

PROBABILISTIC ALGORITHMS FOR TREES

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1. Introduction and Definitions

A *rooted tree*, τ , is a partially ordered set whose Hasse diagram is a tree (in the graph-theoretic sense of the term) having a unique minimal element called the *root*, see Figure 1a. If $|\tau| = n$, a *natural labeling* of τ is a bijection $T: \tau \rightarrow \{1, 2, \dots, n\}$ such that $v < w$ in τ implies $T(v) < T(w)$. One such labeling is given in Figure 1b. In this case, we say T has *shape* τ . We let f_τ represent the number of natural labelings of τ .

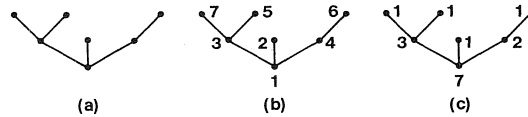
The *hook* of a node $v \in \tau$ is

$$H_v = \{w \in \tau \mid w \geq v\}$$

with corresponding *hooklength* $h_v = |H_v|$. The hooklengths of our example tree are displayed in Figure 1c. The well-known hook formula [3] for the number of natural labelings states that

$$f_\tau = n! / \prod_{v \in \tau} h_v. \tag{1.1}$$

Thus, in our example $f_\tau = 7! / (7)(3)(2)(1)^4 = 120$.



A tree, a labeling and the hooklengths

FIGURE 1

In Section 2 we will give a simple probabilistic proof of (1.1) inspired by an algorithm of Greene, Nijenhuis, and Wilf [1] for standard Young tableaux. The tree version has previously appeared in [5], but is included here for completeness. An algorithmic derivation of the hook-generating function for reverse tree partitions [which specializes to (1.1) as the variable approaches 1] can be found in [6].

A *Fibonacci tree* [9] is a finite lower-order ideal of the infinite poset in Figure 2a. The name derives from the easily proved fact that the number of Fibonacci trees with n nodes is the n^{th} Fibonacci number. For example, Figure 2b shows the five Fibonacci trees with four nodes. Let \mathcal{F}_n be the set of all Fibonacci trees with n nodes, then

$$\sum_{\tau \in \mathcal{F}_n} f_{\tau}^2 = n! \tag{1.2}$$

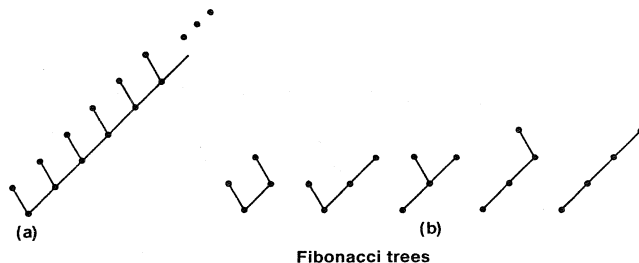


FIGURE 2

Formula (1.2) has a bijective proof due to Bender (reported in [9]). In Sections 3 and 4 below we will give two constructions that build a labeled tree $\tau \in \mathcal{F}_n$ with probability $f_{\tau}^2/n!$, thus proving (1.2) twice. The first algorithm constructs the tree "from without" as done for tableaux in another paper of Greene et al. [2]. The second builds the tree "from within" and is based on work of Pittel [4].

2. Choosing a Labeling Uniformly

Let τ be a fixed shape with n nodes. The following algorithm can be used to choose a labeling of τ .

GNW1. Pick a node $v \in \tau$ uniformly at random, i.e., with probability $1/n$.

GNW2. If v is maximal (a leaf), then let $T(v) = n$ and return to GNW1 with τ and n replaced by $\tau - \{v\}$ and $n - 1$, respectively (unless there are no nodes left, in which case the algorithm halts).

GNW3. If v is not maximal, then choose a different node $w \in H_v$ uniformly at random, i.e., with probability $1/(h_v - 1)$, and return to GNW2 with w in the role of v .

