

A 1.375-Approximation Algorithm for Sorting By Transpositions

Isaac Elias

Royal Institute of
Technology

Tzvika Hartman

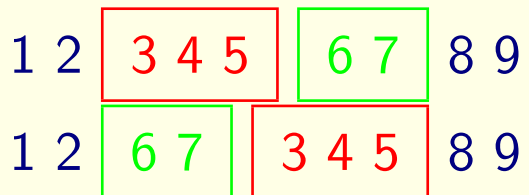
Weizmann Institute of
Science

Sorting By Transpositions

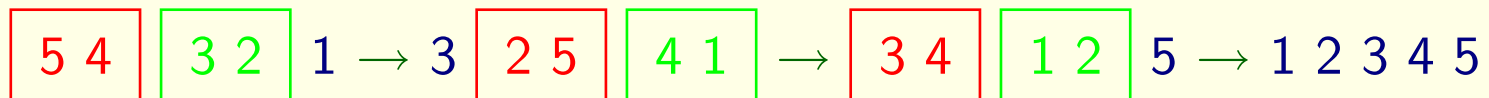
Input: A permutation π .

Output: The least number of transpositions for sorting π .

A transposition is an operation which switches two adjacent blocks in a permutation.



Example of Sorting

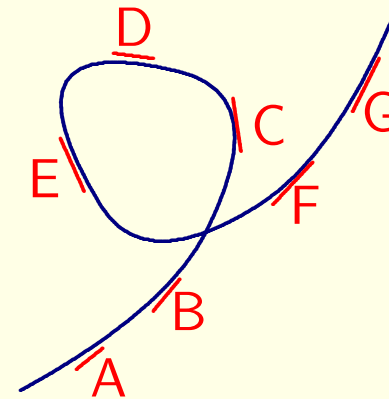


Genome Rearrangements

Elements represent genes on a chromosome.

Segments on the chromosome can be:

- Transposed - caused by transposomes
- Reversed
- Transreversed
- Deleted
- Duplicated

