

#### School of Computing

# Bill Gates and the Pancake Flipping Problem

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Experience the fun of problem solving

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Pancake Flipping

# **Overview:**

□ When did Bill Gates flip pancakes

**The Pancake Flipping Problem** 

□ What we know about Pancake Flipping

**Why Study Pancake Flipping?** 

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# **Did you know that ...**



#### Bill Gates [比尔 盖茨], formerly Microsoft CEO



# What is pancake flipping?





*Source:* Neil Jones and *Pavel Pevzner*, 2004 "Introduction to BioInformatics Algorithms".

Output: Neat, sorted stack of pancakes

pancake flipping





# **Did Bill Gates** *really* **flip pancakes**?

#### THE PANCAKE PUZZLE

"The chef in our place is sloppy, and when he prepares a stack of pancakes they come out all different sizes. Therefore, when I deliver them to a customer, on the way to the table I rearrange them (so that the smallest winds up on top, and so on, down to the largest at the bottom) by grabbing several from the top and flipping them over, repeating this (varying the number I flip) as many times as necessary. If there are "n" pancakes, what is the maximum number of flips (as a function of "n") that I will ever have to use to rearrange them?"





Source: Neil Jones and Pavel Pevzner, 2004 "Introduction to BioInformatics Algorithms".



by Janet Lowe. (see page 19)

http://www.npr.org/templates/story/story.php?storyId=92236781

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# **Pancake Flipping Problem**



Given an initial pancake configuration... You want to get a *"sorted"* configuration ... Constraints: can *only* flip ...







Example ...



Bill Gates & Christos Papadimitriou:, "Bounds For Sorting By Prefix Reversal." *Discrete Mathematics*, Vol 27, pp 47-57, 1979.



#### More pancake-flipping examples...





### **An Initial Algorithm (Greedy)**

#### Simple Idea:

"Sort" the biggest *unsorted* pancake first...





#### **Greedy Algorithm:**

Repeatedly "sort" the biggest pancake;



#### **Try it Online:**

#### http://www.cut-the-knot.org/SimpleGames/Flipper.shtml





### Is Greedy "the best" possible? Answer: NO

#### **A Counter Example:**

Greedy method [5 flips]



Better way [3 flips]



**Question:** Design an algorithm that solve the pancake flipping problems using the *minimum number of flips*.



#### **Pancake Flipping Problem...**

 $\bigcirc$ 

Sometimes, it is good to look from another perspective!





### A Different Perspective: The Solution Space...



Solution now becomes "a path from source node to destination node" in this solution space.

#### **New Problem:**

Want a smart method (algorithm) to search this space to find the optimal flipping solution.





# **A Search Tree Method:**

(systematically search the search space)



**Problem:** Want a smart method (algorithm) to search this space to find the optimal flipping solution.

# A Search Tree Method:



(systematically search the search space)



**PQ:** Can we work backward from the solution to the data? (In AI, this is called "backward chaining"; the previous search [from start towards destination] is called "forward chaining".)







### Real World (of Pancakes...) vs Model world (of Graph...)

Represent pancake sizes by numbers.





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#### The Model World...





#### ? flips

52143

? flips





#### Let's see both together...





#### Is Greedy "the best" possible? Answer: NO

#### **A Counter Example:**

Greedy method [5 flips]

$$324|1 \rightarrow 4231| \rightarrow 13|24 \rightarrow 312|4 \rightarrow 21|34 \rightarrow 1234$$

Better way [3 flips]

$$32|41 \rightarrow 234|1 \rightarrow 4321| \rightarrow 1234$$

**Question:** Design an algorithm that solve the pancake flipping problems using the *minimum number of flips*.





# A Search Tree Method:

(systematically search the search space)



search this space to find the optimal flipping solution.

Pancake Flipping

# **Overview:**

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### Pancake Flipping Problem: Known Results



- For *n* pancakes, at most (5*n*+5)/3 flips are needed [Bill Gates and Papadimitriou, 1979] ~1.666*n*
- 2008 (about 30 years later), at most 18n/11 needed
   [a team from UT-Dallas, 2008] ~1.6363n (diff: < 2%)</p>
- Pancake flipping problem is an open problem in math and an NP-hard problem in computer science



Pancake Flipping

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□ What we know about Pancake Flipping

□ Why Study Pancake Flipping?

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### Why study pancake flipping

- Mathematics Study its properties
  - define *f*(*n*) to be the minimum of number of flip for any configuration of *n* pancakes
- Computing Want an algorithm to solve it
  - Given any *n* pancake configuration, sort it with the *minimum* number of flips



#### **Math vs Computer Science**

#### **Problem:**

We need to drive from NUS-SOC to Kuala Kubu Bahru in West Malaysia?

#### **Mathematicians:**

#### may give a Theorem

There exist a path with total distance no more than 450km.



