

An experimental investigation on the pancake problem

Bruno Bouzy

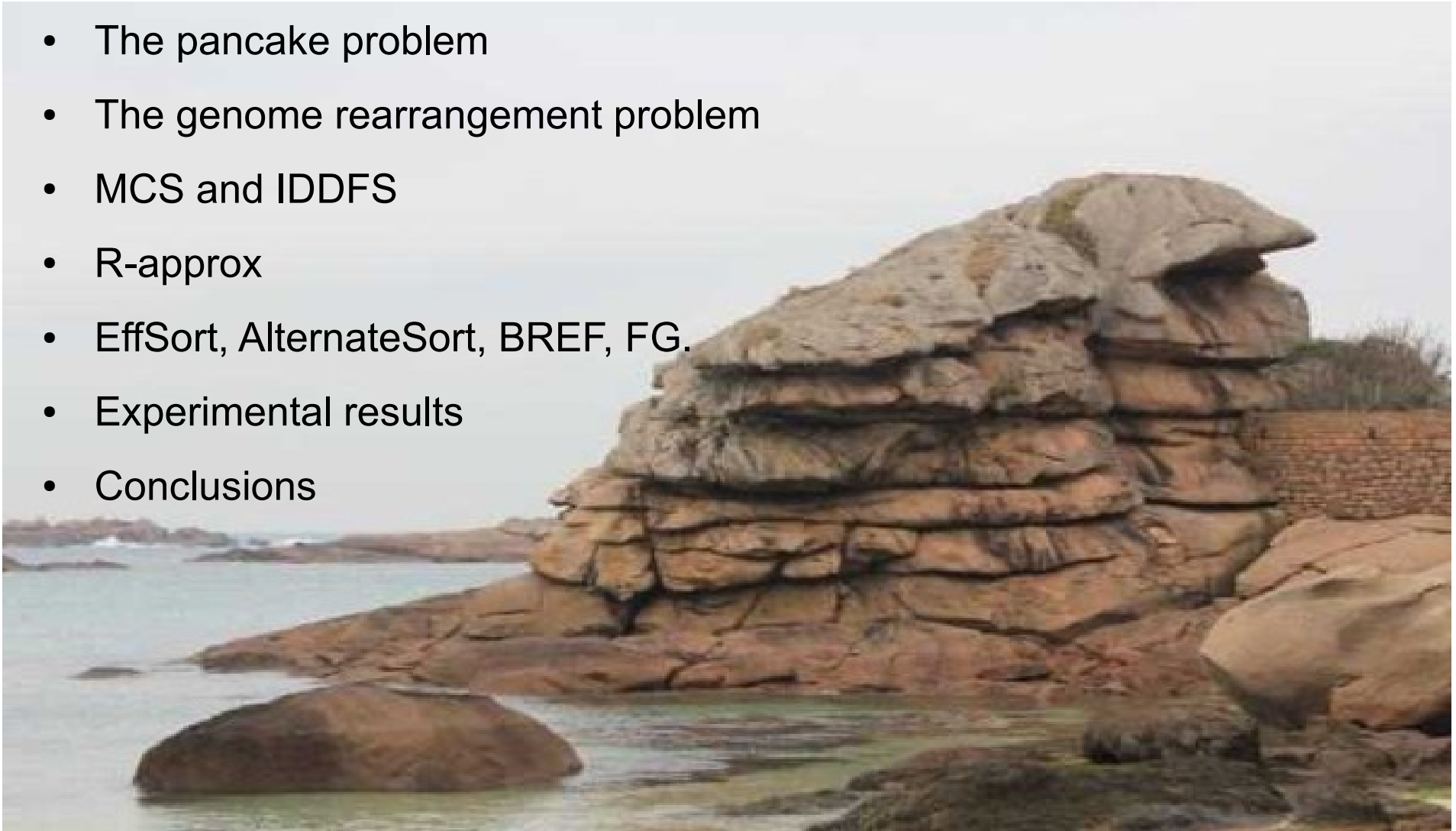
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Outline

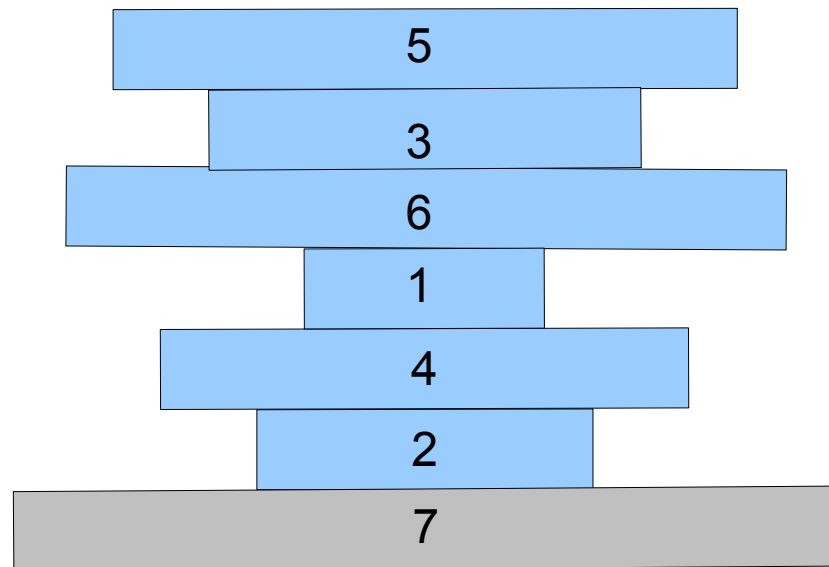
- The pancake problem
- The genome rearrangement problem
- MCS and IDDFS
- R-approx
- EffSort, AlternateSort, BREF, FG.
- Experimental results
- Conclusions



Example

A stack of pancakes.

