

# A New Dynamic Algorithm for the Dyck Language

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**Abstract.** We study Dynamic Membership problems for the two-sides Dyck language, the class of strings of properly balanced parentheses. We present a non-deterministic algorithm and a data structure such that together they solve the following problems: replacements of symbols, insertions, deletions of symbols, and language membership queries.

**Key words:** Dyck Language, balanced braces sequence, dynamic algorithm, persistent structure.

## 1 Introduction

### 1.1 The Dyck Language

The Dyck Language is a language describing balanced bracket sequences. Let  $A = \{a_1, a_2, \dots, a_k\}$  and  $\bar{A} = \{\bar{a}_1, \bar{a}_2, \dots, \bar{a}_k\}$  be two disjoint sets of opening and closing symbols, respectively. For example, the pair  $A = \{([, do, if\}$  and  $\bar{A} = \{), ], od, fi\}$  captures the nested structure of programming languages. The two-sided Dyck language  $D'_k$  over  $A \cup \bar{A}$  is the context-free language generated by the following grammar:

- $\epsilon \in D'_k$
- If  $S \in D'_k$  then  $\forall i : a_i S \bar{a}_i \in S$
- If  $S_1, S_2 \in D'_k$  then  $S_1 S_2 \in D'_k$

### 1.2 The Dynamic Membership Problem

In this paper we consider the problem of maintaining membership in  $D'_k$  of a string from  $(A \cup \bar{A})^n$  dynamically. More precisely, we want to implement a data structure  $\mathcal{D}$  containing a string  $x \in (A \cup \bar{A})^n$  of even length, with the following behaviour:

1. At creation,  $\mathcal{D}$  is initialized by the empty string.
2. The following operations are supported by  $\mathcal{D}$ :
  - member**: return 'yes' if and only if  $x \in D'_k$
  - create**( $\sigma$ ): create a new data structure  $\mathcal{D}$  with  $x = \sigma$
  - split**( $i$ ): split string  $x$  into two strings  $x' = x_1 x_2 \dots x_i$  and  $x'' = x_{i+1} x_{i+2} \dots x_n$
  - merge**( $x', x''$ ): concatenate two strings  $x'$  and  $x''$  into one  $x = x' x''$

### 1.3 Results

Our main model of computation will be a unit-cost random access machine with word-size  $O(\log n)$ , where  $n$  is the size of the input; this model is also known as a random access computer. We present a new data structure solving the Dynamic Membership Problem. Each query is completed in  $O(\log n \log h)$  time, where  $h$  is the maximum nesting depth of brackets in the string and  $O(n \log n)$  memory. In the worst case our result is equal to the one by Frandsen et al. [1].

## 2 Algorithm

### 2.1 A Data Structure for String Equality Verification

We create a non-deterministic structure  $\mathcal{S}$  that handles the following operations for strings over a finite alphabet  $\Sigma$ :

**create**( $\sigma$ ): create a new string  $s = \sigma$  in time  $O(1)$   
**split**( $s, i$ ): create two new strings  $s' = s_1 s_2 \dots s_i, s'' = s_{i+1} s_{i+2} \dots s_n$  in time  $O(\log n)$   
**merge**( $s', s''$ ): create new string  $s = s' s''$  in time  $O(\log(|s' s''|))$   
**equal**( $s', s''$ ): return 'yes' if and only if  $s' = s''$  in time  $O(1)$   
**copy**( $s$ ): create a new copy of string  $s$  in time  $O(1)$

We can use a implicit persistent treap [2, 3] (or other self-balancing binary search trees [4] with implicit keys) with polynomial hashing [5] to comply with the all requirements. Let each  $v$ -th vertex contains a polynomial hash of  $s_{l\dots r}$  if it represents substring  $s_{l\dots r}$ . This simple idea allows you to implement  $\mathcal{S}$ .

### 2.2 Theoretical Background

Let us define inverse string  $s^{-1}$  to be  $\bar{s}_n \bar{s}_{n-1} \dots \bar{s}_1$  with the convention  $\bar{\bar{a}}_i = a_i$  and  $\bar{\epsilon} = \epsilon$ . We say that a string is *reduced* if it contains no neighbouring pair of matching parentheses. Let us introduce a one-sided associative calculus by alphabet  $\Sigma = A \cup \bar{A}$  with relations of matching parentheses:  $\forall i : a_i \bar{a}_i \rightarrow \epsilon$ .

