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# Finite automata for testing composition-based reconstructibility of sequences

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## Abstract

Symbolic sequences uniquely reconstructible from all their substrings of length  $k$  compose a regular factorial language. We thoroughly characterize this language by its minimal forbidden words, and explicitly build up a deterministic finite automaton that accepts it. This provides an efficient on-line algorithm for testing the unique reconstructibility of the sequences.

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## 1. Introduction

The problem of sequence reconstruction from composition has been raised in various contexts, like high throughput DNA sequencing [1], composition-based prokaryotic phylogenetics [10], and string embedding [6]. It considers whether a symbolic sequence can be uniquely recovered from the multiset of all its constituent “ $k$ -tuples.” For example, the oligonucleotide sequences TACTAGACT and TAGACTACT have the same triple composition {ACT, ACT, AGA, CTA, GAC, TAC, TAG}, thus neither is uniquely reconstructible. Given a  $k$ -tuple composition, there exists a linear-time algorithm to determine whether a conforming sequence exists, and if yes it constructs one [2,8]. The number of sequences with a valid composition is given by the “modified BEST formula” [4,5,10], but the calculation can be tough, because the formula is based on the matrix-tree theorem in graph theory, which involves a determinant whose size is, in cases of interest, comparable with the length of the sequence. Alternatively, the set of uniquely reconstructible sequences can be investigated as a formal language. Recently, Kontorovich [6] proved that this language is regular, and conjectured that a finite automaton that accepts it can be efficiently constructed. The present paper supplies a different proof based on the results of Ukkonen [11] and Pevzner [9], and further characterizes this language by its minimal forbidden words. Finally we explicitly build up the associated deterministic finite automaton (DFA), which provides an efficient on-line algorithm for testing the uniqueness of reconstructions of sequences. We have implemented it in a C++ program, and the source code is available on request.

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## 2. Conventions and notation

We start by fixing some notation. The empty string will be denoted by  $\epsilon$ . By convention, we denote by  $\Sigma$  the alphabet in consideration,  $\Sigma^*$  the set of all finite strings over  $\Sigma$ , and  $\Sigma^+ = \Sigma^* \setminus \{\epsilon\}$  the nonempty strings. For the sake of convenience, we lay down the rule that low Latin letters  $a, b, c, d$  denote characters, and high letters  $r, s, \dots, z$  denote strings. We denote by  $|s|$  the length of the string  $s$ , and  $|s|_a$  the number of the character  $a$ 's in  $s$ . The set of all characters occurring in  $s$ , i.e.,  $\{a \in \Sigma: |s|_a \geq 1\}$ , will be denoted by  $\text{alph}(s)$ . For a string  $s = at$ , the character  $a$  is said to be the *head* of  $s$ , denoted by  $\text{head}(s)$ . Similarly,  $b$  is said to be the *tail* of the string  $s = tb$ , denoted by  $\text{tail}(s)$ . For  $s = uvw$ ,  $v$  is said to be a *factor* of  $s$ , and it is called a *left factor* if  $u = \epsilon$ . A factor  $v$  of  $s$  is said to be *proper* if  $v \neq s$ . For unexplained terms in formal language and automata theory, we refer the reader to the standard textbook [3].

Without losing generality, we only consider the problem of duple composition, i.e.,  $k = 2$ . For  $k > 2$ , it can be easily reduced to the former case by considering the set of  $(k - 1)$ -tuples,  $\Sigma^{k-1}$ , as the alphabet.