Each pancake has a size.

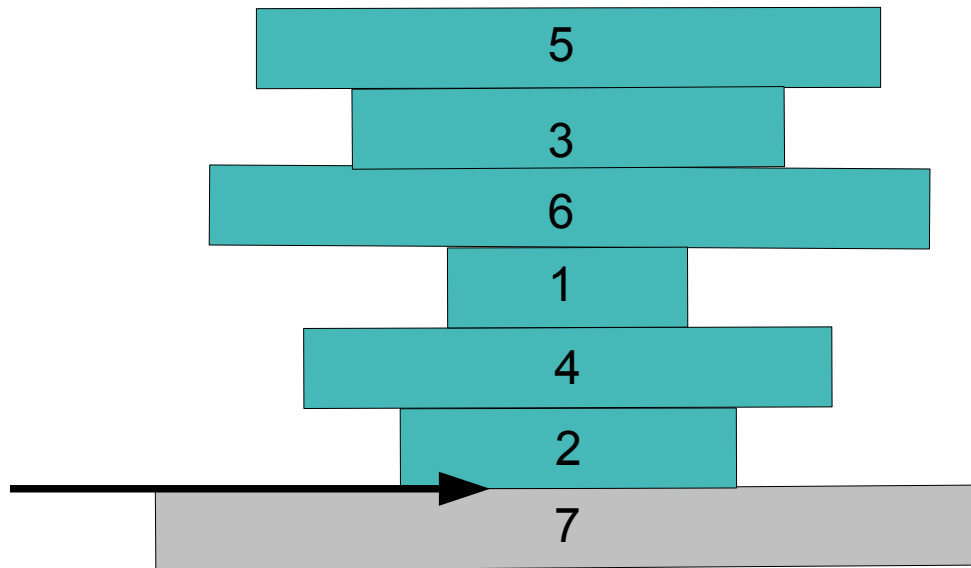


Goal : sort the stack with the greatest pancake at the bottom.

Example

Action :

insert a spatula below a pancake and flip the sub-stack above the spatula.

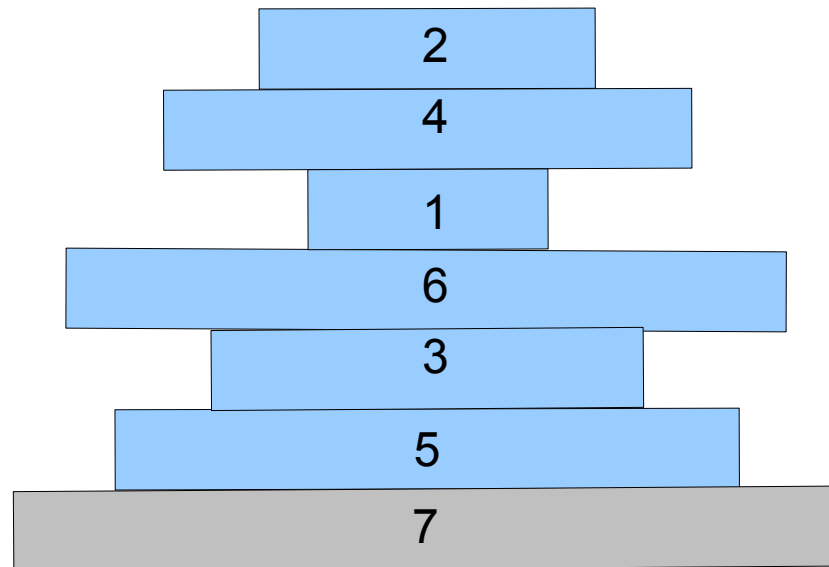


Example

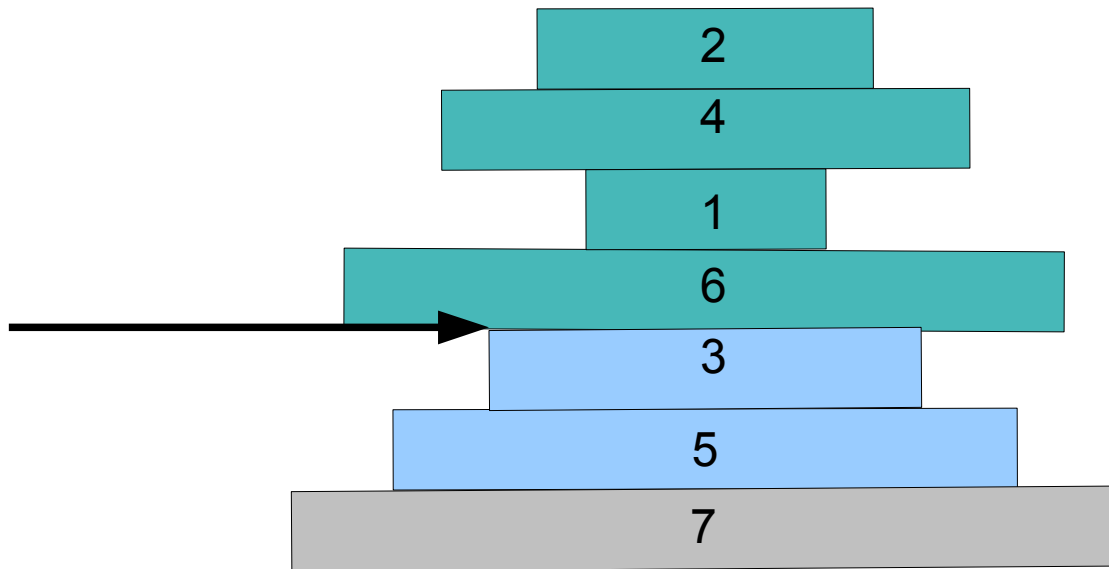
6-flip !

