

How Does Nature Compute?

Lila Kari

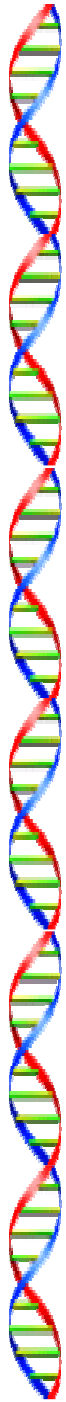
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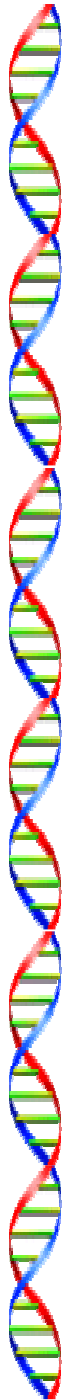
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Computers: What can they accomplish?

- Fly spaceships to Mars
- Control aircraft
- Robot aided manufacturing
- Computer games
- Expedite journal submissions
- Email

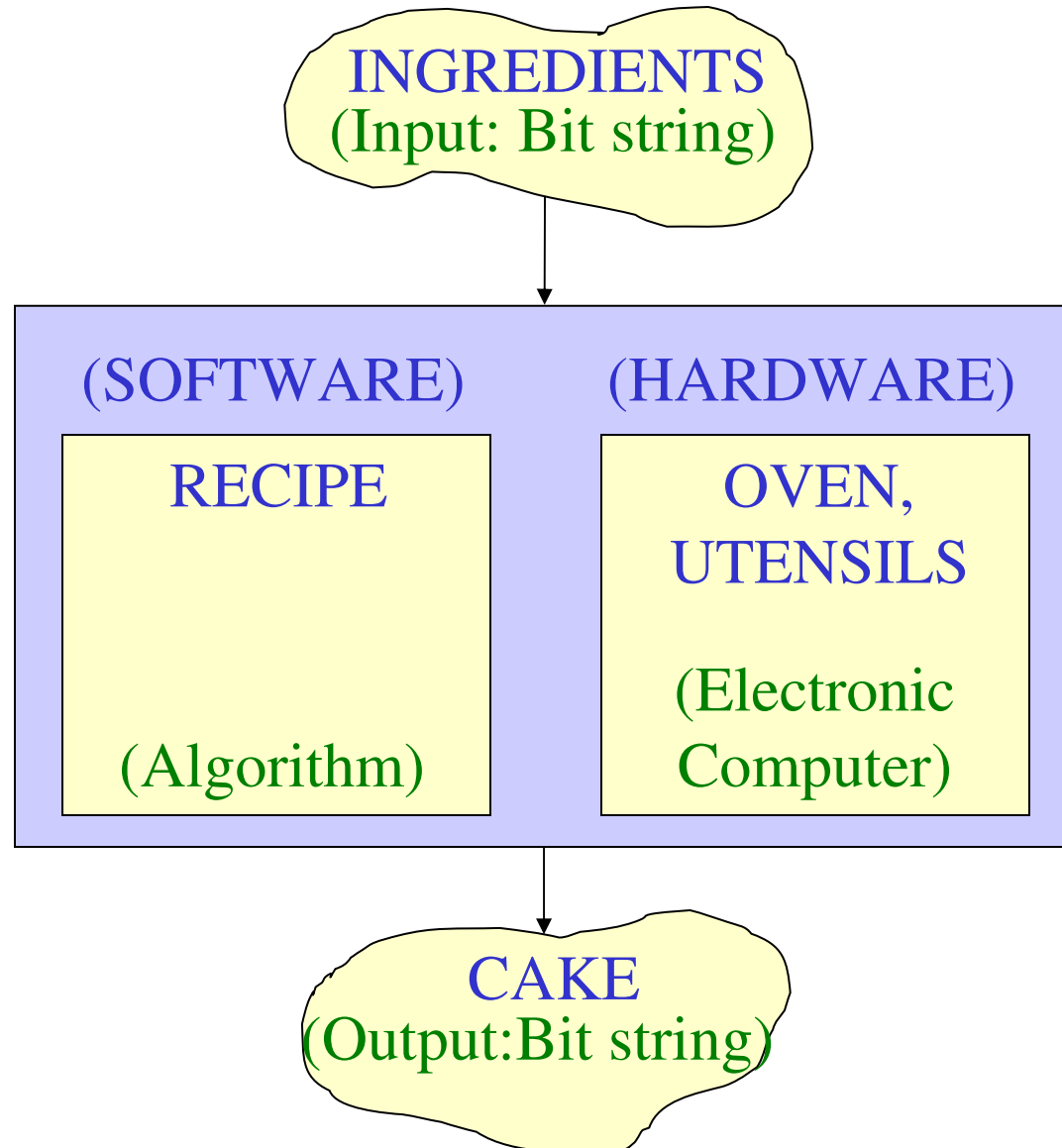
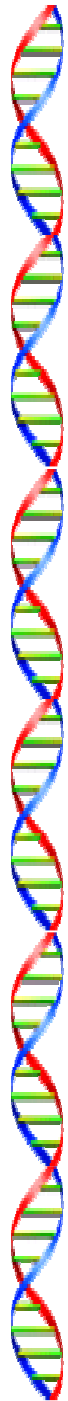