A sequence of nodes generated in the process of finding a vertex to be labeled (in this case by the loop between GNW2 and GNW3) is called a *trial*. An example of a typical trial is given in Figure 3.

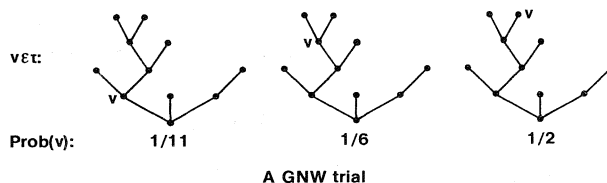


FIGURE 3

Theorem 1: If τ is a fixed rooted tree with n nodes, then GNW1-3 produce all labelings of τ uniformly at random. In fact, the probability of any given labeling is

$$\prod_{v \in \tau} h_v/n!$$

Proof: Let w be any maximal element of τ and let W be the set of vertices on the unique path from w to the root of τ (excluding w itself). Note that these

are the only vertices whose hooklengths are changed if w is removed from τ during GNW2. Therefore, by induction, it suffices to show that the probability that w gets label n is

$$\begin{aligned}
 P(w) &= (1/n) \prod_{v \in W} h_v / (h_v - 1) \\
 &= (1/n) \prod_{v \in W} \left(1 + \frac{1}{h_v - 1} \right).
 \end{aligned}$$

But $1/n$ is the probability of choosing an initial node and each term in the expansion of the product corresponds to the probability of a unique trial ending in w . \square

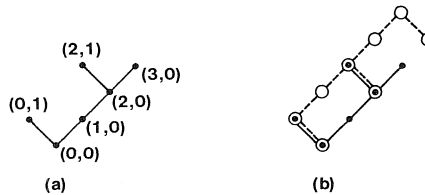
As an immediate corollary we have

Corollary 2: The number of labelings of a given tree τ with n nodes is

$$f_\tau = n! / \prod_{v \in \tau} h_v. \quad \square$$

3. Fibonacci Trees Grown from Without

It will be convenient to introduce coordinates for the infinite tree of Figure 2a. Let the nodes of the "spine" be $(i, 0)$ for $i = 0, 1, 2, \dots$ while the leaves are denoted by $(i, 1)$ for the same range of i . Now, any Fibonacci tree can be specified by its coordinates as is done in Figure 4a.



Coordinates and the associated tree

FIGURE 4

Given any vertex $v = (i, j)$, then v has *associate* $v' = (i, 1 - j)$. If τ is a Fibonacci tree with spine of length s , then the *associated tree* is

$$\tau' = \{v = (i, j) \mid v' \in \tau \text{ or } i = s + 1\};$$

see Figure 4b where the associated tree's nodes are the open circles. Note that τ' is "upside down" with root $r = (s + 1, 1)$.

Now suppose we wish to build a labeled Fibonacci tree, T . Assume that the first $m - 1$ vertices of T have already been constructed and given the labels $1, \dots, m - 1$. Let τ be the current shape of T with associate τ' whose root is r . To add a node labeled m to T we proceed as follows:

WNG1. Choose a $v \in \tau' - \{r\}$ uniformly at random.

WNG2. If $v \notin \tau$, then add v to τ with label m and halt.

WNG3. If $v \in \tau$, say $v = (i, j)$, then return to WNG1 with τ' replaced by $\tau' - \{(i', j') \mid i' \leq i\}$.

Figure 5 presents an example of a trial generated by WGN1-3.

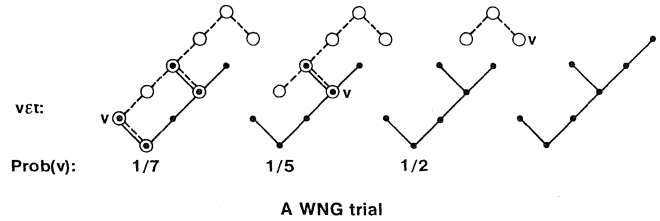


FIGURE 5

If this procedure is used iteratively for $m = 1, 2, \dots, n$ to produce a labeled Fibonacci tree, then let $P(T)$ be the probability that labeling T is created. Thus, the total probability of producing a given shape τ is $P(\tau) = \sum P(T)$, where the sum is over all labelings T of τ .