Pancake 5 is now at the bottom.

Pancake 2 is now at the top.

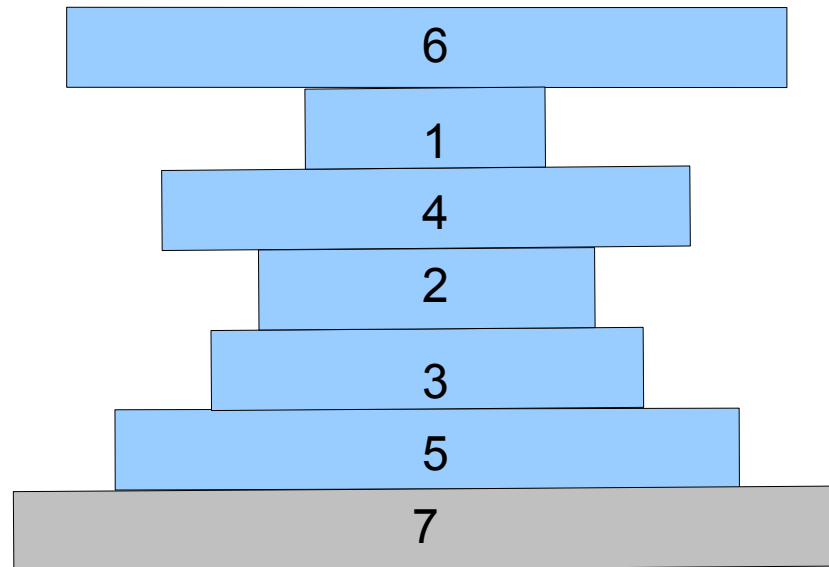


Example

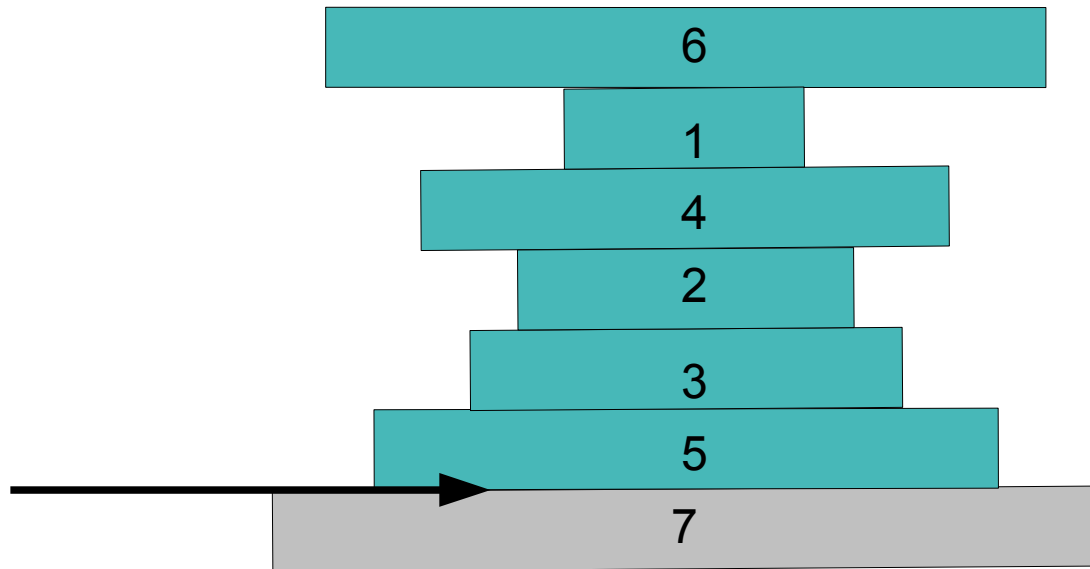


Example

4-flip !