## 3. The complementary language

Ukkonen [11] conjectured and Pevzner [9] proved that any two sequences with the same composition can be transformed into each other by a series of operations called *rotations* and *transpositions*. A rotation

$$R: aubva \rightarrow bvaub$$

applies to a string whose head and tail are the same. This case is simple, and can be eliminated by preceding each string with a special character outside  $\Sigma$ . We will ignore it in the following, such that the frequencies of characters are also conserved, as is usually required in practice. A transposition

$$T: uaxbwaybv \rightarrow uaybwaxbv$$

exchanges a pair of nonoverlapping factors of a string ( $x$  and  $y$ ), given that they are flanked by the same character on either side (respectively  $a$  and  $b$ ). In case  $a = b$  it has a degenerated form

$$T: uaxayav \rightarrow uayaxav.$$

Clearly, these operations do not alter the composition. We can unify them into the form

$$T: uxywv \rightarrow ywxv, \tag{1}$$

where

$$\text{tail}(u) = \text{tail}(w) \quad \text{and} \quad \text{head}(w) = \text{head}(v). \tag{2}$$

We denote by  $L$  the language of uniquely reconstructible sequences, and  $L'$  its complement, then a string  $s$  is in  $L'$  only if it has a form  $s = uxywv$  subject to condition (2). However, this condition is not sufficient even in the constraint  $x \neq y$ . For example, the string 010101 has such a form with  $(u, x, w, y, v) = (0, 10, 10, \epsilon, 1)$  while it is in  $L$ .

**Remark 1.** Generally, the string  $s$  is invariant under the transposition if and only if  $xyw = ywx$ , in other words  $xywyw = ywxw$ . It follows from Proposition 1.3.2 in [7] that the two words  $xw$  and  $yw$  commute if and only if they are the powers of the same word. Therefore we can write

$$x = (rt)^l r, \quad w = t(rt)^m, \quad y = (rt)^n r, \quad l, m, n = 0, 1, \dots$$

To rule out transpositions on strings in  $L$ , we add to the conditions in (2) that

$$\text{head}(xw) \neq \text{head}(yv), \tag{3}$$

while every string in  $L'$  still retains a transposition. This is justified by a slightly stronger proposition:

**Proposition 2.** For any transposition  $T: s \rightarrow s'$  with  $s \neq s'$ , it can be written in the form (1), such that  $u$  is the longest common left factor of  $s$  and  $s'$ .

**Proof.** Suppose  $s$  and  $s'$  have a common left factor  $u' = ua$ , then  $\text{head}(yw) = a$ . We show that the transposition can be written in the form  $T' : u'x'w'y'v \rightarrow u'y'w'x'v$ . There are three cases:

- (1) If  $x \neq \epsilon$  and  $y \neq \epsilon$ , then we can write  $x = ax'$  and  $y = ay'$ , and let  $w' = wa$ .
- (2) If  $x = \epsilon$ , then we can write  $w = ax'$  and  $y = ay'$ , and let  $w' = a$ .
- (3) If  $y = \epsilon$ , then we can write  $x = ax'$  and  $w = ay'$ , and let  $w' = a$ .

It turn out that  $u'y'w'x'v = s'$ .

By repeating this procedure we can extend  $u$  until the given condition holds.  $\square$

As an immediate corollary,

$$L' = \{s = uxwyv : \text{conditions (2) and (3) hold}\}.$$

Each subset  $L'_{abc}$  of  $L'$ , defined by  $\text{tail}(u) = \text{tail}(w) = a$ ,  $\text{head}(w) = \text{head}(v) = b$ ,  $\text{head}(xw) = c$  and  $\text{head}(yv) \neq c$ , clearly constitutes a regular language. Therefore, as regular languages are closed under union and complementation [3, p. 59],  $L'$  and  $L$  must also be regular languages. This is the main theorem in [6]. Moreover, we obtain a right-linear grammar for  $L'$  composed of productions of the following forms:

$$\begin{aligned} U &\rightarrow dU \mid acZ_{acc} \mid aaA_{aa}, \\ Z_{acc} &\rightarrow bZ_{acb}, \\ Z_{acb} &\rightarrow dZ_{acb} \mid aA_{cb}, \\ A_{cb} &\rightarrow dY_b \ (d \neq c) \mid bV \ (b \neq c), \\ Y_b &\rightarrow dY_b \mid bV, \\ V &\rightarrow dV \mid \epsilon, \end{aligned}$$

where  $U$  is the start symbol, and  $a, b, c, d$  run over  $\Sigma$ . It is of little interest to present the routine (but a bit lengthy) proof, instead we note that for given  $a, b$ , and  $c$ , this grammar generates  $L'_{abc}$ .