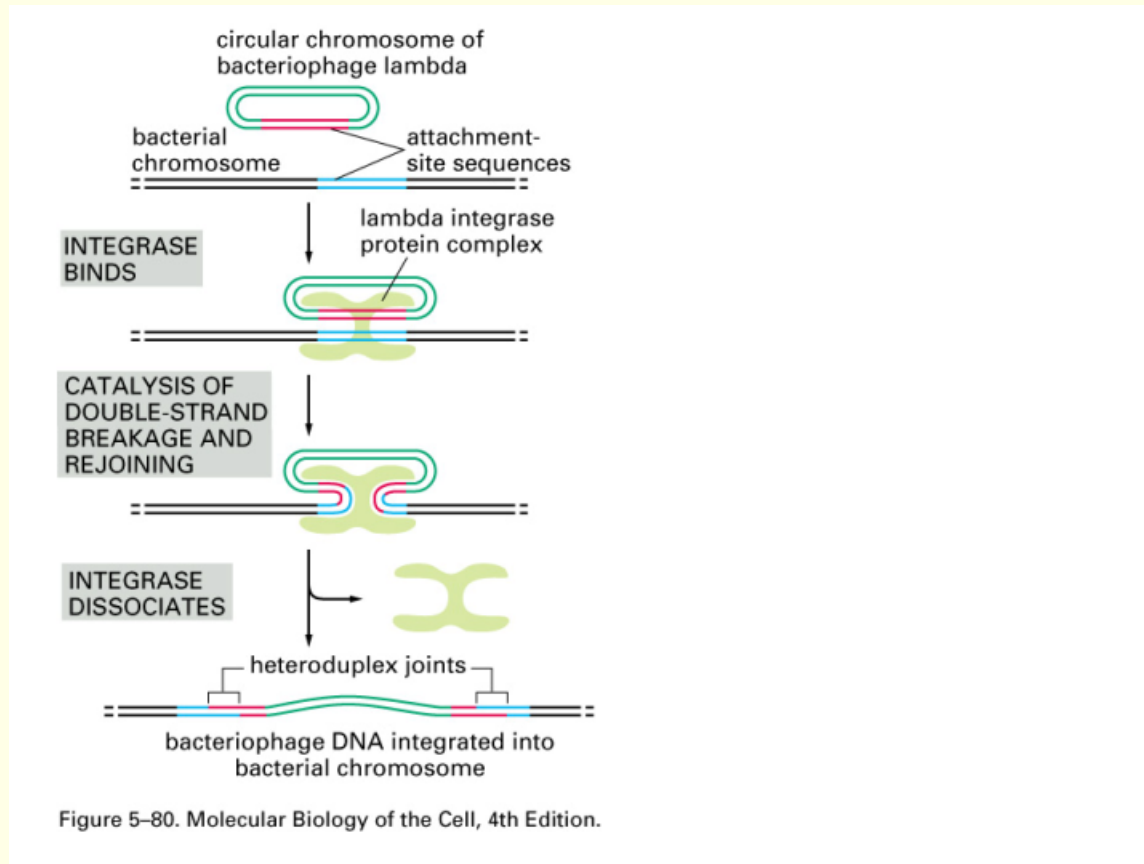


$$\pi = A B C D E F G$$

The Genome Rearrangement Problem

Given two genomes find out what rearrangement events have occurred.

Bacteriophage - Transposome



Bacteria inserts a movable segment - a transposome.

Why study GR?

Evolution

- Rare events - allow for phylogenetic inference further back
- Large scale data: takes the whole genome into consideration
- Better multi-species analysis

Cancer

- Cancer cells undergo many genome rearrangements.
- Used for cancer research, distinguish between benign and malignant tumors, diagnostics, etc.

Previous Results

SBT

1.5-approx

NP/P ?

[Bafna Pevzner, Christie, Hartman]

Transposition Diameter
(longest distance)

$$\leq \frac{2n}{3}$$

[Erikson et.al.]

$$\geq \lfloor \frac{n+1}{2} \rfloor + 1$$

[Christie, Meidanis et.al.]

Our Results

SBT

1.375-approx

Transposition Diameter

$$\geq \lfloor \frac{n+2}{2} \rfloor + 1$$

Diameter for:

Simple permutations

$$\lfloor n/2 \rfloor$$

2-permutations

$$n/2$$

3-permutations

$$\lesssim \frac{11n}{24}$$

The Breakpoint Graph [Bafna Pevzner]

1 2 3 4 5

The Breakpoint Graph [Bafna Pevzner]

1. Add 0 and $n + 1$ to the beginning and end and give each element a left and a right vertex.

0 ▪ ▪1 ▪ ▪2 ▪ ▪3 ▪ ▪4 ▪ ▪5 ▪ ▪6

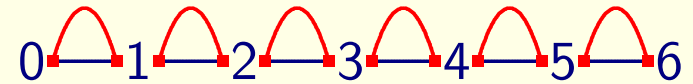
The Breakpoint Graph [Bafna Pevzner]

1. Add 0 and $n + 1$ to the beginning and end and give each element a left and a right vertex.
2. Connect adjacent elements with an edge.



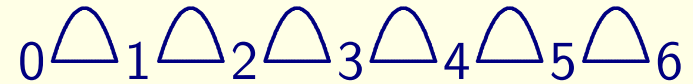
The Breakpoint Graph [Bafna Pevzner]

1. Add 0 and $n + 1$ to the beginning and end and give each element a left and a right vertex.
2. Connect adjacent elements with an edge.
3. Connect successive elements by an arc.



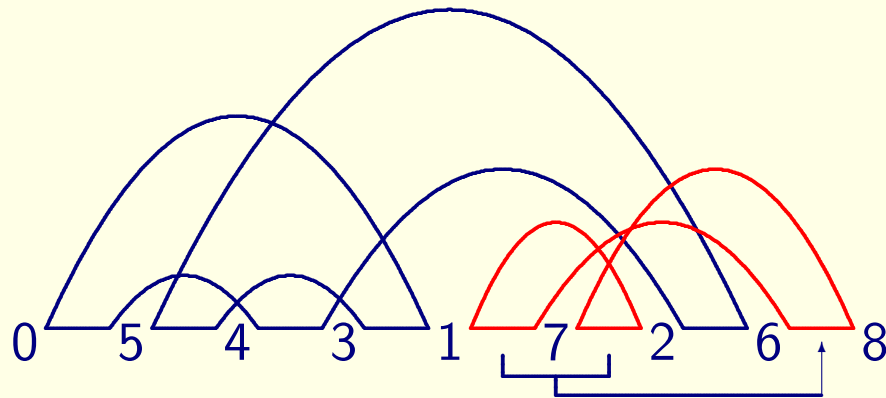
The Breakpoint Graph [Bafna Pevzner]

1. Add 0 and $n + 1$ to the beginning and end and give each element a left and a right vertex.
2. Connect adjacent elements with an edge.
3. Connect successive elements by an arc.

