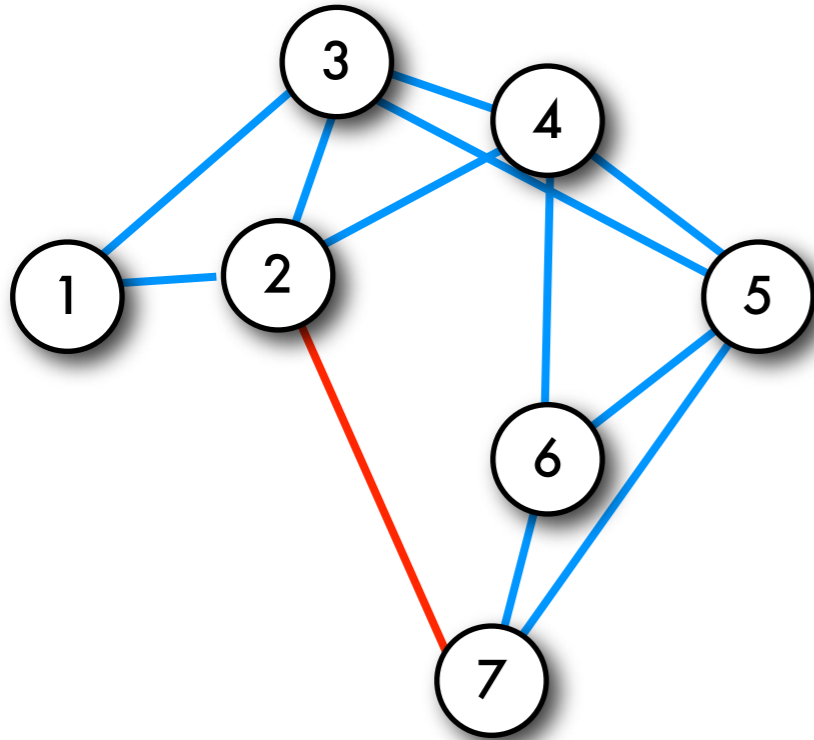


The Interval Ordering Problem

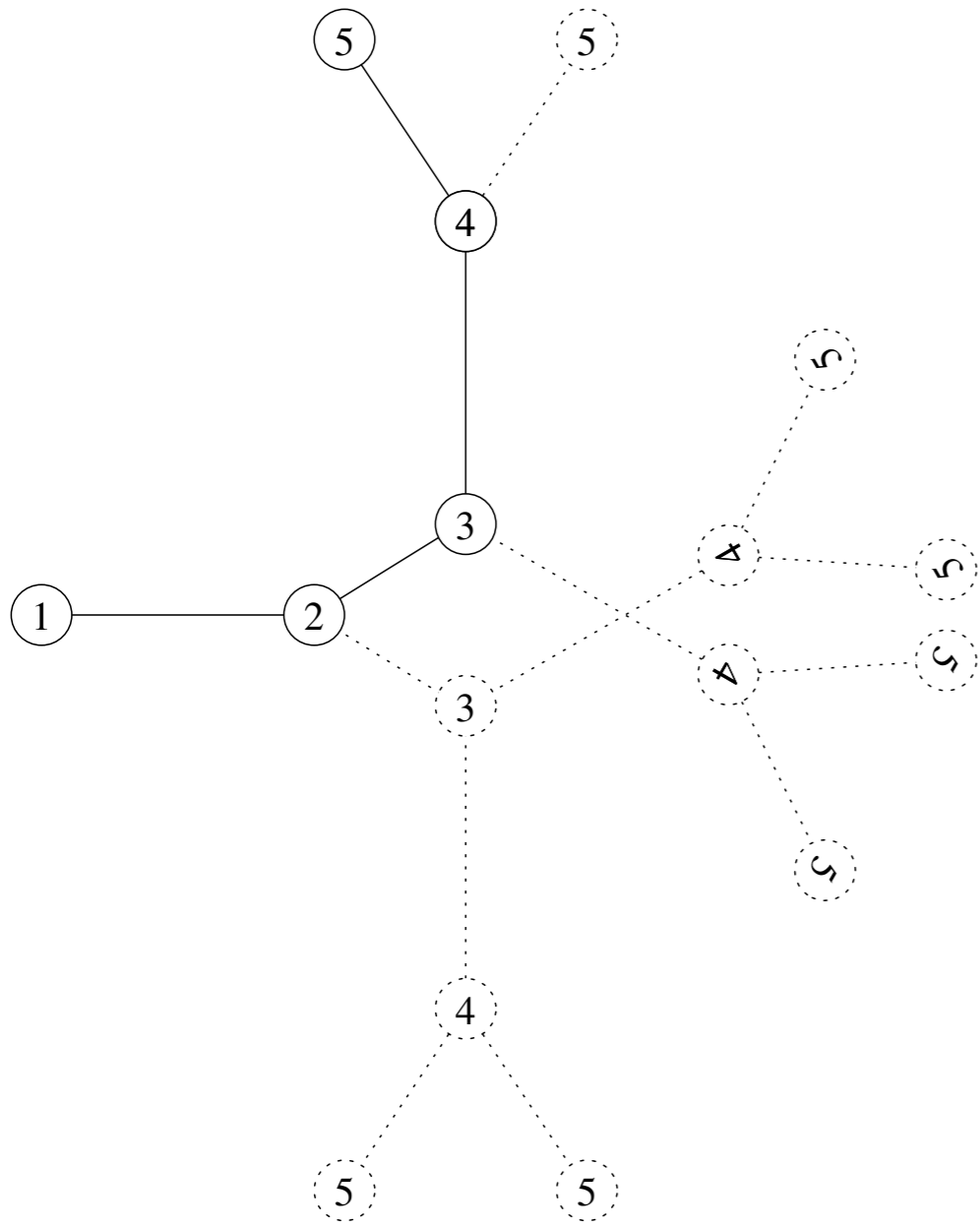
with Maurice Queyranne, Frits Spieksma, Fabrice Talla Nobibon, Gerhard Woeginger