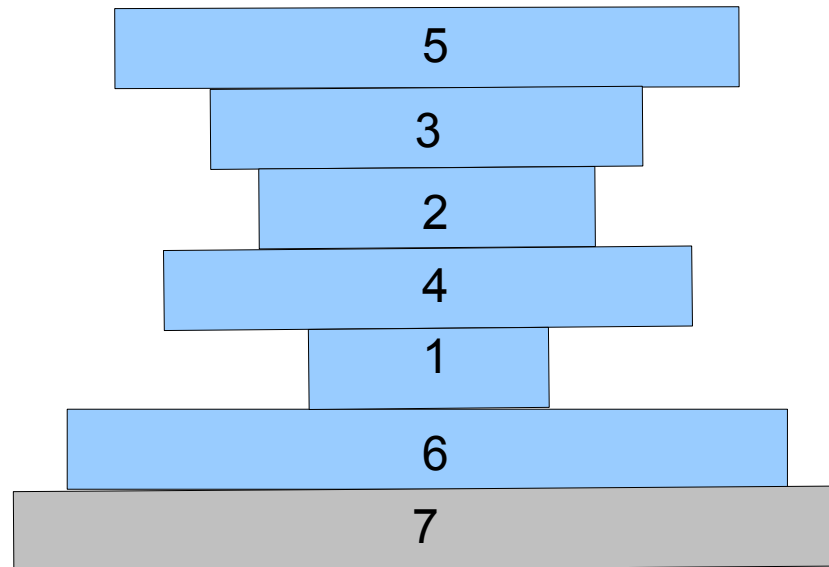


Example

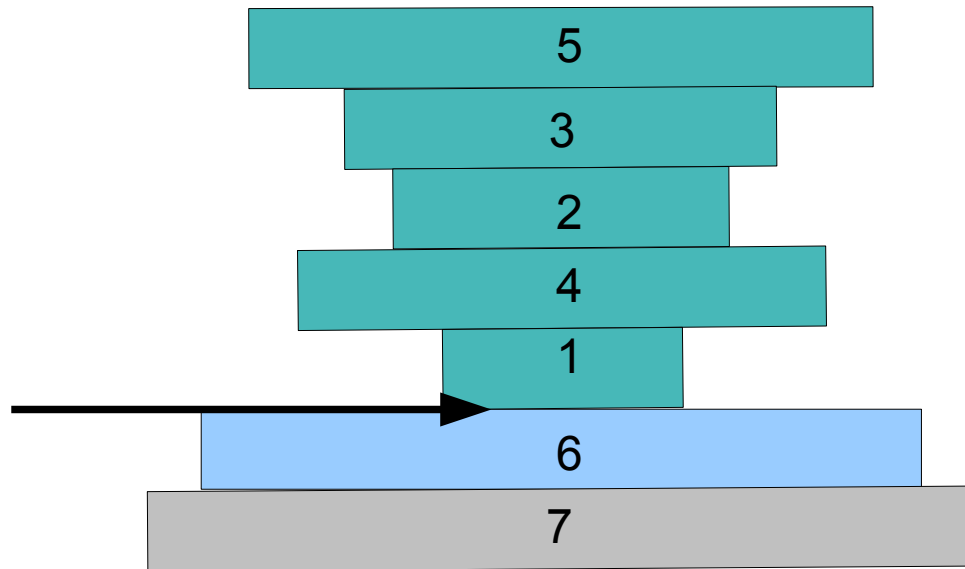


Example

6-flip !

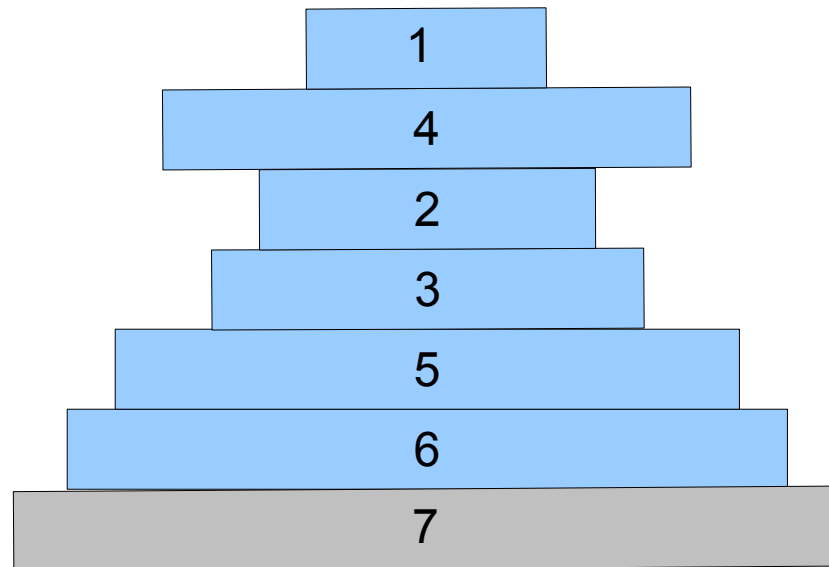


Example

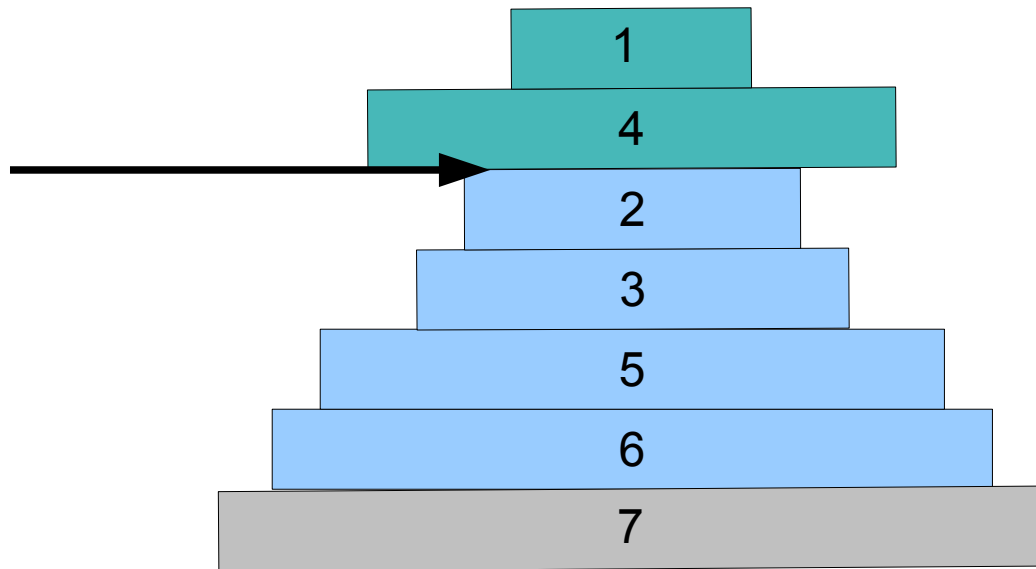


Example

5-flip !