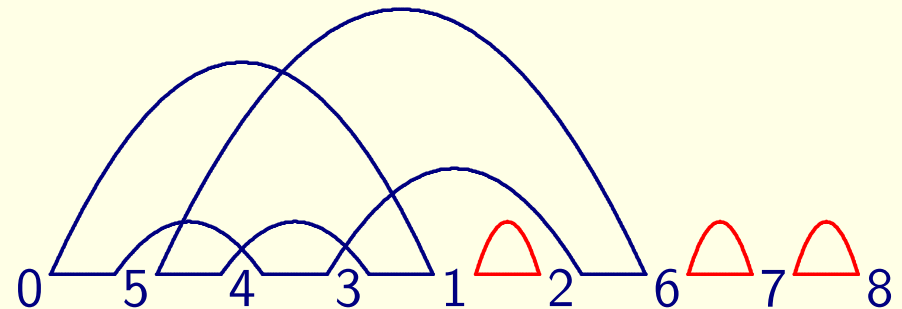


Decomposes into cycles.

Length of cycle = Number of arcs.



One 5-cycle and one 3-cycle.



One 5-cycle and three 1-cycles.

A transposition cuts 3 edges.

A Lower Bound [Bafna Pevzner]

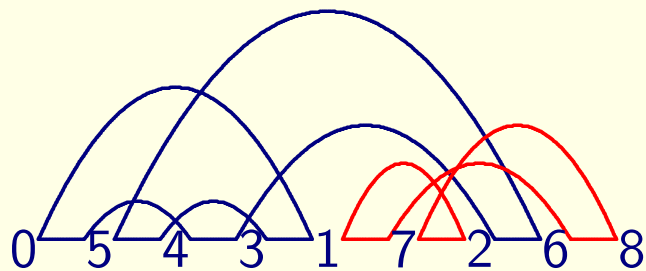
Game Create $n + 1$ odd cycles in as few moves as possible.

k-move the number of odd cycles is increased by k cycles.

Lemma There are only 2 , 0 , and -2 moves.

Lower bound

$$d(\pi) \geq \frac{n + 1 - c_{\text{odd}}(\pi)}{2}$$



$$d(\pi) \geq \frac{8 - 2}{2} = 3$$

Making Approximation Algorithms

Do not use -2 -moves!

Notation (x, y) : Sequence of x moves with y 2 -moves.

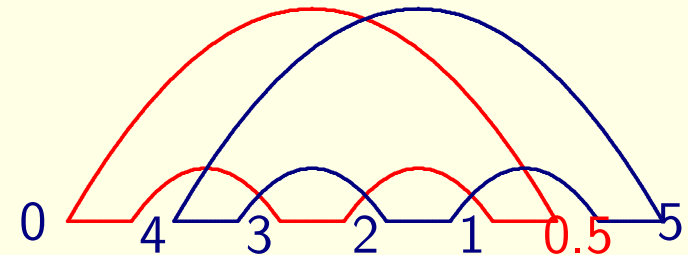
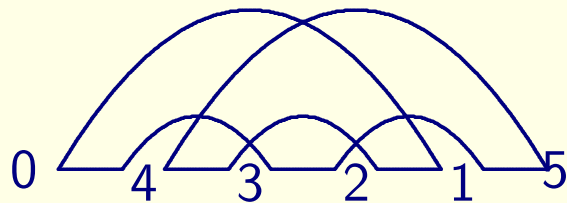
Lemma [BP] There is always $(3, 2)$ -sequence; for every three moves at least two 2 -moves can be performed.

$$\Rightarrow \frac{3}{2} = 1.5\text{-approximation}$$

Our algorithm uses $(11, 8)$ -sequences; for each 11 moves it uses at least 8 2 -moves.

$$\Rightarrow \frac{11}{8} = 1.375\text{-approximation}$$

Adding Elements - Simple Permutations [Lin Xue]



The 5-cycle is broken into two 3-cycles.

All cycles can be broken down into cycles of length ≤ 3 , called simple permutations.

Elements can be added without changing the lower bound.

$$\frac{n + 1 - c_{\text{odd}}(\pi)}{2} = \frac{n' + 1 - c_{\text{odd}}(\pi')}{2} \quad \Rightarrow \quad \frac{4 + 1 - 1}{2} = \frac{5 + 1 - 2}{2}$$

A 1.375-Approximation

Step 1 Simplify π

$$\begin{array}{ccccccc} \pi & \rightarrow & \pi' \dots & \rightarrow & \pi^{(k)} \\ d(\pi) & \leq & \dots & \leq & d(\pi^{(k)}) \\ \frac{n+1-c_{\text{odd}}(\pi)}{2} & = & \dots & = & \frac{n+1+k-c_{\text{odd}}(\pi^{(k)})}{2} \end{array}$$

Step 2 Find sorting using only $(11, 8)$ -sequences for $\pi^{(k)}$

Step 3 Use sorting of $\pi^{(k)}$ to sort π .