A motivation



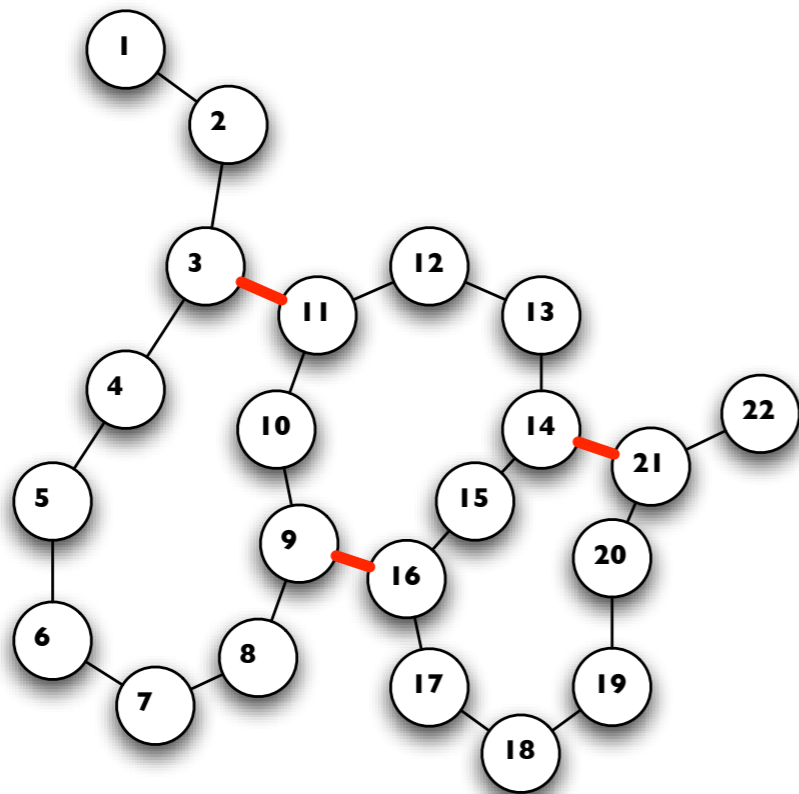
- A molecule consisting of atoms $1, \dots, n$ in unknown positions
- we are given the **distances** between all atom pairs $(i, i+1)$ and $(i, i+2)$
- and the **distances** between some other atom pairs