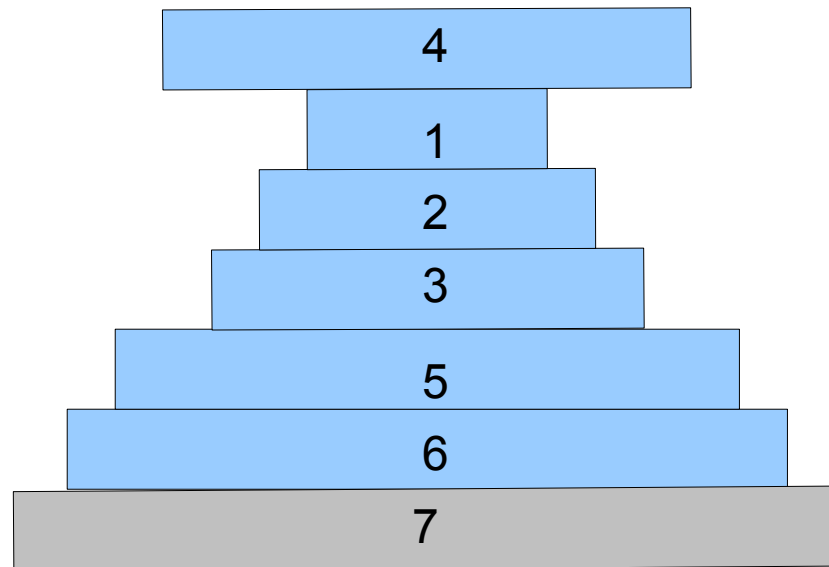


Example

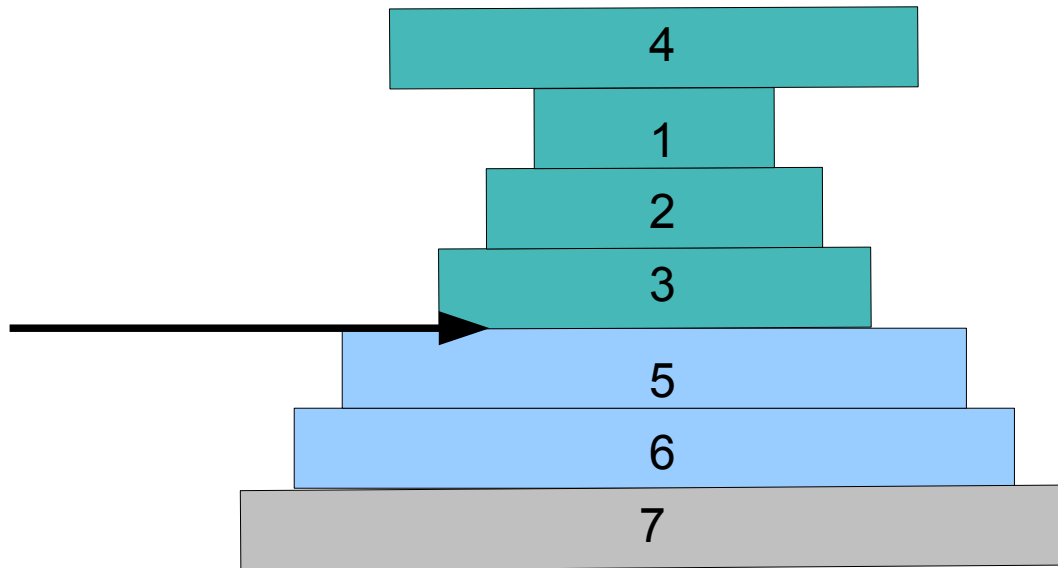


Example

2-flip !