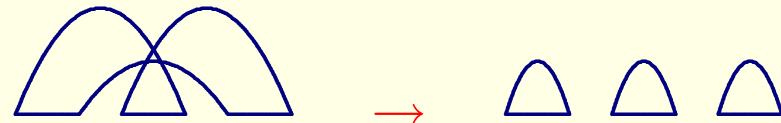
How to do Step 2?

Observations

1. If find a sequence (x, y) s.t. $\frac{x}{y} \leq \frac{11}{8}$ then ok, e.g. $(1, 1)$ and $(4, 3)$.
2. If there is a **2-cycle** then there is a 2-move. **Ok!**
3. There are two types of 3-cycles.

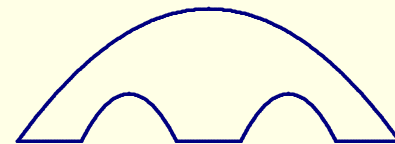
Oriented 3-Cycle

Has a 2-move. **Ok!**



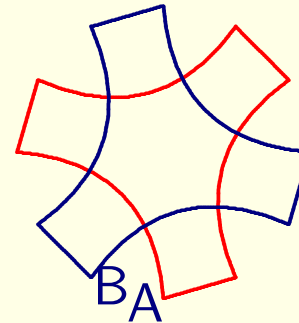
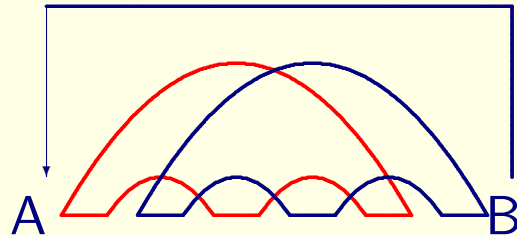
Unoriented 3-Cycle

Does not have 2-move. **Not ok!**



⇒ Only unoriented 3-cycles!

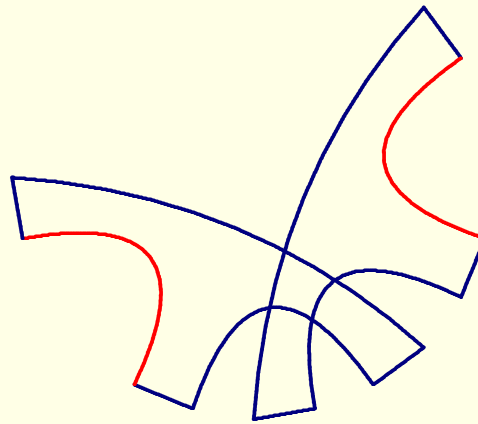
4. [Hart] Sorting linear permutations \Leftrightarrow Sorting circular permutations
Relative structure of the cycles matters (cyclical shift, mirroring).



\Rightarrow **Analyse structures of unoriented 3-cycles.**

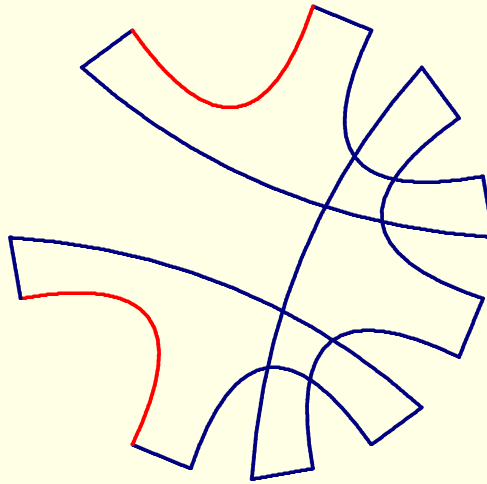
Configurations of Unoriented 3-cycles

Lemma [BP] In a breakpoint graph every arc has to cross another arc.



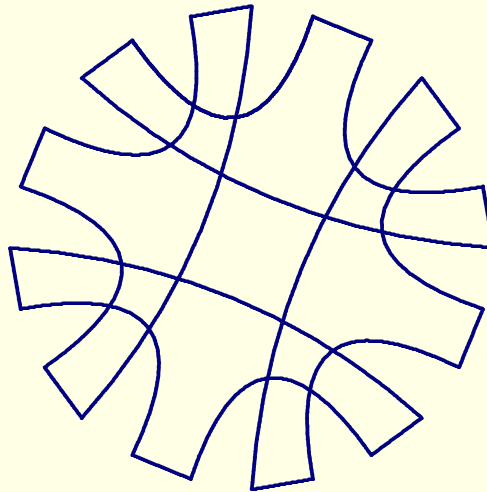
Configurations of Unoriented 3-cycles

Lemma [BP] In a breakpoint graph every arc has to cross another arc.



