

# Combinatorics on words in DNA computing

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# Outline

- 1 DNA computing
  - Introduction
  - DNA molecule
  - The first DNA computation
  - Operations in DNA computing
  - DNA computing revisited
  - What to avoid in a test tube
- 2 Combinatorics on Words
  - Setting in CoW
  - Results
  - More results
  - If there is some time left...

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# Introduction

barriers for “traditional” computers

- ① HUP
- ② von Neumann bottleneck

DNA/biomolecular computers: 1994 Leonard Adleman

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# DNA - basics

deoxyribonucleic acid

1950s: DNA carries genetic information (double helix model)

structure: polymer chains - strands consisting of bases attached to sugar phosphate backbone

4 bases: A, G, T, C

Watson-Crick complementarity: A likes T, C likes G

bases are at a distance of 3,4 ångströms, resulting in 18 Mbits per inch

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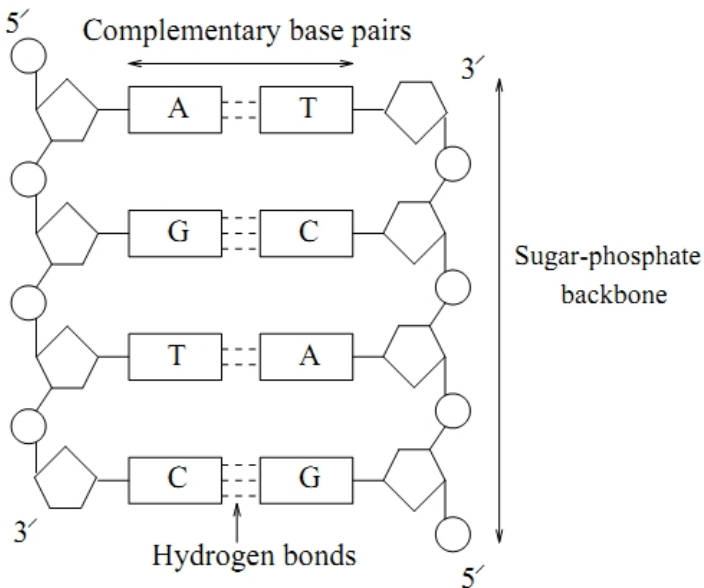
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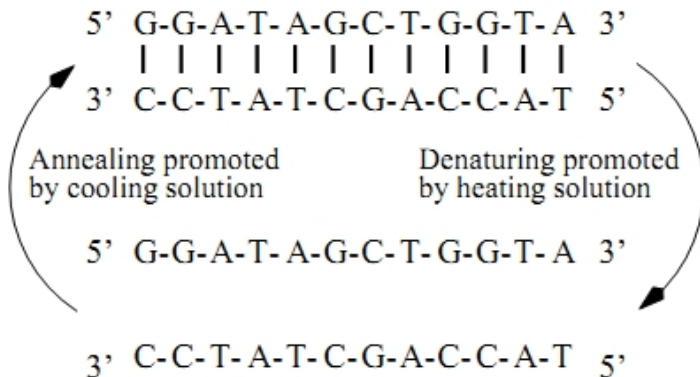
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# Structure of DNA

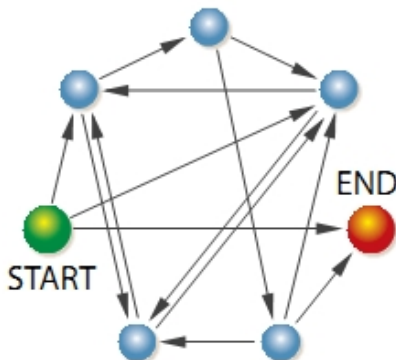


# Watson-Crick complementarity and WK-palindromes



# Hamiltonian Path Problem

Leonard Adleman in 1994: Hamiltonian Path Problem  
(NP-complete) on a graph with 7 vertices