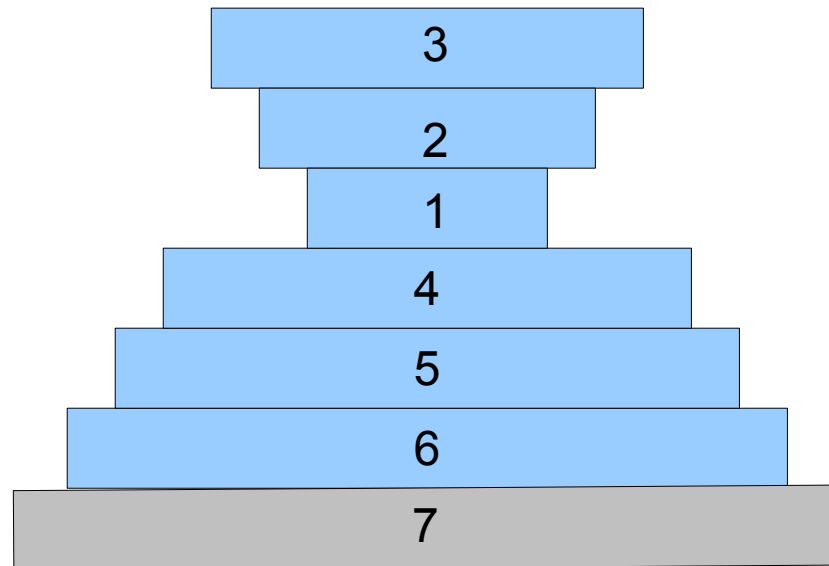


Example

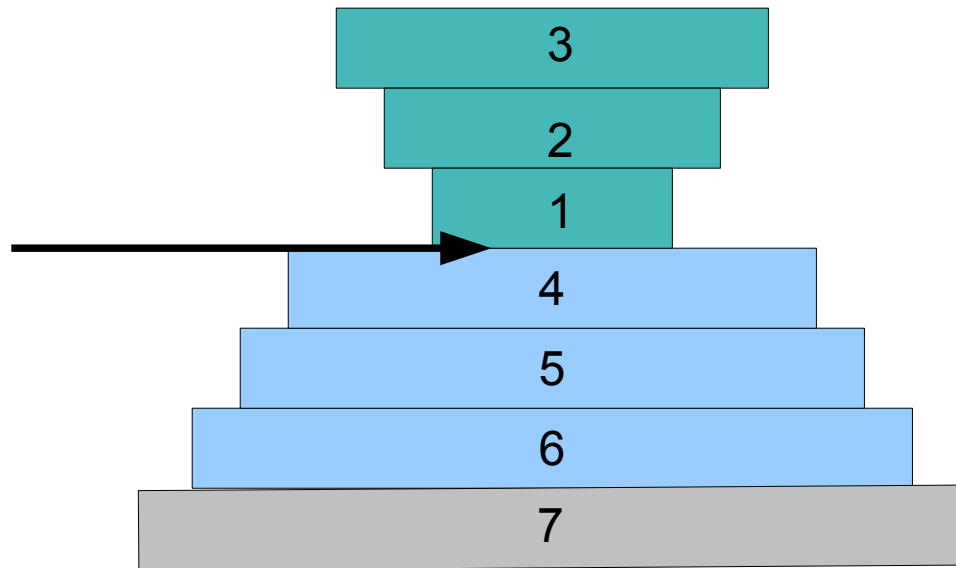


Example

4-flip !

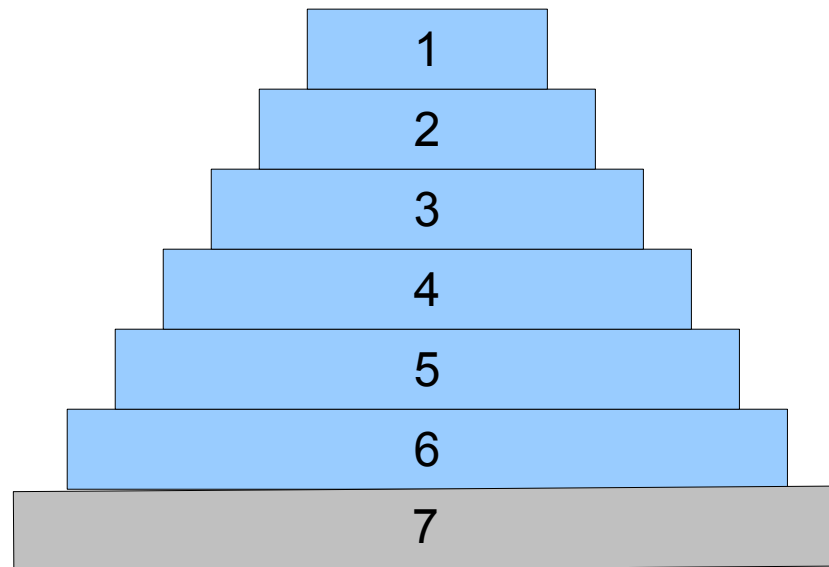


Example



Example

3-flip !



Completed in 7 moves!

3 Linked problems

- Burnt pancake problem :
 - Each pancake is burnt on one side
 - Goal : sorted stack + burnt side down.
- Genome rearrangement problem (Hayes 2007)
 - Action with two spatulas
 - Signed version : each gene has sign (+-1)
 - Unsigned version otherwise
 - Great interest in biology to classify species :
distance(cabbage, turnip) = 3 !

4 classes of problems

#spatulas signed	1	2
yes	Burnt Pancake	Signed Genome
no	Unburnt Pancake	Unsigned Genome

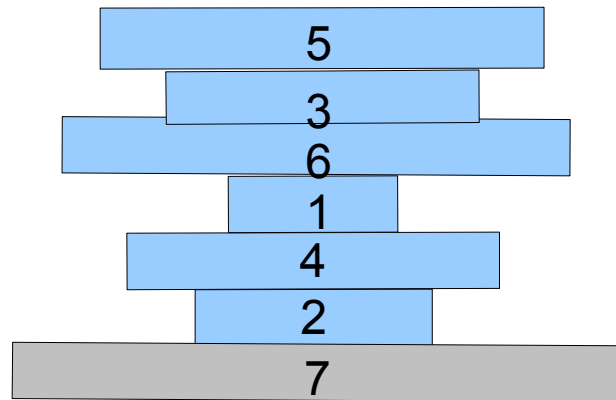
Complexities

- Unsigned genome : **NP-hard** (Caprara 1997)
- Signed genome : **Polynomial** :
 - n^4 (Pevzner & Hennenhalli 1995)
 - n^2 (Bergeron 2005)
 - n (Bader & al 2001) without the solution sequence
- Unburnt pancakes : **NP-hard** (Bulteau & al 2012)
- Burnt pancakes : **?**
 - Polynomial on « simple » instances (Labarre & Cibulka 2011)
 - A study (Cohen & Blum 1995)
 - Another study (Cibulka 2011)