Configurations of Unoriented 3-cycles

Lemma [BP] In a breakpoint graph every arc has to cross another arc.

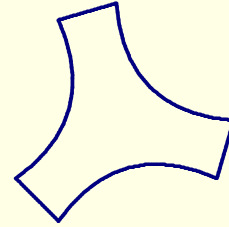


Configurations can be built by adding cycles intersecting with another cycle.

Idea A program that analyses configurations to see if an $\frac{11}{8}$ -seq always exists.

Breadth First Search to Prove Existence of $\frac{11}{8}$ -seq

1. Initiate a queue to contain the configuration with one cycle.



2. While the queue is non-empty do:

(a) Remove the first configuration, A , from the queue.

(b) For each way of adding a cycle; B extension A :

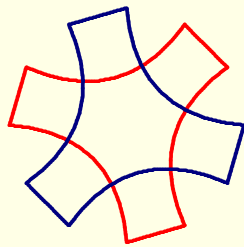
i. If B has $\frac{11}{8}$ -seq then all permutations containing B has one too.

ii. Otherwise add B to the queue and continue analyzing it in next iteration.

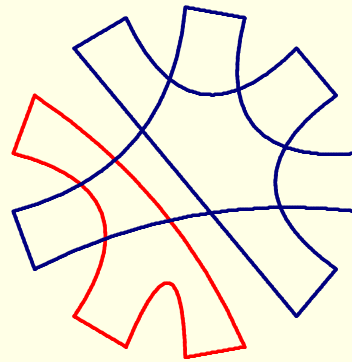
Has to stop otherwise $\frac{11}{8}$ doesn't exist!

Adding Cycles to Configurations without $\frac{11}{8}$ -seq

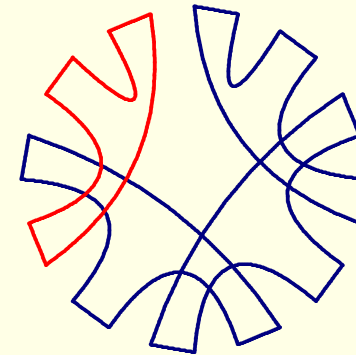
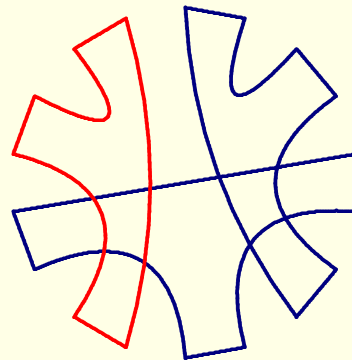
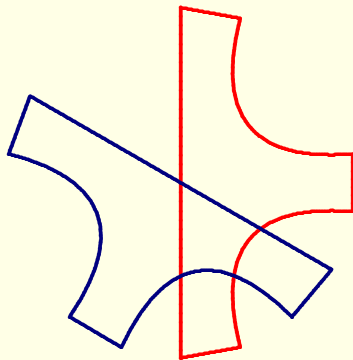
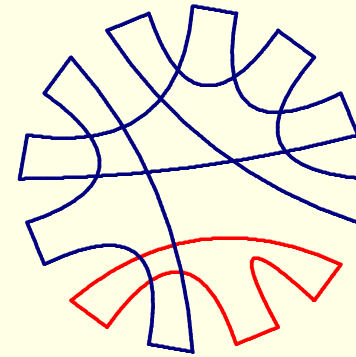
ITER 1



ITER 2

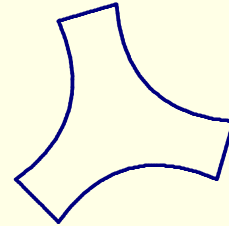


ITER 3



What happens in the BFS?

INIT The queue contains one configuration.



ITER 1 For each configuration in the queue add cycles in all possible ways.

- If extension has $\frac{11}{8}$ -seq then ok.
- Otherwise add to queue and continue adding cycles in ITER 2.

...

ITER 9 The queue is empty (all configurations of 9-cycles have $\frac{11}{8}$ -seq).

Computer aided proof with 80,000 cases.