Theorem 3: If τ is a Fibonacci shape with n nodes, then iteration of WNG1-3 produces all labelings of τ with total probability

$$P(\tau) = f_{\tau}^2/n!$$

Note: It is not true that WNG1-3 produces each labeling of τ with probability $P(T) = f_{\tau}/n!$.

Proof: Let τ have leaves w_1, w_2, \dots, w_k and define the subtrees $\tau_i = \tau - \{w_i\}$ for all i . Let $P(w_i|\tau_i)$ denote the probability that w_i gets labeled n after the algorithm constructs some labeling of τ_i . Hence, by the definitions above and induction,

$$P(\tau) = \sum_i P(\tau_i)P(w_i|\tau_i) = \sum_i (f_{\tau_i}^2/(n-1)!)P(w_i|\tau_i). \quad (3.1)$$

Let the w_i be arranged in order of increasing first coordinate, i.e.,

$$w_1 = (a_1, 1), \dots, w_{k-1} = (a_{k-1}, 1), w_k = (a_k, j),$$

where $a_1 < \dots < a_k$ and j may be 0 or 1. We need a couple of lemmas to help compute the quantities in (3.1).

Lemma 4: Let τ and the w_i be as above, then

$$f_{\tau} = \prod_{i=1}^{k-1} (n - a_i - i).$$

Proof: Using the hook formula (Corollary 2), we see that every term in the $n!$ is canceled by a hook of τ except those in the product above. \square

Lemma 5: Let τ and the w_i be as above, then

$$P(w_i|\tau_i) = (1/n) \prod_{j=1}^{i-1} \left(1 + \frac{2}{n - a_j - j - 1}\right).$$

Proof: Initially we can pick any one of the n nodes in $\tau_i' - \{n\}$. Any trial ending at w_i can only pass through those w_j with $j < i$ and their associates w_j' . Landing on either of these two reduces the number of available nodes in $\tau_i' - \{n\}$ to $n - a_j - j - 1$, accounting for the second term of the binomial above. \square

For notational convenience, let $b_i = n - a_i - i$. Hence, by Lemma 4,

$$f_\tau = b_1 b_2 \dots b_{k-1}$$

and

$$f_{\tau_i} = (b_1 - 1) \dots (b_{i-1} - 1) b_{i+1} \dots b_{k-1}.$$

Also, from Lemma 5,

$$P(w_i | \tau_i) = (1/n) \left(1 + \frac{2}{b_1 - 1}\right) \dots \left(1 + \frac{2}{b_{i-1} - 1}\right).$$

Thus,

$$f_{\tau_i}^2 P(w_i | \tau_i) / (n - 1)! = (1/n!) \left\{ \prod_{1 \leq j < i} (b_j^2 - 1) \right\} \left\{ \prod_{i < j < k} b_j^2 \right\}.$$

Plugging this expression into (3.1), we see that the sum of products telescopes (from the right-hand end) so that

$$P(t) = b_1^2 \dots b_{k-1}^2 / n! = f_\tau^2 / n!$$

as desired. \square

The obvious corollary is

Corollary 6: $\sum_{\tau \in \mathcal{T}_n} f_\tau^2 = n! \quad \square$

We should also note that this algorithm has a "zone effect" similar to the original one for Young tableaux. Specifically, if $v = (a, 1)$ and $w = (b, 1)$ with $a_i < a$, $b < a_{i+1}$, then by Lemma 5 we have $P(v | \tau) = P(w | \tau)$. This observation will be useful in the next section.