a combinatorial structure

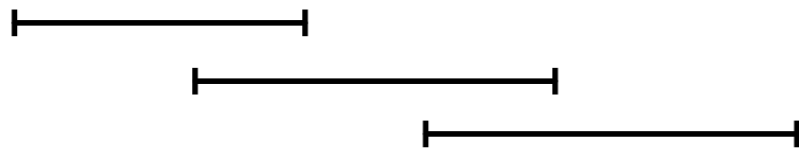


- with the **distances** $(i-1,i)$ and $(i-2,i)$, there are 2 possible positions for atom i relative to atoms $i-1,i-2$.
- A binary string describes all valid embeddings
- the other **distances** are constraints on substrings

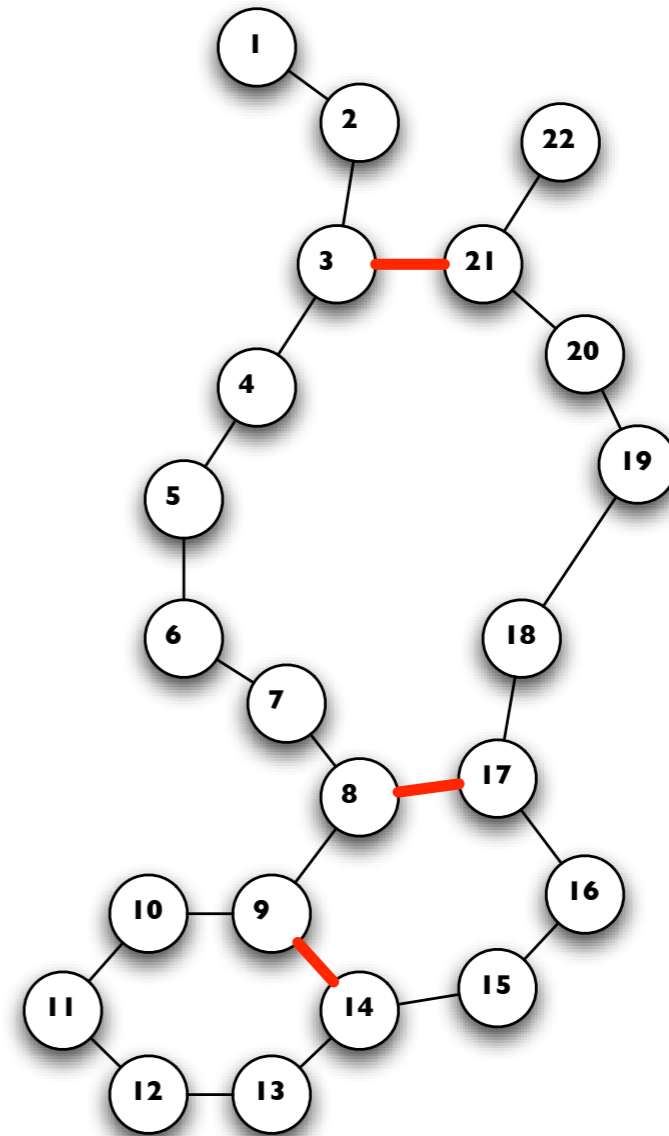
special interval families



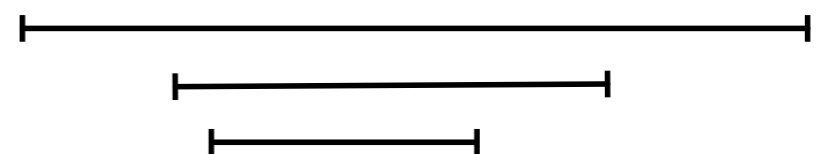
1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22



