

# Finding Maximal Exact Matches in Graph

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# Prerequisites

## BWT Transform: Pseudocode

Assuming our strings' indices are 1-based, we now give a non-efficient algorithm to calculate the BWT transform of a string.

---

```
def BWT(s: str) -> str:
    T = []
    for character in s:
        s = s[len(s)] + s[1..len(s)-1]
        T.push(s)
    sort_lexicographically(T)
    return last_column(T)
```

---



# Prerequisites

## BWT Transform: Example

Transformation				
1. Input	2. All rotations	3. Sort into lexical order	4. Take the last column	5. Output
BANANA\$	BANANA\$ \$BANANA A\$BANAN NA\$BANA ANA\$BAN NANA\$BA ANANA\$B BANANA\$	\$BANANA BANANA\$ A\$BANAN ANA\$BAN ANANA\$B BANANA\$ NA\$BANA NANA\$BA	\$BANANA BANANA\$ A\$BANAN ANA\$BAN ANANA\$B BANANA\$ NA\$BANA NANA\$BA	A\$NNBAA

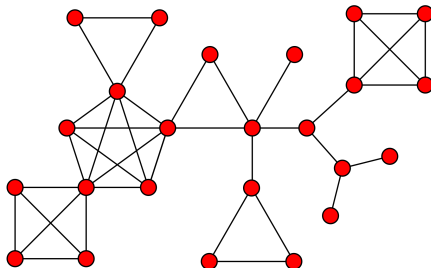


# Prerequisites

## Elastic Founder Graph: Block graph

### Definition (Block Graph)

We call **block graph** an undirected graph in which every biconnected component (i.e. a **block**) is a clique.



### Definition (Elastic Founder Graph)

Consider a block graph  $G = (V, E, I)$  with  $I : V \rightarrow \Sigma^+$ . We call such a graph an **indexable Elastic Founder Graph** if the **semi-repeat-free** property holds: for each  $v$  in block  $V_i$ ,  $I(v)$  occurs in  $G$  only as prefix of paths starting with some  $w \in V_i$ .

# k-MEMs

k-MEMs: LEFTMAX, RIGHTMAX

Let  $Q \in \Sigma^+$  be a query string,  $\kappa$  be a threshold,  $l\text{ext}(i, P, j)$  the left extension of the string  $P[i..j]$  and  $r\text{ext}(i, P, j)$  the right extension.

## Definition (LEFTMAX)

A match  $([x..y], (i, P, j))$  of  $Q[x..y]$  in  $G$  satisfies the *LEFTMAX* property if and only if

$$x = 1 \vee l\text{ext}(i, P, j) = \emptyset \vee Q[x-1] \notin l\text{ext}(i, P, j)$$

We can analogously define the RIGHTMAX property.



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### Definition ( $\kappa$ -MEM)

A match  $([x..y], (i, P, j))$  of  $Q[x..y]$  in  $G$  is called a  $\kappa$ -MEM if it satisfies all the following conditions:

- 1  $LEFTMAX \vee |l_{ext}(i, P, j)| \geq 2$
- 2  $RIGHTMAX \vee |r_{ext}(i, P, j)| \geq 2$
- 3  $y - x + 1 \geq \kappa$

# MEMs in Node Labels

## Node MEMs: definition

### Definition (Node MEM)

A  $\kappa$ -MEM between a query string  $Q$  and the label  $l(v)$  of a vertex  $v \in V$  is called a node-MEM.

To give an algorithm to find node-MEMs we must consider the text

$$T_{\text{nodes}} = \prod_{v \in V} 0 \times l(v).$$

We also need a data structure supporting the following operations over a bitvector  $B$ :

- 1  $r = \text{rank}(B, i)$  in  $O(1)$  with  $r = \sum_{j=1}^i B[j]$ ,
- 2  $j = \text{select}(B, r)$  in  $O(1)$  with  $j \leq i$  the position of the  $r$ -th 1 in  $B$ .



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# MEMs in Node Labels

Algorithm to find k-node MEMs: prerequisites

We assume we have at our disposal the following procedures:

- 1** `mems_using_bidirectional_bwts` which takes as input two bidirectional BWT indices on strings  $T, Q$  and a threshold  $\kappa$  and outputs  $Q'$  MEM strings with  $|Q'| \geq \kappa$  and for each string  $Q'$  four BWT intervals  $([i_T..j_T], [i'_T..j'_T], [i_Q..j_Q], [i'_Q..j'_Q])$  which represent the maximal matches of  $T$  and  $Q$  in  $O(|T| + |Q|)$  time;
- 2** `mems_from_bidirectional_bwt` which takes as input four BWT intervals and outputs the corresponding MEM string  $Q'$  in  $O(|Q'|)$ .