4. Fibonacci Trees Grown from Within

Given $v \in \tau$, then v is a *singleton* if $v' \notin \tau$ and a *doubleton* otherwise. In Figure 6a, the singletons are $(0, 0)$, $(3, 0)$, $(4, 0)$, and $(6, 0)$, with the rest of the vertices being doubletons. If τ has a spine of length s , then the corresponding *extended tree* is

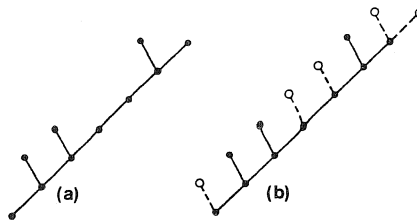
$$\tau'' = \tau \cup \{v' | v \in \tau \text{ is a singleton}\} \cup \{(s + 1, 0)\},$$

see Figure 6b. The elements of $\tau'' - \tau$ are organized into *zones*, which are maximal strings of vertices with consecutive first coordinates. Zones are numbered from the bottom up starting with zone 0, e.g., in Figure 6,

$$Z_0 = \{(0, 1)\}, Z_1 = \{(3, 1), (4, 1)\}, Z_2 = \{(6, 1), (7, 0)\}.$$

In the same way, the doubletons of τ are grouped into *bands* with band i directly below zone i . In our example, the bands are

$$B_0 = \emptyset, B_1 = \{(1, 0), (1, 1), (2, 0), (2, 1)\}, B_2 = \{(5, 0), (5, 1)\}.$$



A tree and the extended tree

FIGURE 6

Finally, it will be convenient to have a total order on the vertices. If $v = (i, j)$ and $w = (x, y)$, then we will write $v \leq_t w$ if $i < x$ or $i = x$ and $j \leq y$.

Now, given a labeled Fibonacci tree T of shape τ on $m - 1$ nodes, we find a node of $w \in \tau'' - \tau$ to label m by constructing a trial as follows. As usual, " := " is the Pascal assignment symbol.

P1. Let $v := (0, 0)$ with probability 1. Let the set of predecessors of v be $P := \emptyset$.

P2. Set $P := P \cup \{v\}$.

P3. Pick w uniformly at random from among the set, D , of possible direct successors of $v = (i, j)$ defined by:

(a) if v is a doubleton, then $D = \{w \in \tau'' - P \mid w \geq_t v\}$.

(b) if v is a singleton, then let B be the band of largest index containing an element of P and let b be the maximum node of B (with respect to \leq_t). In this case

$$D = \{w \in \tau'' - P \mid w >_t b\} - \{w \text{ a singleton} \mid w \leq v\}.$$

If B does not exist, i.e., P consists only of singletons up to this point, then we take $b = (0, 0)$.

P4. If $w \in \tau'' - \tau$, then halt, else return to P2 with $w := v$.

Note that the trials generated by P1-4 do not necessarily respect the partial order in τ and the sequence of D 's computed in P3 is not ordered by containment. For example, if a trial in the tree of Figure 6 has begun $(0, 0)$, $(4, 0)$, then the next node could be any one in τ'' except the two initial nodes and $(3, 0)$. If the trial continues to $(1, 1)$, then any nontrial vertex (i, j) with $i > 1$ is available for the next choice, including $(3, 0)$. However, if the trial begins $(0, 0)$, $(1, 1)$, $(4, 0)$, then the only possible successors are vertices $(3, 1)$, $(4, 1)$, $(5, 0)$, $(5, 1)$, $(6, 0)$, $(6, 1)$, and $(7, 0)$.

Nevertheless, these rules do provide the desired distribution.

Theorem 7: If τ is a Fibonacci shape with n nodes, then iteration of P1-4 produces all labelings of τ with total probability

$$P(\tau) = f_\tau^2/n!$$

Proof: It suffices to show that Lemma 5 is still true when using P1-4. It will be convenient to reformulate the Lemma slightly for this setting. Let λ be a Fibonacci tree with $n - 1$ nodes and leaves w_1, w_2, \dots with first coordinates $a_1 < a_2 < \dots$.

Lemma 8: With λ as above and $w \in \lambda'' - \lambda$ in the k^{th} zone, then the probability of terminating a P1-4 trial at w is

$$P(w) = (1/n) \prod_{w_j \in B_i, i \leq k} \left(1 + \frac{2}{n - a_j - j - 1}\right).$$

Proof: Induct on k . We will provide an explicit proof of the induction step, the anchor step being similar.

