Edit Distances and Factorisations of Even Permutations

Anthony Labarre¹ alabarre@ulb.ac.be

Université libre de Bruxelles (U.L.B.)

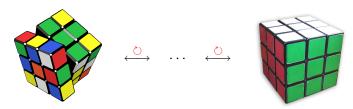
September 15th, 2008

Sixteenth European Symposium on Algorithms (ESA)

¹Funded by the "Fonds pour la Formation à la Recherche dans l'Industrie et dans l'Agriculture" (F.R.I.A.)

Edit distances

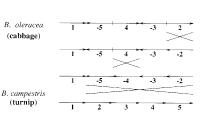
- Edit operations: given fixed set of allowed operations;
- *Edit distance*: minimum number of edit operations needed to transform *X* into *Y*:



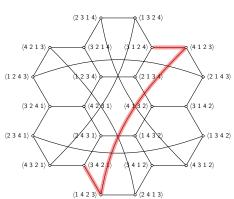
- Many applications:
 - spelling correction (example: type "dsitnace" in Google);
 - genome rearrangements;
 - interconnection networks;

Permutations can model:

genomes and mutations [Hannenhalli and Pevzner, 1999]



devices in interconnection networks

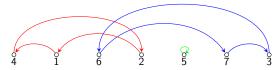


Permutations: basic definitions

- Permutation: linear ordering of $\{1, 2, ..., n\}$;
- Disjoint cycle decomposition:

$$\left(\begin{array}{ccccc} 1 & 2 & 3 & 4 & 5 & 6 & 7 \\ 4 & 1 & 6 & 2 & 5 & 7 & 3 \end{array}\right) = (1,4,2)(3,6,7)(5).$$

• The graph of permutation π , denoted by $\Gamma(\pi)$:



- π is even if $\Gamma(\pi)$ has an even number of even cycles;
- Conjugacy class: permutations with the same decomposition;
- 1-cycles (or fixed points) are often omitted;



- Let:
 - π be a permutation of $\{1, 2, \ldots, n\}$;
 - $S = \{s_1, s_2, ...\}$ be a set of permutations of $\{1, 2, ..., n\}$ (the *edit operations*);
 - ι be the *identity permutation* $\langle 1 \ 2 \ \cdots \ n \rangle$;
- We want to:
 - **①** "**sort** π **by** S": find a sequence of elements of S that sorts π and is as short as possible:

$$\pi \circ x_1 \circ x_2 \circ \cdots \circ x_t = \iota$$
 where $x_1, \ldots, x_t \in S$ and t is minimal

② "**compute the** *S***-distance** $d_S(\pi, \iota)$ ": find the length of such a sequence;

Some edit operations

- From genome rearrangements:
 - reversals:
 - transpositions:
 - block-interchanges:

- $\langle 3 \ 2 \ \underline{5} \ \underline{4} \ \underline{1} \rangle \rightarrow \langle \underline{3} \ \underline{2} \ \underline{1} \ 4 \ 5 \rangle$
- $\langle 3 \mid 25 \mid 41 \rangle \rightarrow \langle 34 \mid 12 \mid 5 \rangle$
- $\langle \boxed{5} \ 4 \ \boxed{3} \ 2 \ 1 \rangle \rightarrow \langle \boxed{3} \ 4 \ 5 \ 2 \ \boxed{1} \rangle$
- From interconnection networks:
 - prefix reversals:
 - prefix transpositions:

$$\langle \boxed{325} \boxed{41} \rangle \rightarrow \langle 41325 \rangle$$

Background

	Operation	Sorting	Distance	Diameter
classical	reversals	NP-hard	NP-hard	n-1
	signed reversals	$O(n^{3/2})$	O(n)	n+1
	block-interchanges	$O(n \log n)$	O(n)	n/2
	transpositions ²	?	?	$\frac{n}{2} \leq ? \leq \frac{2n}{3}$
prefix	reversals	?	?	$\frac{15n}{14} \le ? \le \frac{18n}{11}$
	signed reversals	?	?	$\frac{3n}{2} \leq ? \leq 2(n-1)$
	transpositions	?	?	$\frac{2n}{3} \leq ? \leq n - \log_8 n$

• All three prefix variants are 2-approximable;

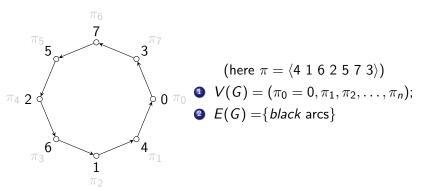
Results

- Expression of the "cycle graph" [Bafna and Pevzner, 1998] of π as an even permutation $\overline{\pi}$;
- Reformulation of **every** edit distance problem on π in terms of particular factorisations of $\overline{\pi}$;
- Simple recovery of previous results;
- New lower bound on the prefix transposition distance, which outperforms previous results;
- Improved lower bound on the maximal value of that distance $\left(\frac{2n}{3} \to \left| \frac{3n+1}{4} \right| \right)$;

