construction of tree structure

## 4 Learning the tree structure

Suppose we want to find the tree structure which best fits a data set. The data set is a matrix of counts $D=\left\{\mathbf{x}_{1} \ldots \mathbf{x}_{N}\right\}$ where each column $\mathbf{x}_{s}$ is a set of samples. For any particular tree structure, we can convert this dataset into a matrix of counts $n_{j c}$, which is the number of times branch $j c$ was taken. Define $D_{j}=\left\{n_{j 1}, \ldots, n_{j C}\right\}$ to be the dataset from the perspective of node $j$.

Formally, we want to do model selection by maximizing the probability of the data, with parameters integrated out (the "structure evidence"):

$$
\begin{align*}
p(D \mid T) & =\int_{\alpha} p(D \mid \alpha, T) p(\alpha \mid T)  \tag{21}\\
& =\prod_{j} \int_{\alpha_{j}} p\left(D_{j} \mid \alpha_{j}\right) p\left(\alpha_{j}\right) \tag{22}
\end{align*}
$$

Because each interior node has separate $\alpha$ parameters, this decouples into a product of integrals, one for each interior node. Intuitively, the structure evidence for an interior node measures how well the data for that node matches an ordinary Dirichlet distribution.

To evaluate this criterion efficiently, one could use Laplace's method at the maximum-likelihood estimate of $\alpha$, or a variational method based on the lower bounds derived by Minka (2000).

Given an efficient way to evaluate the evidence at each interior node, one way to optimize it is to greedily build the tree structure top-down, as illustrated in figure 2. By Theorem 1, we only need to consider binary trees. Start with one interior node. At each step, pick an interior node $r$ with $>2$ children (which must all be leaves), and introduce two new interior nodes, $j_{1}$ and $j_{2}$, which will parent the children. Each possible division of the children is scored using the evidence. Note that when we make a local change to the tree structure, most of the terms in (22) are unchanged. In particular, when we introduce $j_{1}$ and $j_{2}$, we change the term for $r$ and add terms for $j_{1}$ and $j_{2}$. All the other terms are unchanged.

Further development of these ideas is left to the reader.

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