

# Algorithms for Genome Research

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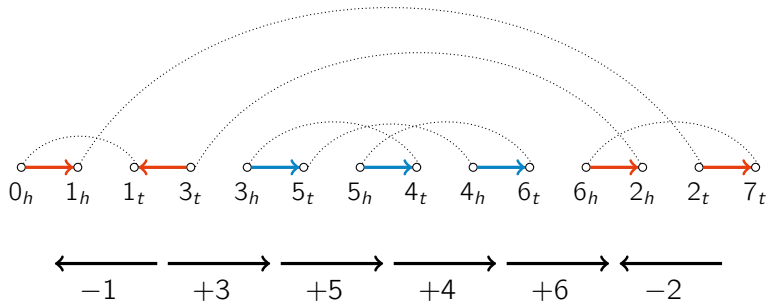
Lecture 2 – Sorting by Signed Reversals II

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# Quick Recap

BP Graph, **oriented** and **unoriented** components:



# Quick Recap

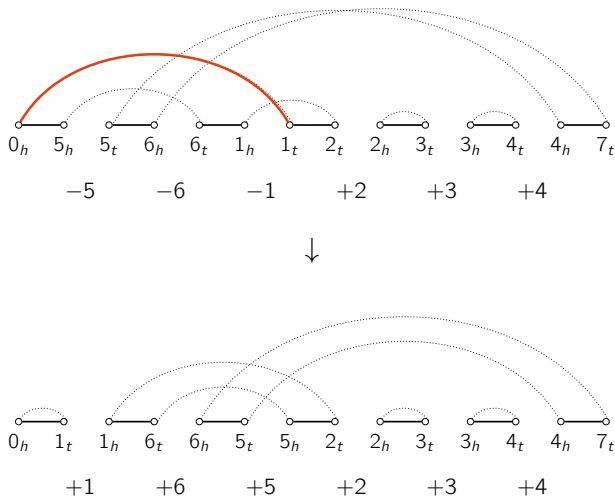
- Sorting is equivalent of increasing # of cycles in BP graph
  - In *oriented (good) components* – at least 1 oriented edge – this is always possible.
  - In *unoriented (bad) components*, not, so we need extra operations.

So we have the following **lower bound**:

$$d_R(\pi) \geq N - C$$

- There are also reversals that increase the number of cycles, but create unoriented components.

# Bad reversal - Example



- Increased number of cycles but created a bad component!

# Finding “good” reversals

- Is it possible to find a reversal that increases the number of cycles **AND** also does not create an unoriented component? **YES!**

# Sorting oriented components

Theorem (Hannenhalli-Pevzer, 95)

If the graph  $BP(\pi)$  has only **oriented components**, then

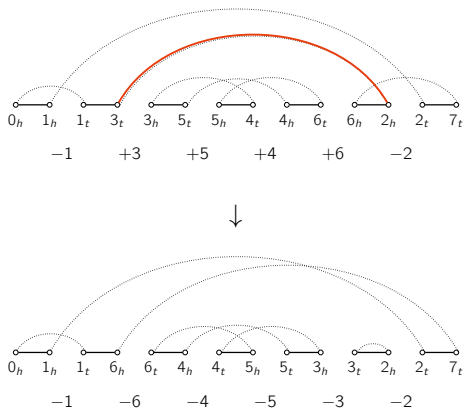
$$d_R(\pi) = N - C$$

where  $N$  is the number of elements of  $\pi$  and  $C$  is the number of cycles of  $BP(\pi)$ .

- This means that there is always at least one “good” reversal, that increases the number of cycles of  $BP(\pi)$  and *does not create any unoriented component*.
- These are called **safe reversals**. How can we find them?

# Safe reversals - Definitions

- The **score** of a reversal is the number of *oriented edges* in the BP graph, *after* the application of the reversal.



The score of this reversal is **two**.