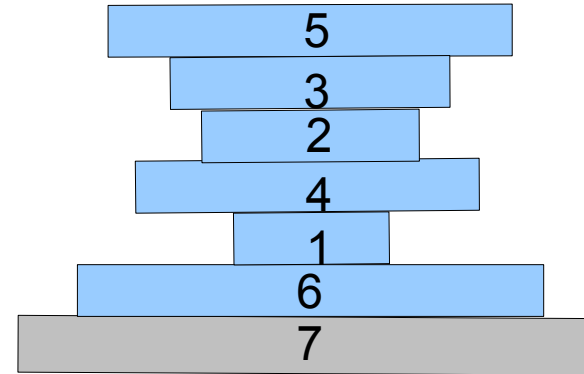
#breakpoints (#bp)

- **Unsigned (unburnt) version :**
 - A breakpoint is situated between two adjacent pancakes (genes) when the difference between the sizes of the two pancakes (genes) is not 1 or -1.
- **Signed (burnt) version :**
 - A breakpoint is situated between two adjacent pancakes (genes) when the difference between the sizes of the two pancakes (genes) is not 1.
- **#breakpoints :**
 - admissible heuristic for the pancake problem (Gates Papadimitriou 1979 ; Helmert 2010)
- **#breakpoints/2 :**
 - admissible heuristic for the genome rearrangement problem

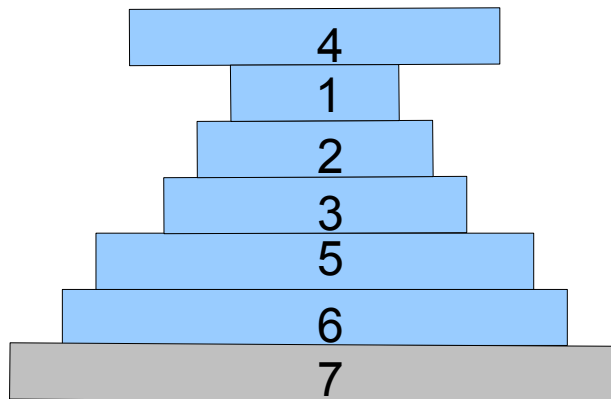
#breakpoints (#bp)



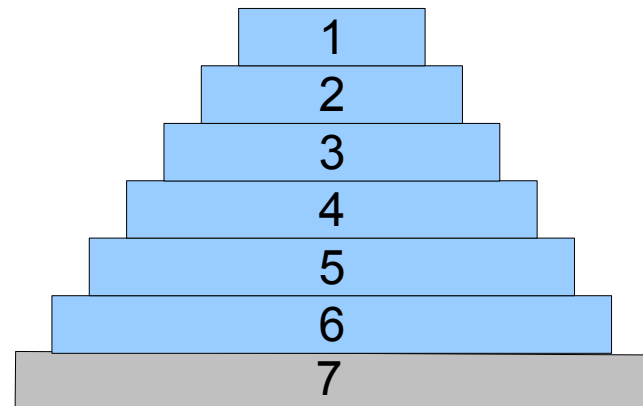
#bp=6



#bp=4



#bp=2



#bp=0

The unburnt pancake problem

- N-pancake problem
 - Stacks of size N
 - Solving a **specific instance**
 - Finding the **diameter** of the graph
- **Lower** and **Upper bounds** on the diameter of the graph
 - Gates & Papadimitriou 1995 : $B^- = (17/16)n$
 - Heydari & Sudborough 1997 : $B^- = (15/14)n$
 - Chitturi & al 2009 : $B^+ = (18/11)n$

The unburnt pancake problem

- **NP-hard** to efficiently sort (Bulteau & al 2012)
 - At most 2 « efficient » moves : binary search.
- **Efficient Sort (EffSort)** (Bulteau & al 2012)
 - Only consider the « efficient » moves (that strictly decrease #bp)
 - And not the « waste » moves (that keep #bp constant)

- **Alternate Sort (AS)** (current work)

```
do {  
    efficient sort (sequence) ; play the sequence ;  
    if not completed, play a waste move ;  
} while not completed ;
```

- **Fixed-Depth Alternate Sort (FDAS)** (current work) :

Alternate Sort using fixed-depth EffSort

The unburnt pancake problem

- The pancake challenge (2004)
 - [Tomas Rokicki](#) entry
 - [Backward Search](#) : the reverse sequence of a solution to P_i is a solution to P_i^{-1}
 - Building sequences hard to solve.
 - Other clever tricks
- [2-approximation algorithm](#) (Fischer & Ginzinger 2005)
 - Type 1, 2, 3, 4 moves
 - $R\text{-approx} = L / \#bp$
 - ≤ 2 (theory)
 - ≈ 1.22 on average (practice)