agreeable intervals



1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22

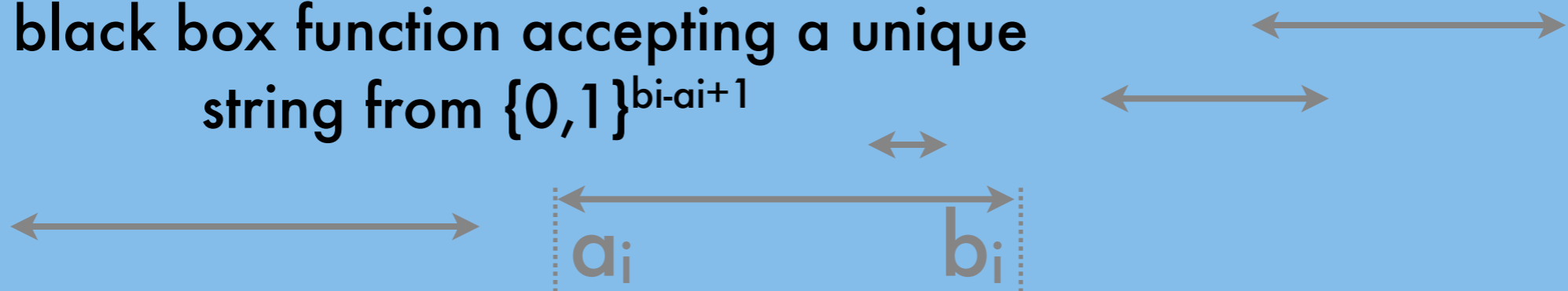


laminar intervals

The Bit String Reconstruction Problem

input $(a_1, b_1, T_1) \dots (a_n, b_n, T_n)$

T_i : black box function accepting a unique string from $\{0, 1\}^{b_i - a_i + 1}$



$y =$

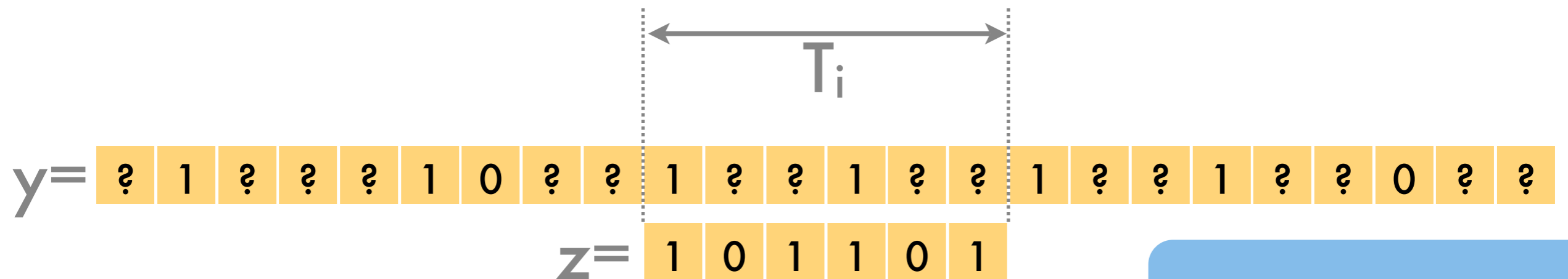
0	1	0	0	1	1	0	0	1	1	0	0	1	1	0	1	0	0	1	1	0	0	1	1
---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---	---

such that $\forall i : T_i$ accepts the substring $y[a_i \dots b_i]$

output

(or announce inconsistency)