# Safe reversals

- **Safe reversals** are reversals that increase the number of cycles of the BP graph by one and do not create new unoriented components.
- Can we always find safe reversals? Yes:

## Theorem (Bergeron, 2001)

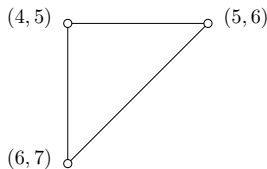
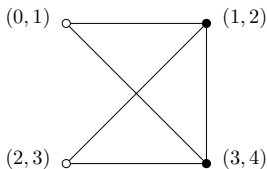
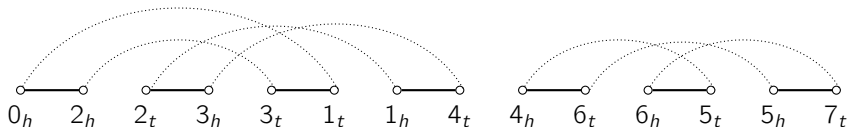
*Among all possible oriented reversals, a reversal of maximal score is always safe.*

- **Algorithm:** Apply maximal score reversals until all components are sorted.



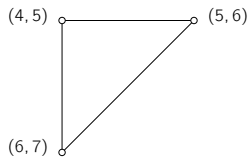
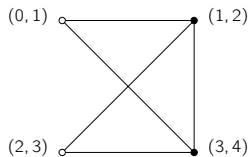
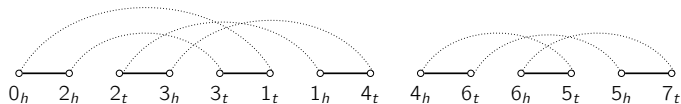
# Finding safe reversals with the Overlap Graph

- The **overlap graph**  $O(\pi)$  is a graph where:
  - Vertices are the grey edges of  $BP(\pi)$ . If the edge is oriented, the vertex is black, otherwise is white.
  - When two grey edges overlap, there is an edge between the corresponding vertices.



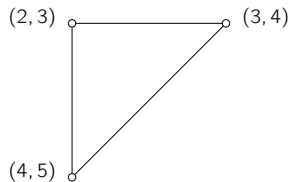
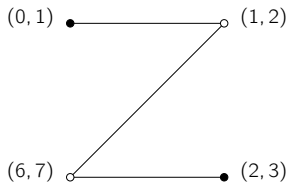
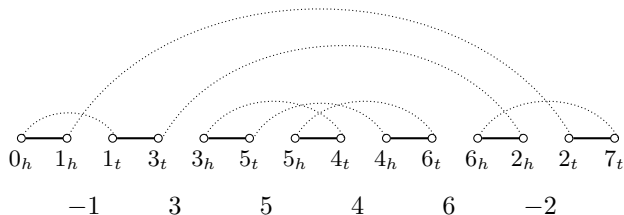
# BP Graph vs Overlap Graph

BP Graph	Overlap Graph
Component	Connected component
Oriented edge	Black vertex, <i>odd degree</i>
Unoriented edge	White vertex, <i>even degree</i>
Oriented component	Component with at least 1 black vertex
Unoriented component	Component with only white vertices



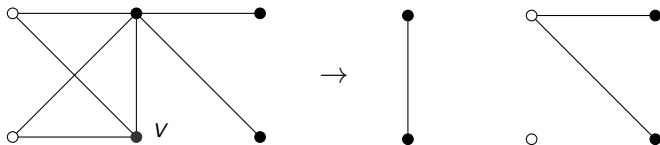
# Another Example

$$\pi = [-1 \ 3 \ 5 \ 4 \ 6 \ -2]$$

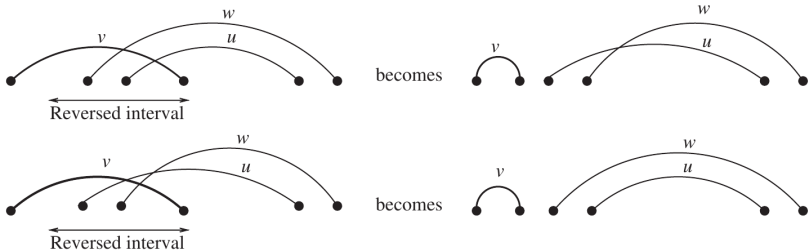


# Effect of Reversal in the Overlap Graph

- A reversal *induced by a vertex  $v$*  is the reversal that is induced by the corresponding grey edge in the breakpoint graph.
  - What happens in  $O(\pi)$  after applying an oriented reversal in a vertex  $v$ ?
- 1 The subgraph induced by  $v$  and its neighbours is **complemented**.