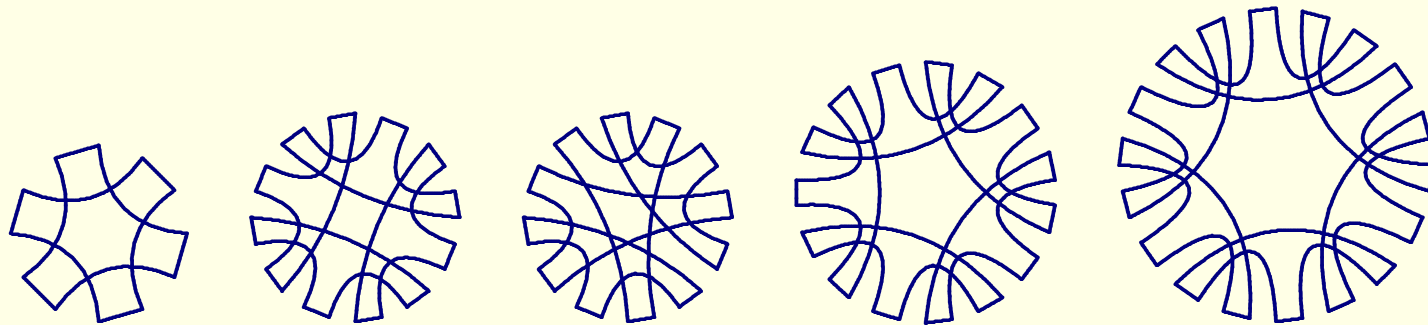
Bad Small Components

We want to show:

Lemma Every 3-permutation with ≥ 8 cycles has an $\frac{11}{8}$ sequence.

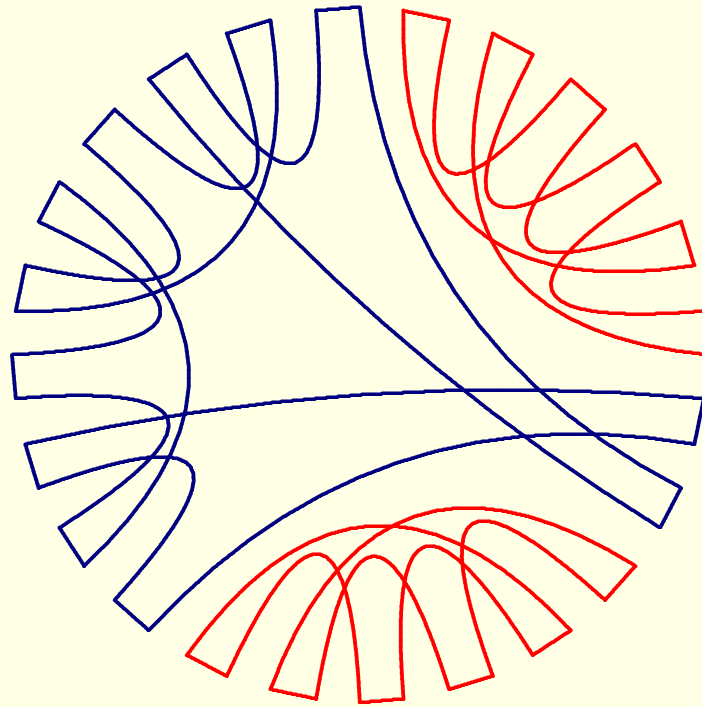
We have shown that components ≥ 9 cycles have an $\frac{11}{8}$ -sequence.

Components < 9 cycles that do not have (x, y) .



New case analysis showing that combinations with ≥ 8 cycles have $(11, 8)$ -seq.

Example of Combination



The Approximation Algorithm

Algorithm *Sort* (π)

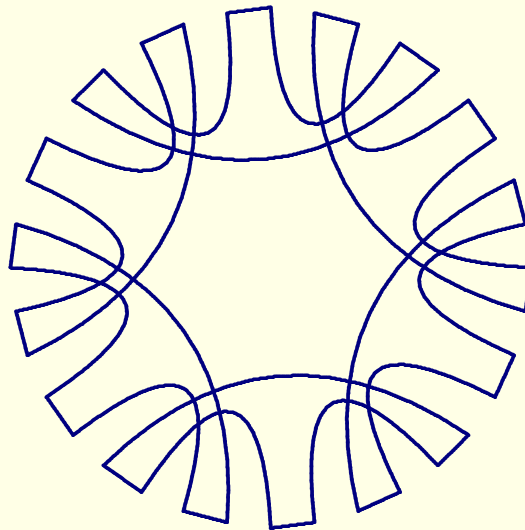
1. Transform permutation π into a simple permutation $\hat{\pi}$.
2. Check if there is a $(2, 2)$ -sequence. If so, apply it.
3. While $G(\hat{\pi})$ contains a 2-cycle, apply a 2-move.
4. While $G(\hat{\pi})$ contains at least 8 cycles apply a $(4, 3)$ or an $(11, 8)$ sequence.
5. While $G(\hat{\pi})$ contains a 3-cycle, apply a $(3, 2)$ sequence.
6. Mimic the sorting of π using the sorting of $\hat{\pi}$.