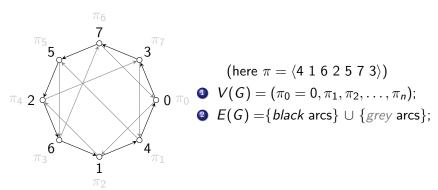
• The "cycle graph" of π , denoted by $G(\pi)$:

$$\pi_{5} = \begin{pmatrix} \pi_{6} & \pi_{7} & \pi_$$

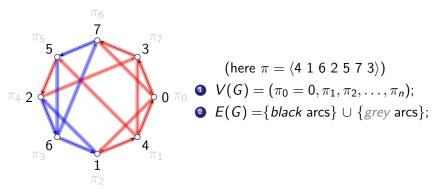
• The "cycle graph" of π , denoted by $G(\pi)$:



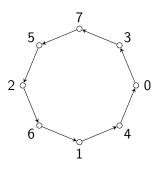
• The "cycle graph" of π , denoted by $G(\pi)$:

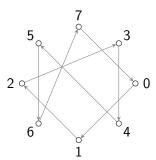


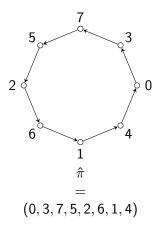
• The "cycle graph" of π , denoted by $G(\pi)$:

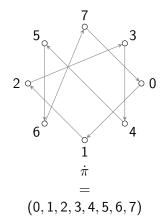


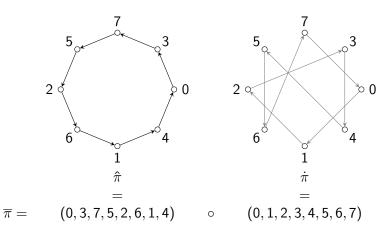
Unique decomposition into "alternating cycles";

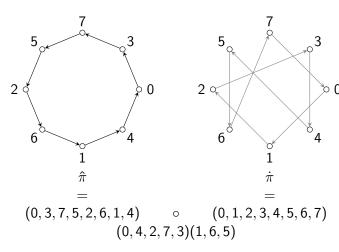




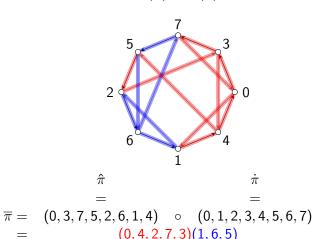








• We have $\overline{\pi} = \hat{\pi} \circ \dot{\pi}$, with $\Gamma(\overline{\pi}) \simeq G(\pi)$; indeed:



A general lower bounding technique

• Note that "sorting by S" is equivalent to "factorising by S":

$$\pi \circ \underbrace{x_1 \circ x_2 \circ \cdots \circ x_t}_{x_1, x_2, \dots, x_t \in S} = \iota \Leftrightarrow \pi = \underbrace{x_t^{-1} \circ x_{t-1}^{-1} \circ \cdots \circ x_1^{-1}}_{x_1^{-1}, x_2^{-1}, \dots, x_t^{-1} \in S}$$

Theorem 1

Let:

1
$$S \subset S_n$$
, with $S = \{s_1, s_2, \ldots\}$,

3 C the set of conjugacy classes that intersect S'.

Then for all π in S_n , every factorisation of π into t elements of Syields a factorisation of $\overline{\pi}$ into t elements of \mathcal{C} .

A general lower bounding technique

Theorem 1 in action

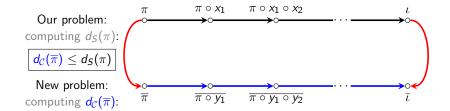
Our problem: $\xrightarrow{\pi} \xrightarrow{\pi \circ x_1} \xrightarrow{\pi \circ x_1 \circ x_2} \cdots \xrightarrow{\iota}$ computing $d_S(\pi)$:

A general lower bounding technique

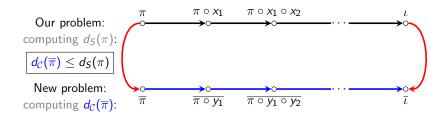
Theorem 1 in action



Theorem 1 in action



Theorem 1 in action



Lemma 2

For all π , σ in S_n :

$$\overline{\pi \circ \sigma} = \pi \circ \overline{\sigma} \circ \pi^{-1} \circ \overline{\pi}$$
$$= \overline{\sigma}^{\pi} \circ \overline{\pi}.$$

A lower bound on the block-interchange distance

Example 3 (lower bound on $bid(\pi)$)

• $S = \{ block-interchanges \}, denoted by <math>\beta(i, j, k, l);$

A lower bound on the block-interchange distance

Example 3 (lower bound on $bid(\pi)$)

• $S = \{ block-interchanges \}, denoted by <math>\beta(i, j, k, l);$ we have:

$$\overline{\beta(i,j,k,l)} = (i-1,k-1)(j-1,l-1) \quad (1 \le i < j \le k < l \le n+1).$$