For both of these algorithms, one may refer to Algorithm 11.3 and Algorithm 11.4 from Genome-Scale Algorithm Design: Biological Sequence Analysis in the Era of High-Throughput Sequencing (Belazzougui et alia).



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## Algorithm to find k-node MEMs

We can now describe the algorithm to find node MEMs.

- 1 We build a bitvector  $B$  to mark the location of 0s in  $T_{\text{nodes}}$  so that with the rank operation we can identify the corresponding node in  $G$ .
- 2 We call `mems_using_bidirectional_bwts` using bidirectional BWT indices on strings  $T_{\text{nodes}}$  and  $Q$ .
- 3 For each string  $Q'$  we get, we call `mems_from_bidirectional_bwt` on its corresponding BWT intervals.
- 4 We use  $B$  to obtain the tuple  $(i, P, j)$  in  $O(1)$  time.

The algorithm described above has complexity

$O(|T_{\text{nodes}}| + |Q| + N_{\kappa})$  with  $N_{\kappa}$  the number of output MEMs.



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# k-Node MEMs spanning exactly L nodes

Algorithm to find k-Node MEMs spanning exactly L nodes: prerequisites

- 1 We call  $P_G^L$  a path of G spanning exactly L nodes.
- 2 We define two new symbols  $c$  and  $d$  not originally part of the alphabet  $\Sigma$  and the two operators:

$$left(u) = \begin{cases} c & \text{if } l\text{ext}(u) = \{c\} \\ \# & \text{otherwise} \end{cases}$$

$$right(u) = \begin{cases} d & \text{if } r\text{ext}(u) = \{d\} \\ \# & \text{otherwise} \end{cases}$$

- 3 We define the text

$$T_L = 0 \times \prod_{u_1, \dots, u_L \in P_G^L} \left( left(u_1) \times l(u_1) \times \dots \times l(u_L) \times right(u_L) \times 0 \right).$$



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- 1 We modify the `mems_from_bidirectional_bwt` so that it makes use of the symbols defined before we get the result using the algorithm we defined for  $\kappa$ -node MEMs.
- 2 The complexity we get is  $O(|T_L| + |Q| + M_{\kappa,L})$  with  $M_{\kappa,L}$  the number of output MEMs.
- 3 Let  $d$  be the maximum in-degree (or out-degree) of a node,  $n$  the total label length of  $G$ . Now we can reformulate the time complexity.
- 4  $|T_L|$  is the concatenation of paths of  $G$  of length  $L$ : for a node  $v$  the number of paths containing  $l(v)$  is at most  $L \times d^{L-1}$ . The complexity can be rewritten as  $O(|Q| + M_{\kappa,L} + n \times L \times d^{L-1})$  which is exponential on  $L$ .



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# MEMs in EFGs

## k-Node MEMs in EFGs: Remark

### Remark

*Given an indexable EFG  $G = (V, E, l)$ , for each  $(v, w) \in E$  string  $l(v)l(w)$  occurs only as prefix of paths starting with  $v$ .*

Rephrasing what is written above, all occurrences of some string  $S$  in  $G$  spanning at least four nodes can be decomposed as  $\alpha l(u_2) \dots l(u_{L-1}) \beta$  such that:

- 1  $u_2 \dots u_{L-1}$  is a path in  $G$  and  $u_2, \dots, u_{L-1}$  are unequivocally identified;
- 2  $\alpha = l(u_1)[i..||u_1||]$  with  $1 \leq i \leq ||u_1||$  for some  $(u_1, u_2) \in E$ ;
- 3  $\beta = l(u_L)$  for some  $(u_{L-1}, u_L) \in E$  or  
 $\beta = l(u_L)(l(u_{L+1})[1..j])$  with  $1 \leq j \leq ||u_{L+1}||$  for some  $(u_{L-1}, u_L), (u_L, u_{L+1}) \in E$ .

Note that  $\alpha, \beta \neq \epsilon$  and  $\beta$  has as prefix a full node label,  $\alpha$  might spell any suffix of a node label.



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# MEMs in EFGs

k-Node MEMs in EFGs spanning more than 3 nodes: preprocessing

First, we take the text  $T'_3 = \prod_{(u,v),(v,w) \in E} (I(u)I(v)I(w) \times 0)$ .

- 1 We mark all implicit or explicit nodes  $\bar{p}$  such that the corresponding root-to- $\bar{p}$  path spells  $I(u)I(v)$  for some  $(u, v) \in E$ , so that we can query in constant time if  $\bar{p}$  is such a node.
- 2 We compute pointers from each node  $\bar{p}$  to an arbitrarily chosen leaf in the subtree rooted at  $\bar{p}$ ;
- 3 for each node  $v \in V$  of the indexable EFG we build trie  $T_v$  for the set of strings  $I(\bar{u}) : (u, v) \in E$ ;
- 4 for each leaf, we store the corresponding path  $uvw$  and the starting position of the suffix inside  $I(u)I(v)I(w)$ .