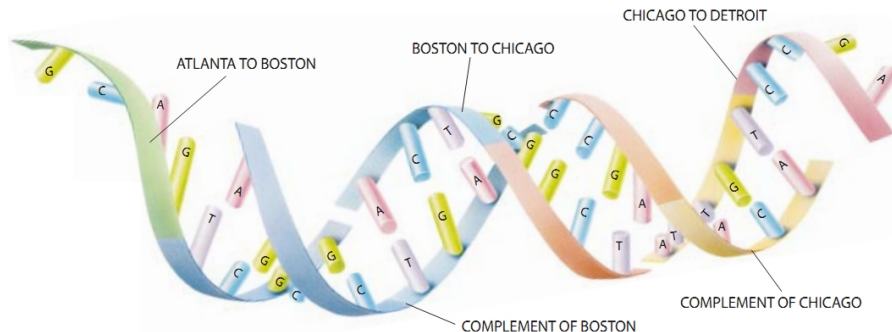


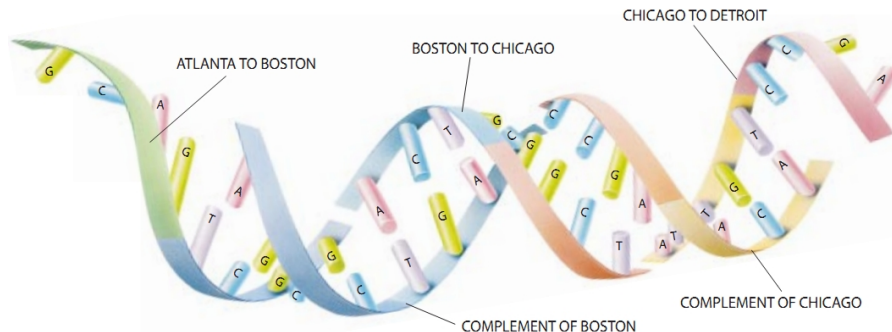
# Adleman's DNA computation I

| CITY    | DNA NAME | COMPLEMENT |
|---------|----------|------------|
| ATLANTA | ACTTGCAG | TGAACGTC   |
| BOSTON  | TCGGACTG | AGCCTGAC   |
| CHICAGO | GGCTATGT | CCGATACA   |
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| FLIGHT            | DNA FLIGHT NUMBER |            |
| ATLANTA - BOSTON  | GCAGTCGG          |            |
| ATLANTA - DETROIT | GCAGCCGA          |            |
| BOSTON - CHICAGO  | ACTGGGCT          |            |
| BOSTON - DETROIT  | ACTGCCGA          |            |
| BOSTON - ATLANTA  | ACTGACTT          |            |
| CHICAGO - DETROIT | ATGTCCGA          |            |





200-node instance would require  $10^{24}$  Earth masses of DNA

# The likely frame

The scale of this ligation reaction far exceeded what was necessary for the graph under consideration. For each edge in the graph, approximately  $3 \times 10^{13}$  copies of the associated oligonucleotides were added to the ligation reaction. Hence it is likely that vast numbers of DNA molecules encoding the Hamiltonian path were created. In theory the creation of a single such molecule would be sufficient. Hence, for this graph, sub-attomol quantities of oligonucleotides would probably have been sufficient. Alternatively, a much larger graph could have been processed with the pmol quantities employed here.

# Manipulation with DNA

- synthesis
- denaturing, annealing and ligation
- separation - affinity purification
- detect
- gel electrophoresis
- PCR - polymerase chain reaction
- cutting (using restriction enzymes)

# Very general model of a DNA calculation

1. select the language - code
2. do the computation
3. decode

# Advantages and drawbacks

## Advantages:

- Size: the information density could go up to 1 bit per cube nm
- High parallelism:  $10^9$  calculations per ml of DNA per second
- Energy efficiency:  $10^{19}$  operations per Joule

## Drawbacks:

- required mass of DNA
- reagents
- need of manipulation by a human
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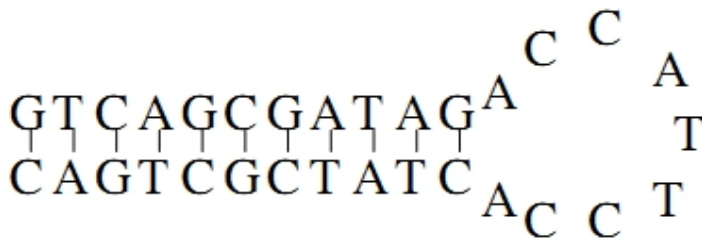
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# What problems can be solved by a DNA computer

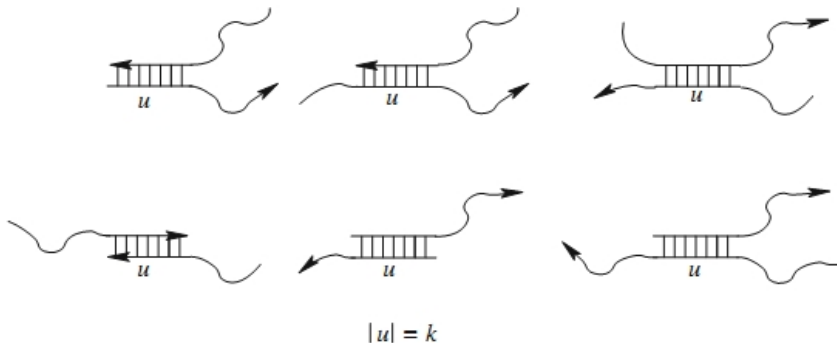
Problems solved that can be found in literature:

- TSP - travelling salesman problem
- addition
- SAT - satisfiability problem
- DES cracking
- maximal clique problem
- ...

# Intramolecular hybridization - hairpins



# Intermolecular hybridization



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# Representing DNA strands

DNA strands are considered in their  $5' \rightarrow 3'$  orientation as finite words over  $\Delta = \{A, G, C, T\}$

involutive antimorphism WK:  $A \leftrightarrow T, C \leftrightarrow G$

an encoding (i.e. initial set of words) for a computation is a language  $L \subset \Delta^*$

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# Questions

How to choose the initial coding set  $L$ ?