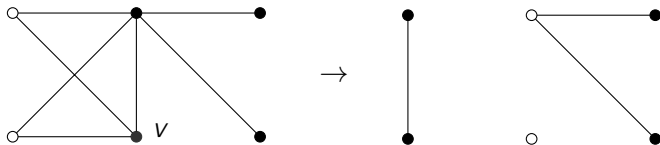
**Why?**



*A. Bergeron/Discrete Applied Mathematics 146 (2005) 134–145*

# Effect of Reversal in the Overlap Graph

- 2 All neighbours of  $v$  have their orientation inverted.



**Why?**

## Reversal Score with $O(\pi)$

We know how the overlap graph changes with a reversal, then it is possible to find an equation for the reversal score of any vertex  $v$ :

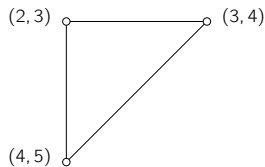
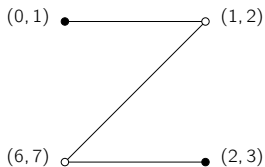
### Definition (Reversal score)

The score of a reversal induced by a vertex  $v$  in the overlap graph is given by

$$s(v) = T + U - O - 1$$

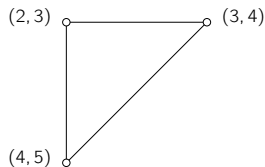
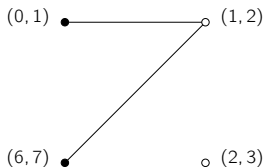
where  $T$  is the number of oriented vertices in the graph,  $U$  and  $O$  are the number of unoriented and oriented vertices adjacent to  $v$ , respectively.

## Reversal Score - example



For  $v = (2,3)$ , we have  $T = 2$ ,  $U = 1$ ,  $O = 0$ . Therefore  $s(v) = T + U - O - 1 = 2$ .

After applying the reversal, we have the following graph:



and we see that the score (number of oriented vertices) is indeed 2.

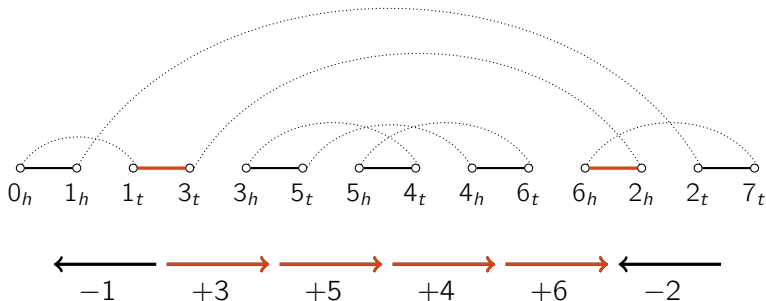


# Sorting Example

$$\pi = (0 \quad 3 \quad 1 \quad 6 \quad 5 \quad -2 \quad 4 \quad 7)$$

# Sorting Unoriented Components

- Let's analyse the effect that reversals have on cycles of  $BP(\pi)$ .
- Reversals change # of cycles by  $-1$ ,  $0$ , or  $+1$ .
- What happens exactly when we apply a reversal **defined** by two black edges?