The Brute Force Search



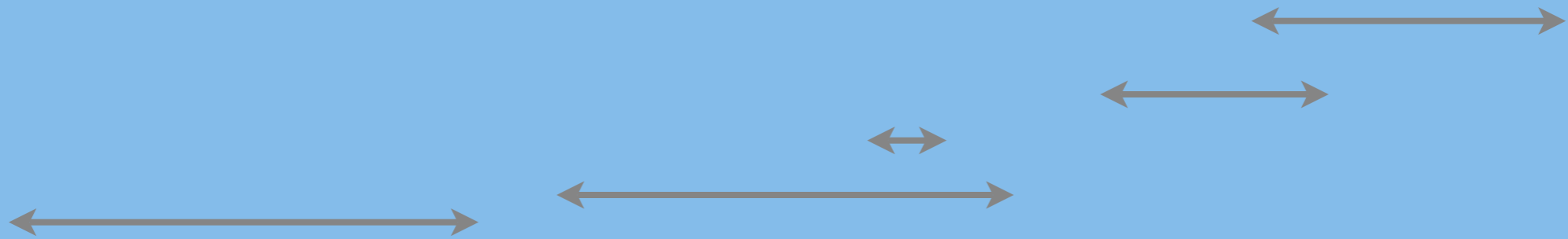
- $y \in \{?, 0, 1\}^m$, start with $y = ?^m$
- for all constraints (a_i, b_i, T_i) in **some order**
 - $w = y[a_i, b_i]$, $k = \text{number of ? in } w$
 - try all 2^k substitutions of ? by 0 or 1 in w
 - until we find a z accepted by T_i
 - replace in y the portion $y[a_i, b_i]$ by z

which order
on those
constraints
leads to the
smallest
running
time ?