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First, we take the text  $T'_3 = \prod_{(u,v),(v,w) \in E} (I(u)I(v)I(w) \times 0)$ .

- 1 We mark all implicit or explicit nodes  $\bar{p}$  such that the corresponding root-to- $\bar{p}$  path spells  $I(u)I(v)$  for some  $(u, v) \in E$ , so that we can query in constant time if  $\bar{p}$  is such a node.
- 2 We compute pointers from each node  $\bar{p}$  to an arbitrarily chosen leaf in the subtree rooted at  $\bar{p}$ ;
- 3 for each node  $v \in V$  of the indexable EFG we build trie  $T_v$  for the set of strings  $I(\bar{u}) : (u, v) \in E$ ;
- 4 for each leaf, we store the corresponding path  $uvw$  and the starting position of the suffix inside  $I(u)I(v)I(w)$ .



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Let  $\bar{p}$  be the suffix tree node of  $T'_3$  reached from the root by spelling  $Q[1..y]$  in the suffix tree until we cannot continue with  $Q[y+1]$ :

- 1 If we cannot continue with a 0,  $Q[1..y]$  spans no more than 3 nodes, so we can discard it. We can now consider matching  $Q[2..y]$  in  $G$  taking the suffix link of  $\bar{p}$ .
- 2 If we can continue with a 0 and the occurrences of  $Q[1..y]$  span no more than 2 nodes, we proceed as in the previous step.
- 3 In this case,  $Q[1..y] = \alpha l(u_2)l(u_3)$  for exactly one  $u_2 \in V$ , with  $(u_2, u_3) \in E$ , we follow the suffix link walk from  $\bar{p}$  until we find the marked node  $\bar{q}$  corresponding to  $l(u_2)l(u_3)$ : from  $\bar{q}$  we try to match  $Q[y+1..]$  until failure, matching  $Q[y+1..y']$  and reaching node  $\bar{r}$ .



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# MEMs in EFGs

## Algorithm complexity

### Theorem (Algorithm complexity)

*Let alphabet  $\Sigma$  be of constant size, and let  $G = (V, E, I)$  be an indexable Elastic Founder Graph of height  $H$ , that is, the maximum number of nodes in a block of  $G$  is  $H$ . The algorithm to find  $\kappa$ -node-MEMs spanning  $L > 3$  nodes has time complexity  $O(nH^2 + |Q| + M_\kappa)$  with  $n = \sum_{v \in V} |v|$  and  $M_\kappa$  the number of output MEMs.*

### Proof.

It derives from the complexity of the algorithm to find  $\kappa$ -node MEMs spanning exactly  $L$  nodes given before.



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## Corollary

*The results we have given for  $\kappa$ -node MEMs,  $\kappa$ -node MEMs spanning exactly  $L$  nodes and EFGs hold when  $Q[1..m]$  is replaced by a set of queries of total length  $m$ .*

## Corollary

*The algorithms we have given before (including the corollary above) can be modified to report only MEMs that occur in text  $T$  formed by concatenating the rows (ignoring gaps and adding separator symbols) of the input MSA of the indexable EFG.*

This can be done in additional  $O(|T| + r \log r)$  time and  $O(r \log n)$  bits of space, and with multiplicative factor  $O(\log \log n)$  added to the running times of the respective algorithms, where  $r$  is the number of equal-letter runs in the BWT of  $T$ .



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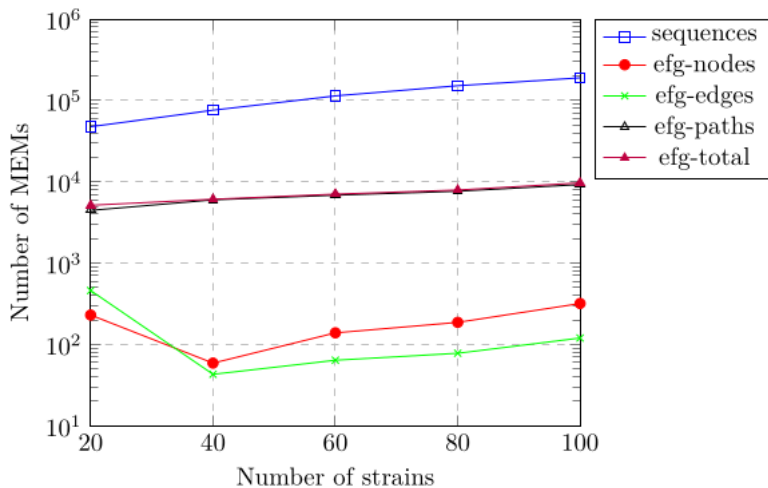
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# Experimental results

Number of MEMs with different indices and varying number of covid19 strains



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Number of BWT runs with different indexes and varying number of covid19 strains

