

Exercises – Algorithms for Genome Rearrangement

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<http://wiki.techfak.uni-bielefeld.de/gi/Teaching/2015summer/gr>

Exercise List 4 — 04.05.2014

Discussion of exercises on: 11.05.2014

Exercise 1

(2 Points)

Given the signed permutation

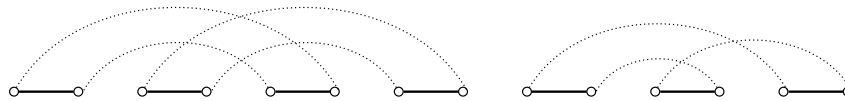
$$\pi = (+3 +5 +4 +6 +2 +1 +7 -8 +9)$$

- How many components does the graph $BP(\pi)$ has, and of which type?
- What is the reversal distance?
- Find a sequence of reversals that transform all unoriented components into oriented components.

Exercise 2

(2 Points)

Consider the following breakpoint graph.

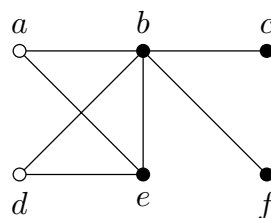


- What is the reversal distance for this breakpoint graph?
- Find a permutation π that has this breakpoint graph.

Exercise 3

(2 Points)

Suppose that a given permutation has the following overlap graph:



- What is the vertex with maximum score? Apply the reversal defined by this vertex, update the overlap graph, and repeat the process until the permutation is sorted.
- Can you find a breakpoint graph that corresponds to the overlap graph in the figure?

Exercise 4

(2 Points)

The number of possible (unsigned) permutations over $\{1, 2, \dots, n\}$ is $n!$. Obviously, there exist bijective mappings between the numbers $1, 2, \dots, n!$ and permutations over $\{1, 2, \dots, n\}$. Find such a mapping that is computable in both directions in polynomial time. *Tip: Google is your friend.*

Factoradic numbers:

$(i_k i_{k-1} \dots i_2 i_1 i_0)$ represents the integer $i_k k! + i_{k-1} (k-1)! + \dots + i_2 2! + i_1 1! + i_0 0!$.
Now, a permutation π is represented by an array A of size n where position $A[i]$ the number of indices $i' > i$ with values $\pi[i']$ smaller than $\pi[i]$:

```
1 2 3 4 - 0 0 0 0
1 2 4 3 - 0 0 1 0
1 3 2 4 - 0 1 0 0
1 3 4 2 - 0 1 1 0
1 4 2 3 - 0 2 0 0
1 4 3 2 - 0 2 1 0
2 1 3 4 - 1 0 0 0
2 1 4 3 - 1 0 1 0
2 3 1 4 - 1 1 0 0
2 3 4 1 - 1 1 1 0
2 4 1 3 - 1 2 0 0
2 4 3 1 - 1 2 1 0
3 1 2 4 - 2 0 0 0
```

...

The entries in the i th last column range from 0 to $i - 1$. All rows are different.

Interpreted as factoradic numbers, we get a unique mapping.

Quadratic-time calculations are simple. Linear-time calculations are more difficult, one direction is shown in the following, the other direction is more complicated.

See <http://en.wikipedia.org/wiki/Permutation> (Numbering permutations):

```
function permutation(k, s) {
    var int factorial:= 1;
    for j = 2 to length(s) {
        factorial := factorial* (j-1);
        swap( s[j - ((k / factorial) mod j)], s[j]);
    }
    return s;
}
```

Exercise 5

(3 Points)

After a *cycle merge* reversal, that is, a reversal defined by black edges in two different cycles, the two cycles are merged into one. Prove that this new cycle is always oriented.