#### 4. Minimal forbidden words

A language is said to be *factorial* (or *factorizable*) if it contains all factors of its members. Clearly  $L$  is factorial. A factorial language can be determined by its *minimal forbidden words* (MFWs, also known as distinct excluded blocks, or DEBs) [13]. A MFW of a language is a string that does not belong to the language while all its proper factors do. They help to understand the structure of the language.

**Theorem 3.** *A string  $r$  is a MFW of  $L$  if and only if  $r = axwyb$ , such that*

- (1)  $\text{head}(w) = b$  and  $\text{tail}(w) = a$ ;
- (2)  $x \neq \epsilon$  or  $y \neq \epsilon$ ;
- (3)  $x, w, y \in L$ ;
- (4)  $\text{alph}(x)$ ,  $\text{alph}(w)$ , and  $\text{alph}(y)$  are mutually disjoint;
- (5)  $|w|_a = 1$  and  $|w|_b = 1$ .

**Proof.** The sufficiency is trivial. And the necessity of the first three conditions is evident, so we only need to justify the last two conditions.

Suppose  $y$  contains a character  $b'$  which occurs in  $xw$ , then  $y$  has a left factor  $y'b'$ , and we can write  $xw = x'w'$ , with  $\text{head}(w') = b'$ . It is the case that  $r$  has a left factor  $r' = ax'w'y'b' \in L'$ , which contradicts that  $r$  is a MFW. Therefore,  $\text{alph}(xw) \cap \text{alph}(y) = \emptyset$ . Clearly, reversing a string does not alter its membership of  $L$ , i.e.,  $L$  is reversal. So similarly we have  $\text{alph}(x) \cap \text{alph}(w) = \emptyset$ . Hence condition (4) holds.

Suppose  $|w|_a > 1$ , then we can write  $w = zay'a$ . Let  $x' = xz$ , then  $axw = ax'ay'a$ .

If  $x \neq \epsilon$ , by  $\text{alph}(x) \cap \text{alph}(w) = \emptyset$  we have  $\text{head}(x'a) \neq \text{head}(y'a)$ . It follows  $ax'ay'a \in L'$ , which contradicts that  $r$  is a MFW. Therefore  $|w|_a = 1$ . Similarly  $|w|_b = 1$  if  $y \neq \epsilon$ .

If  $x = \epsilon$ , then  $\text{head}(x'a) = \text{head}(w) = b$ . It follows from condition (2) that  $y \neq \epsilon$ , hence  $|w|_b = 1$ , and  $\text{head}(y'a) \neq b$ . Again it follows  $ax'ay'a \in L'$  and results in a contradiction. Therefore  $|w|_a = 1$ . Similarly  $|w|_b = 1$  if  $y = \epsilon$ .  $\square$

We can enumerate the MFWs of  $L$  by recursion on  $|\Sigma|$ . For the simplest nontrivial case, say  $\Sigma = \{0, 1\}$ , the MFWs can be represented by a regular expression  $001^+0 + 01^+00 + 110^+1 + 10^+11$ .

### 5. The finite automata

Technically we can construct the finite automaton that accepts  $L$  from the grammar of  $L'$  or the MFWs of  $L$ , but it is more convenient to design it directly as follows.

**Input alphabet:** Without losing generality, we let

$$\Sigma = \{1, 2, \dots, m\}.$$

**States:**

$$Q = P \times N \times C,$$

where

$$P = \Sigma \cup \{0\}, \quad N = (\Sigma \cup \{\epsilon\})^{m+1}, \quad \text{and} \quad C = \{\text{WHITE}, \text{BLACK}\}^m.$$