- 1 simulation
- 2 algorithm
- 3 theory

Bio-operations vs.  $L$ : after each bio-operation the language changes - do the properties hold?

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# Constraints on $L$

to avoid possible hybridizations, we impose several constraints on  $L$

- $L$  is  $\Theta$ - $k$ - $m$ -subword code if  
 $\forall u \in \mathcal{A}^k, 1 \leq i \leq m, \mathcal{A}^* u \mathcal{A}^i \Theta(u) \mathcal{A}^* \cap L = \emptyset$
- $L$  is  $\Theta$ - $k$ -code if  $\forall u, v \in \mathcal{A}^k \cap L, u \neq \Theta(v)$
- $L$  is bond-free if  $\forall u, v \in \mathcal{A}^k \cap L, H(u, \Theta(v)) > d$ , where  $H$  is the Hamming distance
- frequency of  $C$  and  $G$  in each word should be  $\frac{1}{2}$
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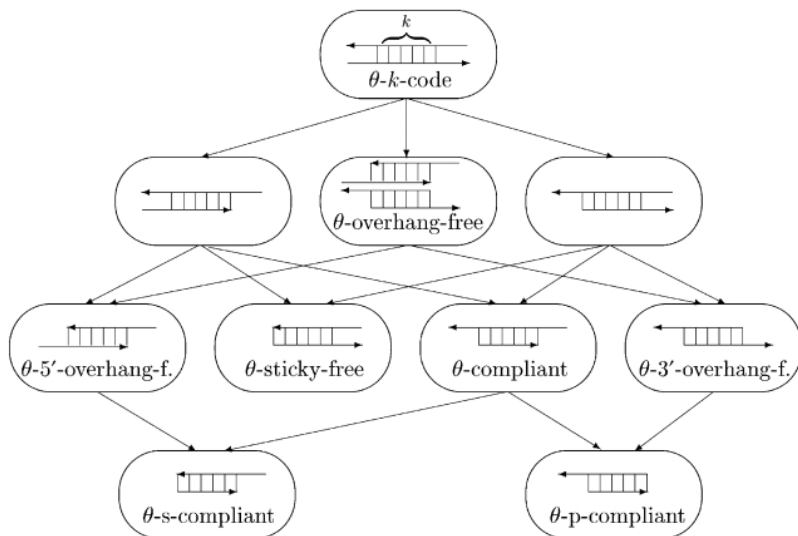
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# Conjugates and $\Theta$ -conjugates

## Proposition (Lyndon and Schützenberger, 1962)

*Let  $u, v, w \in \mathcal{A}^*$  such that  $uv = vw$ . Then there exist  $p, q \in \mathcal{A}^+$  such that  $u = pq$ ,  $w = qp$  and  $v = p(qp)^i$  for  $i \geq 0$ .*

## Proposition (Kari and Mahalingam, 2007)

*Let  $u, v, w \in \mathcal{A}^+$  such that  $uv = \Theta(v)w$ , where  $\Theta$  is an involutive morphism or antimorphism over  $\mathcal{A}$ .*

- ① *If  $\Theta$  is an antimorphism, then there exist  $x, y \in \mathcal{A}^*$  such that either*
  - *$u = xy$ ,  $w = y\Theta(x)$  and  $v = \Theta(x)$ ; or*
  - *$u = x = \Theta(w)$ ,  $y = \Theta(y)$  and  $v = y$ .*
- ② *If  $\Theta$  is a morphism, then there exist  $x, y \in \mathcal{A}^*$  such that  $u = xy$  and one of the following hold:*
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# Set of $\Theta$ -palindromes

## Proposition

*The set of all  $\Theta$ -palindromes ( $\Theta$  being an involutive antimorphism) is not regular.*

## Lemma (Pumping lemma for regular languages)

*Let  $L$  be a regular language. Then there exists an integer  $p \geq 1$  depending only on  $L$  such that every  $w \in L$  of length at least  $p$  can be written as  $w = xyz$  satisfying the following conditions:*

- ①  $|y| \geq 1$ ,
- ②  $|xy| \geq p$ ,
- ③ for all  $i \geq 0$ ,  $xy^iz \in L$ .

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*Let  $u \in \mathcal{A}$  be a non-empty word such that  $u \neq \Theta(u)$ . Then the following statements are equivalent*

- ①  $\sqrt{u}$  is the product of two non-empty  $\Theta$ -palindromes.*
- ② There exists a non-empty  $\Theta$ -palindrome  $v$  such that  $\Theta$ -commutes with  $u$ .*
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## Second untitled frame

### Proposition

*Let  $\Theta$  be either a morphic or an antimorphic involution and let  $\Sigma$  be such that for all  $a \in \Sigma$ ,  $a \neq \Theta(a)$ .*

*(...)*

# References

Lila Kari

Thank you.