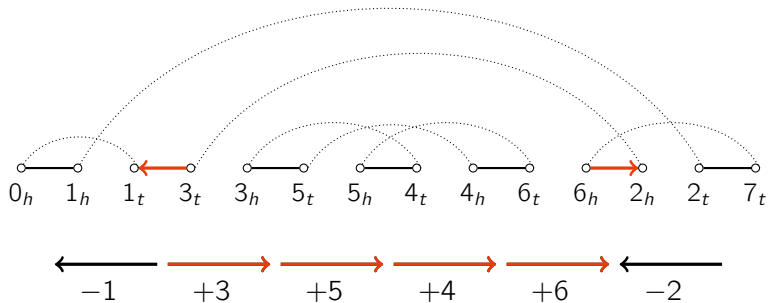


# Reversals and effect on cycles

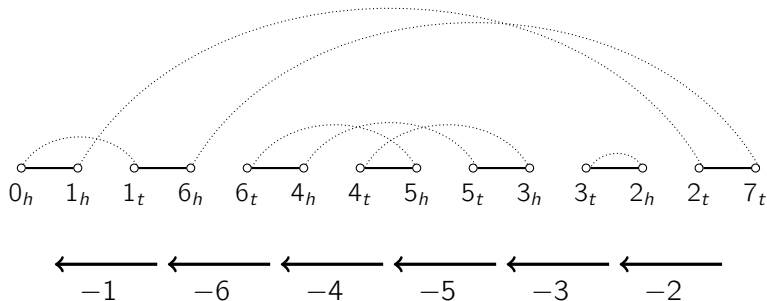
- 1 Edges are on the **same cycle**:
  - **Type I**: Divergent edges: breaks the cycle.  $\Delta C = +1$ .
  - **Type II**: Convergent edges:  $\Delta C = 0$ , may change cycle orientation.
- 2 Edges on **different cycles**:
  - **Type III**: Merges the two cycles.  $\Delta C = -1$ .

So far, we only used **Type I** operations, to sort oriented components.

# Type I - Same Cycle, divergent

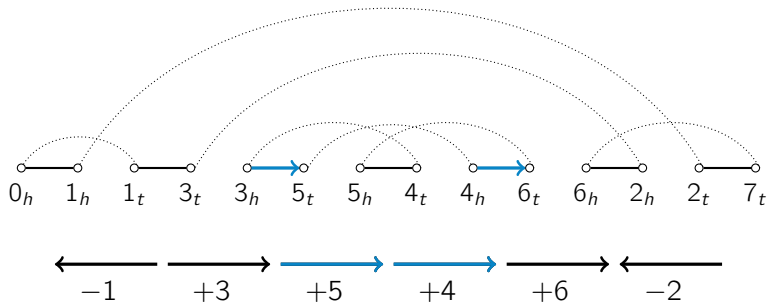


## Type I - Same Cycle, divergent

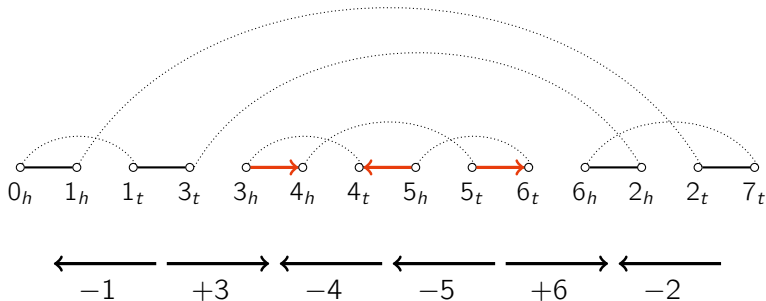


This reversal increases the number of cycles by one,  $\Delta C = +1$ .