Let  $\mu$  be a map from  $\Sigma^*$  to  $\Sigma^*$  such that  $\mu(s)$  is derived from  $s$  in one-sided associative calculus and  $\mu(s)$  is not reducible.

**Lemma 1.** *The map  $\mu(s)$  is a function. In other words, for any string  $s$  there exists a unique  $\mu(s)$  regardless of the order of reducing actions.*

**Lemma 2.** *Let  $s$  be a substring of string  $t$ . If any  $a_i \bar{a}_j$  is a substring of  $\mu(s)$  then  $t \notin D'_k$ .*

**Lemma 3.** *Suppose that any substring  $s$  of a string  $t$  does not satisfies the hypotheses of Lemma 2; then  $\mu(t) = \bar{a}_{i_1} \bar{a}_{i_2} \dots \bar{a}_{i_k} a_{j_1} a_{j_2} \dots a_{j_m}$ . This means that  $\mu(t)$  consists of a prefix of closing braces and a suffix of opening braces.*

We say that a string is *complementary* if it satisfied the hypotheses of Lemma 3. For any complementary string  $t$  we can denote two functions  $\bar{\phi}(t), \phi(t) : \Sigma^* \rightarrow \Sigma^*$  as follows:

$$\bar{\phi}(t) = \bar{a}_{i_1} \bar{a}_{i_2} \dots \bar{a}_{i_k}, \phi(t) = a_{j_1} a_{j_2} \dots a_{j_m}$$

**Lemma 4.** *Let  $s = xy$  such that  $x, y, s \in \Sigma^*$ . The string  $s$  is complementary if and only if  $x$  and  $y$  is complementary strings, and one of the following conditions:*

1. *If  $\phi(x) = x_1 x_2$ , where  $x_1, x_2 \in \Sigma^*$  and  $(x_1)^{-1} = \bar{\phi}(y)$ , then  $\bar{\phi}(s) = \bar{\phi}(x)$  and  $\phi(s) = x_2 \phi(y)$ .*
2.  *$\bar{\phi}(y) = \bar{y}_1 \bar{y}_2$ , where  $\bar{y}_1, \bar{y}_2 \in \Sigma^*$  and  $(\bar{y}_2)^{-1} = \phi(x)$ , then  $\bar{\phi}(s) = \bar{\phi}(x) \bar{y}_1$  and  $\phi(s) = \phi(y)$ .*

All proof contains in the Frandsen et al. [1] and Harrison [6].

### 2.3 The Main Data Structure

Let  $\mathcal{D}$  be the structure under string  $s$ . We maintain a balanced binary tree in  $\mathcal{D}$  whose  $v$ -th vertex represents substring  $s_{l..r}$  and where each internal node represents the concatenation of its children's strings  $s_{l..m}$  and  $s_{m+1..r}$ . If  $s_{l..r}$  is complementary string, then we will save  $\phi(s_{l..r})$  and  $\bar{\phi}(s_{l..r})$  in  $v$ . In other case, we will save only one bit about uncomplementary.

We can easily compute the vertex data. If substrings  $s_{l..m}$  or  $s_{m+1..r}$  is not complementary, then  $s_{l..r}$  is not complementary by the Lemma 4. After, we need to check two conditions from statement of Lemma 4 using the Data Structure for String Equality Verification  $\mathcal{S}$ . If each of them unsatisfied, then  $s_{l..r}$  is not complementary. In other case, we will compute  $\phi(s_{l..r})$  and  $\bar{\phi}(s_{l..r})$  by the formulas from Lemma 4.

Calculating this information takes  $O(\log |\mu(s_{l..m})| + \log |\mu(s_{m+1..r})|)$  time using our data structure  $\mathcal{S}$ . If the maximum nesting depth of brackets is  $h$ , then updating of one vertex can be completed in  $O(\log h)$  time. Therefore, member, split and merge requests can be completed in time  $O(\log n \log h)$  per operation and  $O(n \log n)$  memory total, because of structure  $\mathcal{S}$  is persistent.

### References

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