Two general tools

- **Iterative Deepening Depth-First Search (IDDFS)** (Korf 1985)
 - Needs an Heuristic = #breakpoints (Helmert 2010)
 - Solves random instances of size 60 or 80 in a few seconds on average
 - Cannot solve some specific complex stacks of size 30.
 - Time to solve size N stacks : exponential in N
 - While not completed, a lower bound is provided
 - Optimal solution at completion
- **(Nested) Monte-Carlo Search ((N)MCS)** (Cazenave 2009)
 - Needs domain-dependant simulations
 - Solves random instances of size up to 512.
 - The time to solve a stack is polynomial in N
 - While not completed, an upper bound is provided
 - Stops only if the optimal length == heuristic value
 - Otherwise, no way to see if the solution is optimal

Domain-dependant tools

- Simulators
 - Fixed-Depth Alternate Sort
 - Fischer and Ginzinger (**FG**) algorithm
 - **BREF**

```
while not completed {  
    choose an efficient move at random ;  
    If (success) play it ;  
    else { choose a waste move at random ; play it ; }  
}
```
- Backward Solutions (**BS**) (Rokicki 2004)

Settings of experiments

- For a **specific problem**, finding an optimal solution.
- For a **set of X problems** drawn at random
 - R-approx = $L / \#breakpoints$
 - Less than one hour to solve a set a problems
 - Standard deviation of r-approx on one problem = 0.05
 - ==> 2-sigma rule : precision of $(0.05 \cdot 2 / 10 =) 0.01$ on r-approx averaged on 100 problems

Results of experiments (1)

- IDDFS is **optimal** and $R > 1$? (normal because #bp is a lower bound)
- IDDFS cannot solve instances of size > 64
- MCS can solve instances of sizes up to **256**
- For $N \leq 64$, MCS is **inferior** to IDDFS

		IDDFS		NMCS		+ BREF	
N	L	R	T	L	R	T	level
8	6.5	1.09	0	6.5	1.09	0	4
16	14.5	1.05	0	15.5	1.09	0.05	4
32	31.0	1.03	0	36.5	1.21	0.6	3
64	63.0	1.02	5	76	1.24	1.2	2
128	–	–	∞	163	1.29	0.3	1
256	–	–	∞	333	1.31	9	1

Results of experiments (2)

- BS enhancement : $l = \min(l_1, l_2)$
 - l1 = length of the forward simulation
 - l2 = length of the **backward simulation** on the inverse problem
- Left : MCS + BREF
- Right : **MCS + BREF + BS**

		NMCS			+BREF			+ BS		
N	L	R	T	level	L	R	T	level		
8	6.5	1.09	0	4	6.6	1.10	0	4		
16	15.5	1.09	0.05	4	15.3	1.09	0.1	4		
32	36.5	1.21	0.6	3	35.2	1.15	0.8	3		
64	76	1.24	1.2	2	75.2	1.20	2	2		
128	163	1.29	0.3	1	159	1.26	0.5	1		
256	333	1.31	9	1	335	1.32	0.1	0		

Results of experiments (3)

- Fischer and Ginzinger (FG)
- FG simulation (R-approx=2 in theory and 1.22 in practice)
- Left : MCS + BREF + BS
- Right : MCS + FG + BS

		NMCS	+BREF	+ BS		NMCS	+FG	+ BS
N	L	R	T	level	L	R	T	level
8	6.6	1.10	0	4	6.7	1.10	0	4
16	15.3	1.09	0.1	4	14.8	1.05	0.6	4
32	35.2	1.15	0.8	3	31.6	1.05	4.2	3
64	75.2	1.20	2	2	68.5	1.10	2.5	2
128	159	1.26	0.5	1	155	1.23	0.8	1
256	335	1.32	0.1	0	336	1.32	0.2	0

Results of experiments (4)

- Fixed-depth Alternate Sort (FDAS)
- Left : MCS + FG + BS
- Right : MCS + FDAS
- Solving size 512 stacks

N	MCS + FG + BS				MCS + FDAS			
	L	NMCS R	+FG T	+ BS level	L	NMCS R	+ FDAS T	level
8	6.7	1.10	0	4	6.6	1.09	0	3
16	14.8	1.05	0.6	4	14.5	1.04	0.04	3
32	31.6	1.05	4.2	3	31.3	1.04	0.8	3
64	68.5	1.10	2.5	2	65.1	1.04	18	3
128	155	1.23	0.8	1	139	1.10	10.5	2
256	336	1.32	0.2	0	300	1.18	1.5	1
512	—	—	—	—	614	1.20	9	1

Conclusions

- The (unburnt) pancake problem is experimentally [revisited](#).
- Associated with :
 - [BREF](#), [FG](#), [FDAS](#) as simulators,
- MCS extends the results obtained by IDDFS
 - Sizes of stacks [up to 512](#) with $r\text{-approx}=1.20$
 - $R\text{approx} = 1.4$ for N in $[16, 64]$
 - $R\text{approx}$ in $[1.10, 1.20]$ for N in $[128, 512]$

Future work

- The **burnt** pancake problem
 - IDDFS : Improve the Cibulka's **heuristic function**
 - MCS : assess **Cohen & Blum** algorithm as simulator
 - Find new results on the **diameter**
- The **unburnt** pancake problem
 - Find new results on the **diameter**
 - Generate **complex stacks**

Thank you for your attention!

Questions ?

bruno.bouzy@parisdescartes.fr