Can we do better?

If we analyse even bigger components can we do better?

Probably not! It seems as if the unoriented necklace can not be sorted better than with $(11, 8)$ -sequences.

A new lower bound is probably needed!



Diameter for 3-permutations

Definition The longest sorting distance for any permutation made up only of 3-cycles.

Today: Every 3-permutation can be sorted with $(11, 8)$ sequences.

If a permutation has k 3-cycles it can be sorted using

$$\sim \frac{k}{8} \cdot 11 \text{ moves.}$$

All cycles length 3 so $n = 3 \cdot k$.

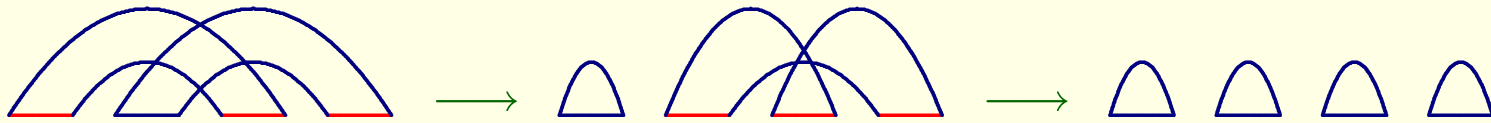
Hence upper bound

$$\lesssim \frac{11k}{8} = \frac{11n}{24}$$

Diameter for 2-permutations

Definition The longest sorting distance for any permutation made up only of 2-cycles.

All arcs must cross other arc.



Creates 4 1-cycles in 2 moves.

The identity has $n + 1$ cycles.

Every 2-permutation is sorted using

$$= \frac{n + 1}{4} \cdot 2 = \frac{n + 1}{2} \text{ moves.}$$

Diameter for Simple Permutations

Definition The longest sorting distance for any permutation made up only of 2-cycles and 3-cycles.

Same kind of proof as for 2-permutations.

About 10 cases to analyse.

Same diameter as for 2-permutation:

$$\sim \frac{n+1}{2} \text{ moves.}$$

General Diameter

Definition The longest sorting distance for any permutation.

Earlier conjecture the reversed permutation hardest to sorted.

We show that:

$$\pi = 0 \ 4 \ 3 \ 2 \ 1 \ 5 \ 13 \ 12 \ 11 \ 10 \ 9 \ 8 \ 7 \ 6 \ 14 \quad \text{2-permutation} \quad n+1$$

requires one more move than the reversed to be sorted.

Reversed require $\sim \frac{n}{2} + 1$ moves and π requires $\frac{n}{2} + 2$ moves.

Many interesting open questions.

Cycles seem to partion into groups and combinations of these groups are hard to sort.

Results

SBT 1.375-approx $\geq \lfloor \frac{n+2}{2} \rfloor + 1$
Transposition Diameter

Diameter for:
Simple permutations $\lfloor n/2 \rfloor$
2-permutations $n/2$ only 2-cycles
3-permutations $\lesssim \frac{11n}{24}$ only 3-cycles

Diameters for circular permutations.

Acknowledgments

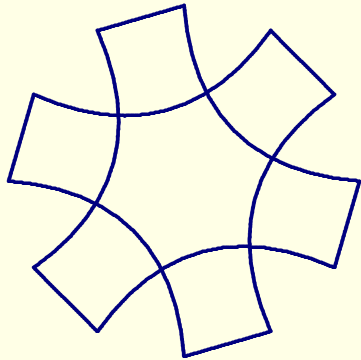
Our advisors
Prof. Jens Lagergren
and
Prof. Ron Shamir

Elad Verbin

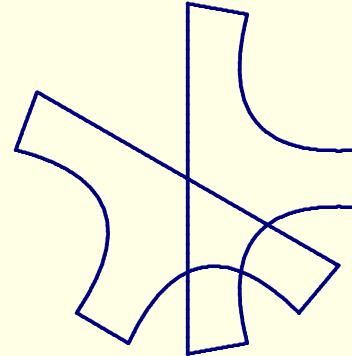
Thanks!