## Type II - Same Cycle, convergent

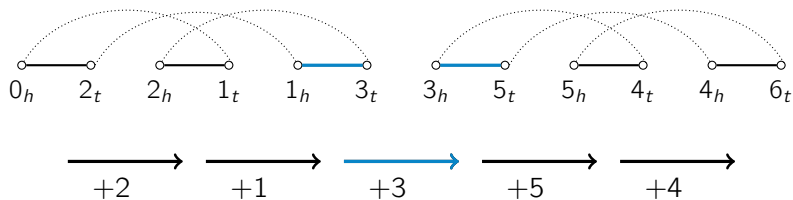


## Type II - Same Cycle, convergent



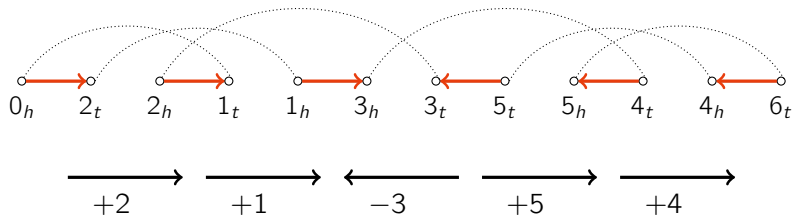
Does not change number of cycles ( $\Delta C = 0$ ), but the cycle is **oriented**.

# Type III - Different Cycles





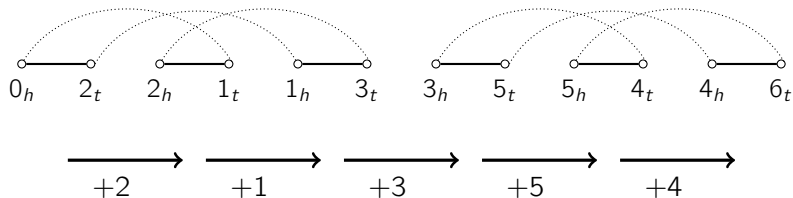
## Type III - Different Cycles



Merges the two cycles, decreasing the number of cycles by one ( $\Delta C = -1$ ), but the new cycle is **oriented**.

# Extra Operations

- How many extra operations do we need to sort unoriented components?



java InversionVisualisation L2/2unoriented.txt

## Extra Operations

- Applying one reversal in each cycle, orients both cycles, with 2 extra operations:

$$d = N - C + 2$$

- Applying one reversal merging both cycles, creates one new **oriented** cycle. Only one operation, but also one less cycle:

$$d = N - (C - 1) + 1 = N - C + 2$$

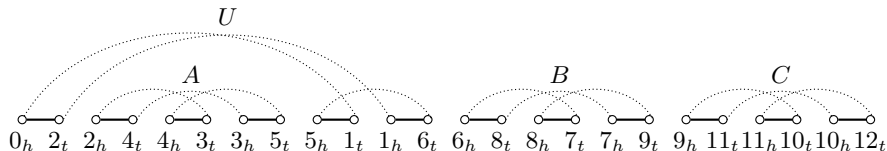
- In both cases, 2 extra operations. Does this mean that

$$d = N - C + K$$

where  $K$  is the number of unoriented components? **Almost...**

# Definitions

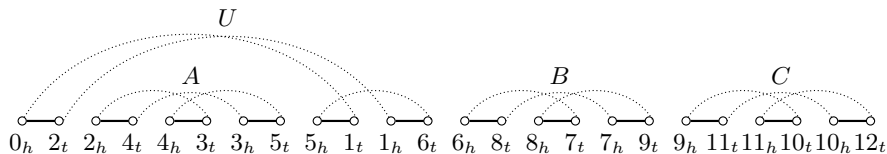
- A Component  $U$  **separates** two other components  $A$  and  $B$  if any edge from a vertex from  $A$  to  $B$  would cross an edge of  $U$ .



- $U$  separates  $A$  and  $B$ . (Also  $A$  and  $C$ ).

# Definitions

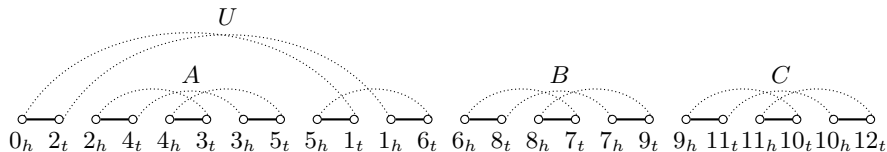
- A **hurdle** is an unoriented component that does **not** separate other two unoriented components.