- We have $S' \subseteq \mathcal{C}$, where \mathcal{C} contains all pairs of 2-cycles;
- We have $d_{\mathcal{C}}(\overline{\pi}) = \frac{|\overline{\pi}| c(\Gamma(\overline{\pi}))}{2}$;
- Therefore, we recover the result of [Christie, 1996]:

$$\forall \ \pi \in S_n : bid(\pi) \geq \frac{n+1-c(\Gamma(\overline{\pi}))}{2}.$$

A lower bound on the transposition distance

Example 3 (lower bound on $td(\pi)$)

• $S = \{\text{transpositions}\}\$, denoted by $\tau(i, j, k)$;

$$\left(\begin{array}{ccccc} 1 & \cdots & i-1 & i & i+1 & \cdots & j-2 & j-1 \\ 1 & \cdots & i-1 & j & j+1 & \cdots & k-1 \end{array}\right. \begin{vmatrix} j & j+1 & \cdots & k-1 \\ i & i+1 & \cdots & j-2 & j-1 \\ k & \cdots & n \\ \end{array}\right).$$

A lower bound on the transposition distance

Example 3 (lower bound on $td(\pi)$)

• $S = \{\text{transpositions}\}\$, denoted by $\tau(i, j, k)$; we have:

$$\overline{\tau(i,j,k)} = (i-1,k-1,j-1) \quad (1 \le i < j < k \le n+1).$$

- We have $S' \subseteq C$, the set of all 3-cycles;
- We have $d_{\mathcal{C}}(\overline{\pi}) = \frac{|\overline{\pi}| c_{odd}(\Gamma(\overline{\pi}))}{2}$;
- Therefore, we recover the result of [Bafna and Pevzner, 1998]:

$$\forall \ \pi \in S_n : td(\pi) \geq \frac{n+1-c_{odd}(\Gamma(\overline{\pi}))}{2}.$$

A new lower bound on the prefix transposition distance

Example 3 (lower bound on $ptd(\pi)$)

• $S = \{ prefix transpositions \}; we get:$

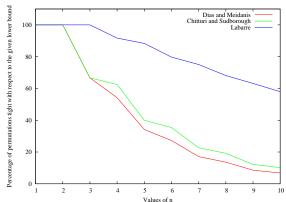
$$\overline{\tau(1,j,k)} = (0, k-1, j-1) \quad (1 < j < k \le n+1).$$

- We have $S' \subseteq \mathcal{C}$, the set of all 3-cycles that contain 0;
- We can compute $d_{\mathcal{C}}(\overline{\pi})$, and this yields the following *new* lower bound :

$$\forall \ \pi \in S_n : ptd(\pi) \geq \frac{n+1+c(\Gamma(\overline{\pi}))}{2} - c_1(\Gamma(\overline{\pi})) - \begin{cases} 0 & \text{if } \pi_1 = 1, \\ 1 & \text{otherwise.} \end{cases}$$

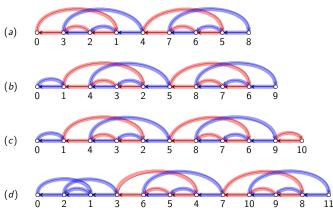
Quality of the results

- Block-interchanges: the lower bound is the exact distance;
- Prefix transpositions: the new result:
 - always outperforms [Dias and Meidanis, 2002];
 - "often" outperforms [Chitturi and Sudborough, 2008]:



The prefix transposition diameter of S_n

• These permutations satisfy $ptd(\pi) \ge \lfloor \frac{3n+1}{4} \rfloor$, thereby improving on the lower bound of 2n/3 by [Chitturi and Sudborough, 2008]:



Future work

- Complexity/approximation issues (transpositions, prefix operations);
- Can the $\overline{\pi}$ model provide *upper* bounds?
- Extending the $\overline{\pi}$ model to \emph{signed} permutations and/or other structures;

Thank you!



Bafna, V. and Pevzner, P. A. (1998). Sorting by transpositions.

SIAM Journal on Discrete Mathematics, 11(2):224-240 (electronic).



Chitturi, B. and Sudborough, I. H. (2008). Bounding prefix transposition distance for strings and permutations.

In HICSS, page 468, Los Alamitos, CA, USA.



Christie, D. A. (1996). Sorting permutations by block-interchanges.

Information Processing Letters, 60(4):165–169.



Dias, Z. and Meidanis, J. (2002). Sorting by prefix transpositions.

In SPIRE, volume 2476 of LNCS, pages 65-76. Springer-Verlag.



Elias, I. and Hartman, T. (2006).

A 1.375-approximation algorithm for sorting by transpositions. *IEEE/ACM Trans. Comput. Biol. Bioinform.*, 3(4):369–379.



Hannenhalli, S. and Pevzner, P. A. (1999).

Transforming cabbage into turnip: Polynomial algorithm for sorting signed permutations by reversals.

Journal of the ACM, 46(1):1-27.