The Interval Ordering Problem

fixed a function f

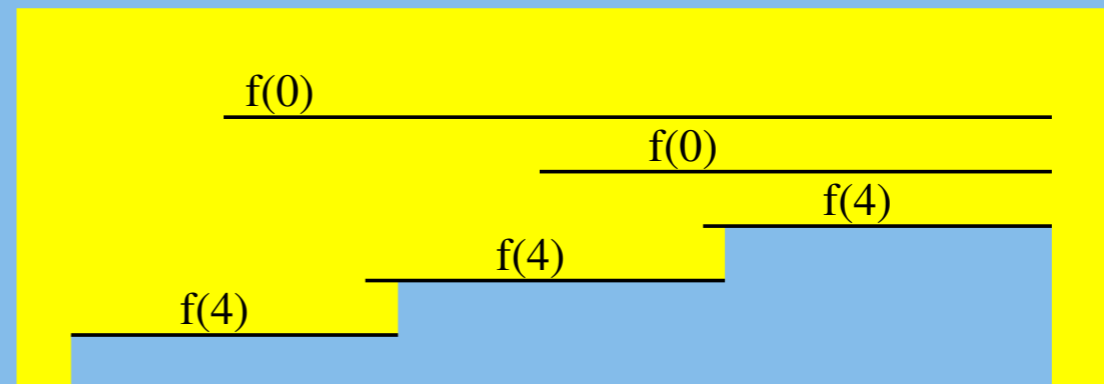
input n intervals I_1, \dots, I_n



output an order on these intervals

minimizing

$$\sum_k f(I_k \setminus (I_1 \cup \dots \cup I_{k-1}))$$

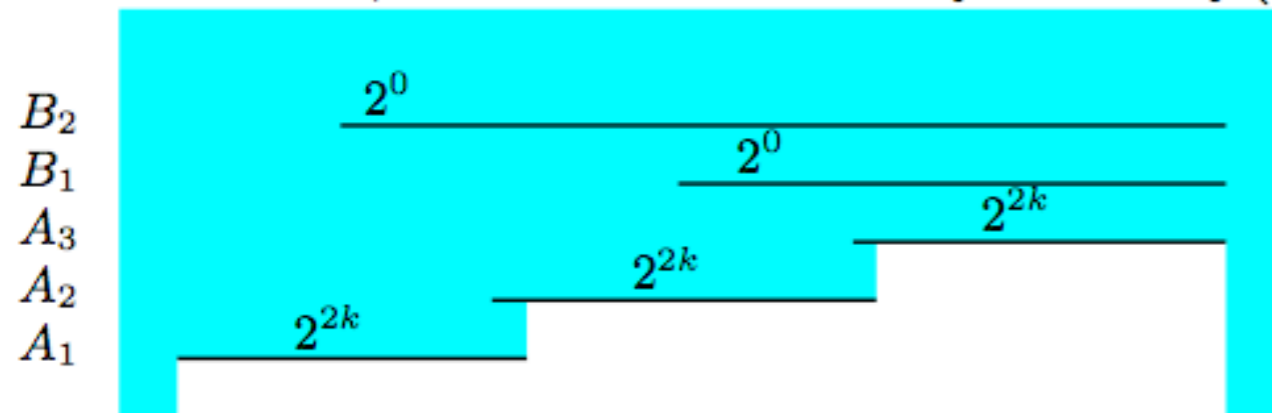


greedy ordering with respect to exposed length

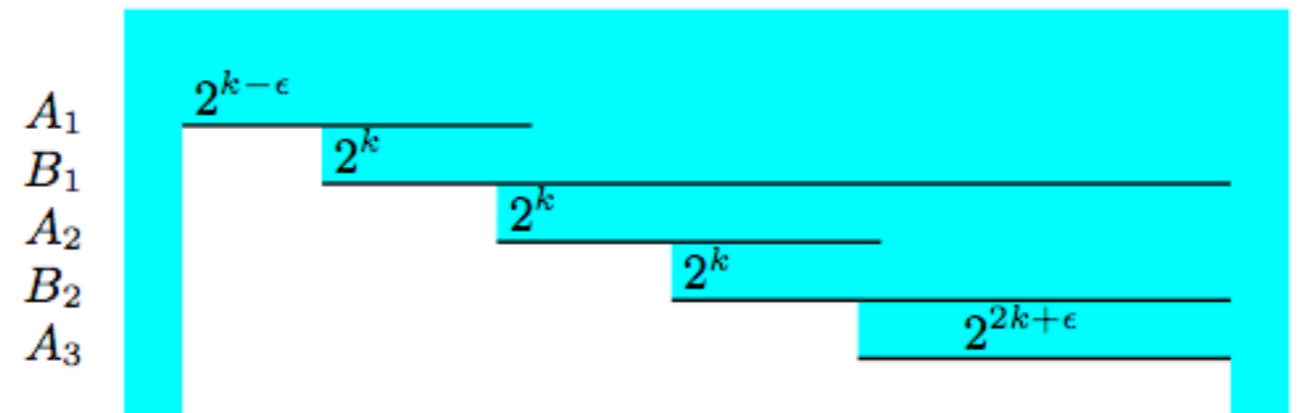
How bad is the greedy ordering?

for $f(x)=2^x$...

Example 2. Consider a family of instances, where each instance consists of $2k-1$ intervals: $A_1 = [0, 2k)$, $A_2 = [2k - \epsilon, 4k)$, $A_3 = [4k - \epsilon, 6k)$, ..., $A_k = [2k(k-1) - \epsilon, 2k^2)$, $B_1 = [k - \epsilon, 2k^2)$, $B_2 = [3k - \epsilon, 2k^2)$, $B_3 = [5k - \epsilon, 2k^2)$, ..., $B_{k-1} = [2k^2 - 3k - \epsilon, 2k^2)$, for some constants $k, \epsilon > 0$ with the cost function $f(x) = 2^x$.



greedy ordering with respect to exposed length



optimal ordering

A greedy sequence is $(A_1, A_2, \dots, A_{k-1}, A_k, B_{k-1}, B_{k-2}, \dots, B_1)$ and achieves a cost of $k2^{2k} + k - 1$, whereas the optimal solution is $(A_k, B_{k-1}, A_{k-1}, B_{k-2}, \dots, A_2, B_1, A_1)$ and has the cost of $2^{2k+\epsilon} + (2k-3)2^k + 2^{k-\epsilon}$. The ratio between both costs can be made arbitrarily large, by choosing appropriate k and small $\epsilon > 0$.

... arbitrary bad!

What do we know?

- f arbitrary, (I) agreeable : dynamic programming in $O(n^3)$
 - f continuous, convex, (I) agreeable : dynamic pro. in $O(n^2)$
 - $f(x)-f(0)$ sub-additive, (I) laminar : inner to outer in $O(n \log n)$
 - variant: minimize $\max_k f(\text{exposed part of } I_k)$: greedy in $O(n^2)$
 - $f(x)=2^x$, (I) arbitrary : open
- +
- f arbitrary, I arbitrary cannot be constant approximated (unless $P=NP$)
-