The trials t : $v_0 = (0, 0)$, v_1, \dots, w are of two types, those that pass through an element of B_k and those that do not. The latter are in bijective probability preserving correspondence with trials v_0, v_1, \dots, w' , where $w' \in Z_{k-1}$. In the former case, if $v_j \in B_k$ is the first such node then v_0, \dots, v_{j-1}, w' is a legal trial having the same probability as the initial segment of t . We will show below that the sum of the probabilities P of all possible final segments v_j, v_{j+1}, \dots, w is independent of both the particular node of B_k and the

initial history of t . Thus, by induction, it suffices to demonstrate that

$$1 + P|B_k| = \prod_{w_j \in B_k} \left(1 + \frac{2}{n - a_j - j - 1}\right).$$

But the right side above telescopes to $(s + |B_k|)/s$, where s is the denominator corresponding to the largest leaf in B_k that has coordinates $(a, 1)$, say. It is easy to see that if we consider the subtree $\sigma = \{(i, j) \in \lambda \mid i > a\}$ then $s = |\sigma| + 1$. Hence, to finish the proof of the theorem, we need only show

Lemma 9: Let $t, v = v_j, P$, and σ be as above. Then P is independent of the set of nodes on t prior to v and of v itself (as long as $v \in B_k$). In fact, $P = 1/(|\sigma| + 1)$.

Proof: Let $\{v = u_1 \leq_t u_2 \leq_t \dots \leq_t u_m\}$ be the set of all possible vertices that could appear on t from v up to (but not including) w , i.e., the set of all elements above v that are either elements of B_k or singletons not previously on t . Because of these restrictions, the set of direct successors, $D(u_i)$, does not depend on the previous u_j chosen and, in fact, we have

$$\begin{aligned} D(u_i) &= \{u_j \mid j > i\} \cup \{v \in \sigma'' \mid v \text{ is not a singleton in } \sigma\} \\ &= D(u_{i-1}) - \{u_i\}. \end{aligned}$$

Thus,

$$|D(u_m)| = |\{v \in \sigma'' \mid v \text{ is not a singleton in } \sigma\}| = |\sigma| + 1$$

and

$$|D(u_i)| = |D(u_{i-1})| - 1.$$

Hence,

$$P = \frac{1}{|D(u_1)|} \left(1 + \frac{1}{|D(u_1)| - 1}\right) \dots \left(1 + \frac{1}{|\sigma| + 1}\right) = \frac{1}{|\sigma| + 1}$$

as desired. \square

Of course, Theorem 7 gives another proof of Corollary 6.

5. Remarks and Open Questions

Another point of similarity between Fibonacci trees and standard tableaux is the formula

$$\sum_{\tau \in \mathcal{F}_n} f_\tau = I_n, \tag{5.1}$$

where I_n is the number of involutions in the symmetric group S_n . The correspondence of Bender [9] mentioned in the introduction also proves (5.1). Is there a probabilistic way to demonstrate this, either for trees or tableaux?

A third family of posets that displays behavior similar to that of standard tableaux and rooted trees are the shifted standard tableaux [3]. The shifted analog of the hook formula (1.1) has been proved probabilistically by one of us [7]. It would be interesting to find an aleatory proof of the "sum of squares" equation in the shifted case (see [8] for the exact formula).

Finally, tableaux and shifted tableaux are intimately connected with representations of S_n . Ordinary tableaux give the degrees of ordinary irreducible representations (using matrices in GL_n), while their shifted cousins are related to projective ones (those using PGL_n , the projective linear group). In this setting, the analog of (1.2) expresses the fact that the sum of the

squares of the irreducible degrees equals the order of the group. Can (1.2) itself be recast in this light? Specifically, is there a group of matrices G such that the degrees of the irreducible representations $\rho: S_n \rightarrow G$ are given by the f_τ ?

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Announcement

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