**Initial state:**

$$q_0 = (0, \epsilon^{m+1}, \text{WHITE}^m).$$

**Final states:**

$$F = \{(p, n, c) \in Q: c \neq \text{BLACK}^m\}.$$

**Transition function:**  $\delta: Q \times \Sigma \rightarrow Q$  is defined by the following algorithm:

```

1   $\delta((p, n, c), a)$ 
2  if  $n_p \neq \epsilon$  and  $n_p \neq a$ 
3    then  $i \leftarrow p$ 
4    repeat
5       $c_i \leftarrow \text{BLACK}$ 
6       $i \leftarrow n_b$ 
7    until  $i = p$ 
8  if  $c_a = \text{BLACK}$ 
9    then  $c \leftarrow \text{BLACK}^m$ 
10  $n_p \leftarrow a, p \leftarrow a$ 

```

**Theorem 4.** The DFA  $M = (Q, \Sigma, \delta, q_0, F)$  accepts  $L$ .

Before stating the formal proof, we roughly describe the function of each component of the state variable. We use  $p$  to register the last read character, and every input string is preceded with a special character  $p_0 = 0$ . The vector  $n$  implements a singly linked data structure, where an element  $n_a$  gives the character following the most recent occurrence of  $a$ . This implies that a simple linear search in  $n$  can always reach  $p$ , thus the loop on lines 2–6 of the algorithm never falls infinite. The vector  $c$  attributes a “color” to every character, initially all are WHITE. If  $b$  occurs in a factor  $aza$  of the input string, with two  $a$ ’s followed by distinct characters, then  $c_b$  turns BLACK through the loop. Such a factor  $aza$ , with the follower different from  $\text{head}(za)$ , will be called a *bead*. If  $b$  occurs after this bead, the string will be in  $L'$ , thus a character colored BLACK will be forbidden.

**Proof.** We prove it by induction on the length  $l$  of the input, along with an auxiliary proposition: After reading any string  $s \in L$ ,  $c_a = \text{BLACK}$  if and only if  $a$  occurs in a bead.

The basis is evident. Suppose for any string  $s$  of length  $k$  the proposition holds. For any  $t$  of length  $k + 1$  we write  $t = sb$ .

If  $s \notin L$ , then  $t \notin L$ , and by the inductive hypothesis  $c = \text{BLACK}^m$  before  $b$  is read. According to the transition function  $c$  will remain  $\text{BLACK}^m$ , thus  $t$  will be rejected by  $M$ .

If  $s \in L$ , then  $c \neq \text{BLACK}^m$  before  $b$  is read. If the condition on line 1 holds, then  $t$  has a bead  $pzp$ . The loop body starts a walk from  $p$ , and lets  $c_d = \text{BLACK}$  for every character  $d$  visited. If  $d$  in the bead is not reached, then by the rule for assignment of  $n$  on the last line of the algorithm, it must be in another bead  $axa$  with  $a \neq p$ , and by the inductive hypothesis  $c_d = \text{BLACK}$  already. Therefore, when line 7 of the algorithm is reached, the condition holds if and only if  $b$  has occurred in a bead, i.e.,  $c$  will get  $\text{BLACK}^m$  if and only if  $t \in L'$ .  $\square$

## 6. Discussion

As pointed out in [8], the sequence reconstruction problem in consideration is equivalent to the problem of uniqueness of Eulerian trail in a directed pseudo-graph, since it can be naturally represented by a sequence over the set of vertices  $V = \Sigma^{k-1}$ . For a graph with an Eulerian trail  $t = sb$ , the state variable  $n$  represents a spanning tree towards  $b$ , with edges  $\{(a, n_a) : a \in V, a \neq b\}$ .

Under reasonable assumptions, the time complexity of the present algorithm is linear for fixed  $\Sigma$  and  $k$ . Since the number of  $(k - 1)$ -tuples occurred in the sequence is usually small relative to the total number of possible ones, the state variables can be stored in a dynamic data structure to save space. For example, it can be implemented as a hash table, so that the expected running time is still linear. Furthermore, the algorithm is on-line, and can halt on the first occurrence of a forbidden word. Utilizing it, investigation on real biological sequences [10] and preliminary numerical experiments [12] have revealed some interesting features of the distribution of probability of uniqueness of sequence reconstruction with respect to  $k$ .

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