A $(3, 2)$ -sequence = 1.5 Approximation

There are two configurations with two cycles:

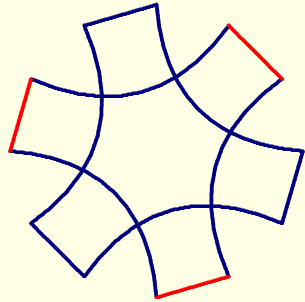


Interleaving cycles
 $(3, 2)$ -sequence exist!

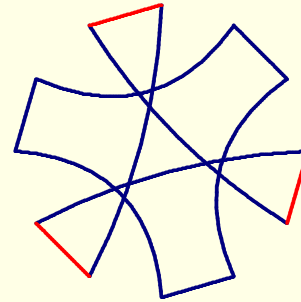


Two intersecting cycles
 $(3, 2)$ -sequence **does not** exist.

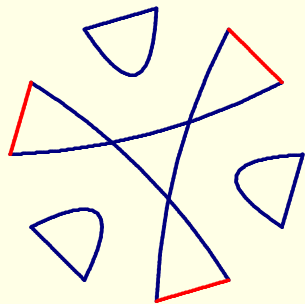
Sorting Two Interleaving



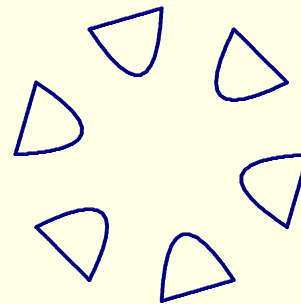
→
0-move



→
2-move



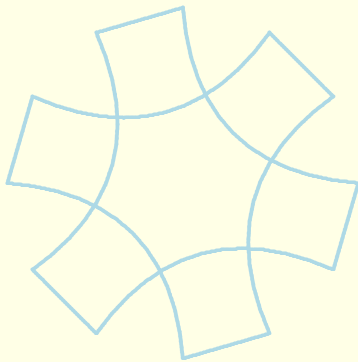
→
2-move



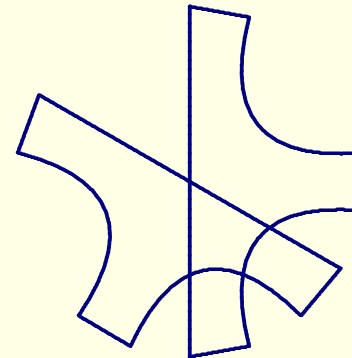
(3,2)-seq.

A $(3, 2)$ -sequence = 1.5 Approximation (cont.)

There are two configurations with two cycles:



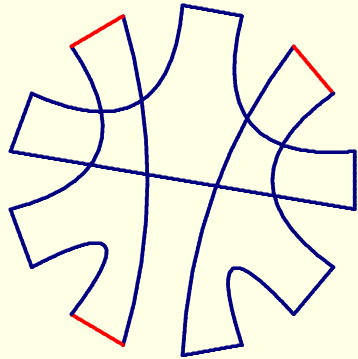
Interleaving cycles
 $(3, 2)$ -sequence exist!



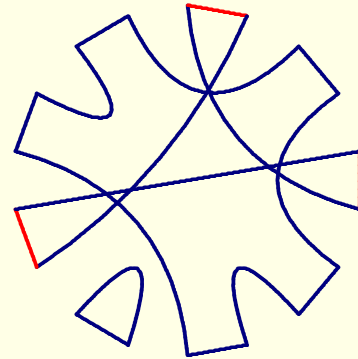
Two intersecting cycles
 $(3, 2)$ -sequence does not exist.
But every extension has a
 $(3, 2)$ -sequence.

⇒ There is always a $(3, 2)$ -sequence.

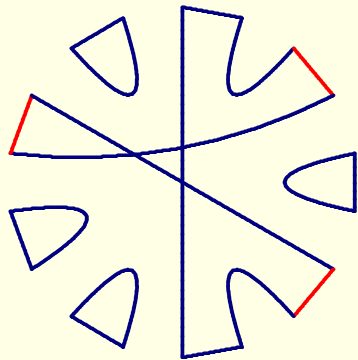
Sorting Extension of Two Intersecting



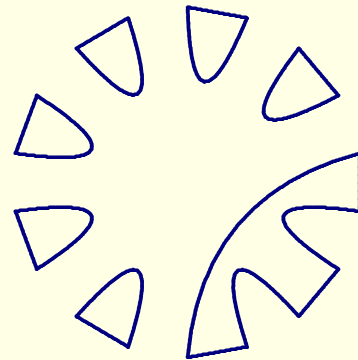
→
0-move



→
2-move



→
2-move



(3,2)-seq.