- $A, B$  and  $C$  are hurdles.

# Definitions

- A **super-hurdle** is a hurdle that, when removed, causes the creation of a new hurdle.



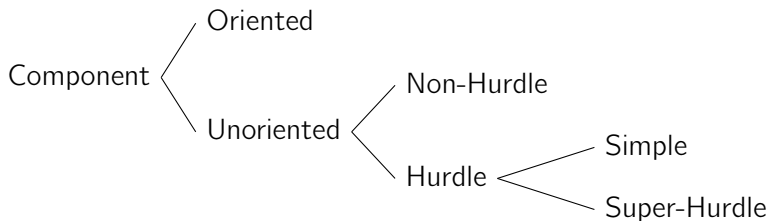
- $A$  is a super-hurdle.  $B$  and  $C$  are called *simple* hurdles.

- Why are these definitions important? Because except for one very rare special case, we have

$$d = N - C + H$$

where  $H$  is the number of hurdles.

# BP Graph – Component Types





# Reversal Types

- **Type I: Oriented Reversal:**  $\Delta C = +1$ .
  - Edges on same cycle, divergent.
- **Type II: Hurdle Cutting:**  $\Delta C = 0, \Delta H = -1$ .
  - Edges on same cycle (hurdle), convergent.
- **Type III: Hurdle Merging:**  $\Delta C = -1, \Delta H = -2$ .
  - Edges on different cycles (hurdles).

# Separating component

- Why a separating component is not a Hurdle?
- Because it can be oriented by a **Hurdle Merging** of two hurdles that is separates.

```
java InversionVisualisation L2/sep-hur-example.txt
```

# Super-hurdles: Problems might occur

- Cutting a super-hurdle is bad.
- Merging hurdles that are separated from a super-hurdle can cause the separating component to become a hurdle.

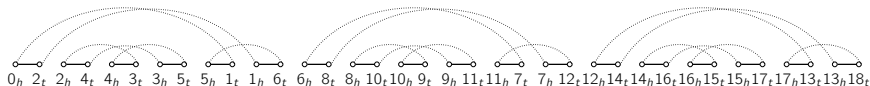
```
java InversionVisualisation L2/sep-hur-example.txt
```

## Super-hurdles: Problems might occur

- How to avoid those problems?
- When there is an odd # of hurdles, cut a simple hurdle.
- When there is an even # of hurdles, merge opposite hurdles.
- Can we always do that? No... meet the **fortress**!

# Fortresses

- A **fortress** is a permutation that has an odd number of hurdles, and all are super-hurdles.



In this kind of permutation, there is no way to avoid an **extra operation**, a hurdle cut that creates a new hurdle.

```
java InversionVisualisation L2/fortress.txt
```

# Reversal Distance - Complete equation

## Theorem (Reversal Distance, HP 95)

*The reversal distance of a permutation  $\pi$  is given by*

$$d(\pi) = N - C + H + F$$

*where:*

- *$N$  is the number of genes*
- *$C$  is the number of cycles in  $BP(\pi)$*
- *$H$  the number of hurdles in  $BP(\pi)$*
- $F = \begin{cases} 1, & \pi \text{ is a fortress} \\ 0, & \text{otherwise} \end{cases}$

# Reversal Distance - Complete Algorithm

```
1: procedure ReversalSort( $\pi$ )
2:   while  $\pi \neq$  identity do
3:     if  $\exists$  oriented component in  $BP(\pi)$  then
4:        $\rightarrow$  Apply a max score oriented reversal – Type I
5:     else if even # of hurdles then
6:        $\rightarrow$  Apply a Hurdle Merging on opposite hurdles – Type III
7:     else if  $\exists$  simple hurdle then
8:        $\rightarrow$  Apply a Hurdle Cutting on a simple hurdle – Type II
9:     else
10:       $\rightarrow$  Merge any two super hurdles (Fortress)
11:    end if
12:  end while
13: end procedure
```