### But, a computer scientist wants the exact route (and more)

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>

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from National University of Singapore School of Com. Hon Wai to Kuala Kubu Bharu, Selangor, Malaysia **Raub District** Malaysia Gambang 4 h 58 min (440 km) 🙆 Kuala Kubu Bharu Pekan via Lebuhraya Utara-Selatan/AH2/E2 4 h 20 min without traffic Kuala Lumpur A This route has tolls. Muadzam Shah A This route crosses a country border. National University of Singapore School of 🔶 1 h 15 min Computing Endau 13 Computing Drive, Singapore 117417 Port Dickson Taman Mersing Segamat Negara Johor Get on AYE from Kent Ridge Dr and Clementi Rd District Endau-Rompin 6 min (2.1 km) Bagansiapiapi 🚘 4 h 58 min 440 km Kluang Take Lebuhraya Utara-Selatan/AH2/E2, ELITE/E6 4 Yong Peng and AH2/E1 to Route 3208 in Bukit Beruntung, Batu Pahat Pulau Rupat Rawang, Malaysia. Take exit 118-Bkt. Beruntung from Lebuhraya Utara - Selatan/AH2/E1 Dumai 4 h 1 min (409 km) Pulau Bengkalis Nusajaya National Universi Take Route 1 to Jalan Bukit Kerajaan in Pekan Duri of Singapore Sch Kuala Kubu Bharu, Kuala Kubu Baru Pulau Pedang Pulau Pulau Batam 31 min (29.1 km) Rangsang Kuala Kubu Bharu Minas Fast Selangor, Malaysia Map data @2016 Google, Urban Redevelopment Authority Privacy maps.google.com.sg Send feedback 50 km

Terms



### Why study pancake flipping

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#### Applications

- sorting by prefix reversal
- used to study evolution of species in biology





#### **Application of Sorting by Reversals**

▲HITS

#### **SIAM Journal on Computing**

SIAM J. Comput. / Volume 25 / Issue 2

#### Genome Rearrangements and Sorting by -Reversals

SIAM J. Comput. Volume 25, Issue 2, pp. 272-289 (1996)

Issue Date: 1996

ABSTRACT REFERENCES (28) CITING ARTICLES

Vineet Bafna and Pavel A. Pevzner

Sequence comparison in molecular biology is in the beginning of a major paradigm shift-a shift from gene comparison based on local mutations (i.e., insertions, deletions, and substitutions of nucleotides) to chromosome comparison based on global rearrangements (i.e., inversions and transpositions of fragments). The classical methods of sequence comparison do not work for global rearrangements, and little is known in computer science about the edit distance between sequences if global rearrangements are allowed. In the simplest form, the problem of gene rearrangements corresponds to sorting by reversals, i.e., sorting of an array using reversals of arbitrary fragments. Recently, Kececioglu and Sankoff gave the first approximation algorithm for sorting by reversals with guaranteed error bound 2 and identified open problems related to chromosome rearrangements. One of these problems is Gollan's conjecture on the reversal diameter of the symmetric group. This paper proves the conjecture. Further, the problem of expected reversal distance between two random permutations is investigated. The reversal distance between two random permutations is shown to be very close to the reversal diameter.

**Important Application in Computational Biology:** Used to study the evolution from one species to another.

# **Curious?** See more details in the extra slides at the end.



# **Relevant Skills and Courses**

- Pancake flipping is a *model* for
  - sorting by prefix-reversals
- Many CS problems are model in *similar* ways
  - sending files over internet (routing problems)
  - time table scheduling (graph colouring, 图着色问题)
- Courses to learn these things
  - CS1231 (Discrete Mathematics, 离散数学) [Blogs: 1, 2,]
  - CS3230 (Analysis of Algorithms, 算法设计与分析)
  - new!



### **More on Pancake Flipping**

Have some fun with pancake flipping:

http://www.cut-the-knot.org/SimpleGames/Flipper.shtml

Read about Pancake Flipping:

http://www.amazon.com/Bill-Gates-Speaks-Greatest-Entrepreneur/dp/ 0471401692

Listen to the story on NPR (July-2008):

http://www.npr.org/templates/story/story.php?storyId=92236781

Youtube Video:

https://www.youtube.com/watch?v=oDzauRFiWFU

How are the following people connected to Bill Gates? Jacob Goodman, Harry Lewis, Harry Dweighter, David Cohen, Manuel Blum



#### **More on Genome Rearrangments**

Genome Rearrangement:

http://www.cut-the-knot.org/SimpleGames/Flipper.shtml

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Thank you!

#### If you want to contact me, email, FB: leonghw@comp.nus.edu.sg

or

Google "Leong Hon Wai"



#### The Remaining slides are

#### FOR YOUR FUN READING ONLY



#### **Activity: Fun with Pancake Flipping**

Try your hand on this example. How many flips are needed?







# What are Reversals in the study of Genomics?

Example: (2 genomes G and H, from different species)  $G = \{ 1, -5, 4, -3, 2 \}$   $H = \{ 1, 2, 3, 4, 5 \}$ 

A genome consists of a sequence of genes;
Represent each gene by a integer {1,2, ..., n}.
Each gene has a "*direction*" indicated by the sign (like -3).
Then a genome is a "like" a stack of pancakes of diff sizes.
(each pancake is burnt on one side, indicate "direction")

1()

9

# **Reversals in the Gene Sequence**

2

3

8

1, 2, 3, 4, 5, 6, 7, 8, 9, 10

\* Consider the block of genes (4, 5, 6, 7, 8,)

# **Reversals and Breakpoints**







\* Consider the block of genes (4, 5, 6, 7, 8,)
\* The reversal introduced two *breakpoints* 
\* Order of genes reversed, and direction inverted (4, 5, 6, 7, 8,) → (-8, -7, -6, -5, -4,)

www.bioalgorithms.info

# Reversals: Example













• Gene order comparison:



Evolution is manifested as the divergence in gene order

# Transforming Cabbage into Turnip (with 3 reversal operations)





- What are the similarity blocks and how to find them?
- What is the architecture of the ancestral genome?
- What is the evolutionary scenario for transforming one genome into the other?

# History of Chromosome X



Rat Consortium, Nature, 2004