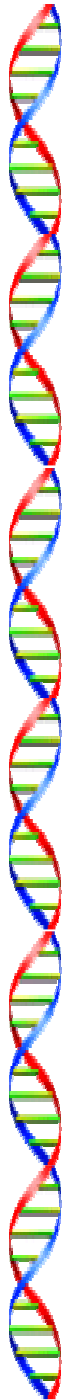
Computers:

What do they actually do?

- Computers = a collection of **switches (bits)** that are on (**1**) or off (**0**).
- Can execute only **simple operations**
 - § Flipping a bit's value
 - § Zeroing a bit
 - § Testing a bit

How do they do it?





Formal Models of Computing: Turing Machines

- Data

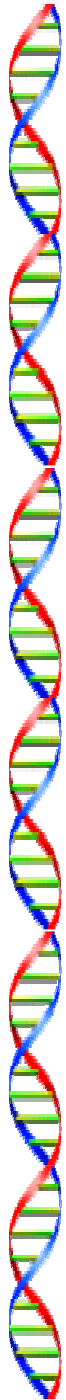
- § String of symbols written on a **tape**

- Operations

- § **Read** a square

- § **Overwrite** the symbol with another

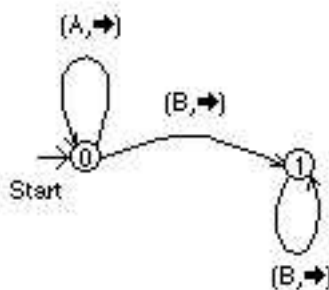
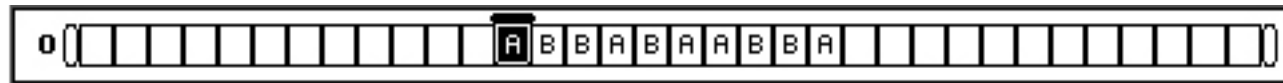
- § **Move** left or right



Turing Machine

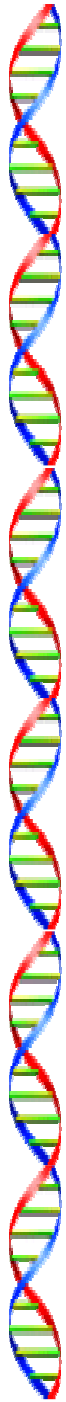
Computation = Finite list of instructions

“If you are in state **S** and read input symbol **X** then write **Y** and move Left/Right”



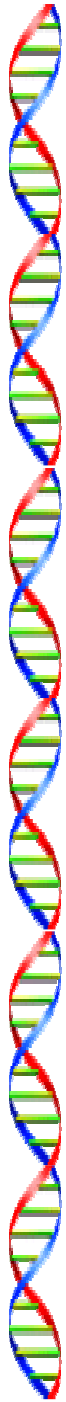
TO RUN: enter a sequence of A's and B's, position read head on leftmost symbol, and start.

	State	Significance
Start	0	Has not yet seen a B
	1	Has seen at least one B



Turing Machine

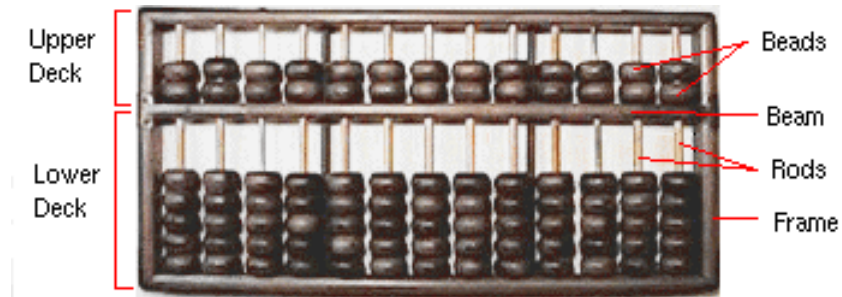
- Turing machines are capable of **universal computation** (everything that can be computed can be computed by a TM)
- The abstract notion of **computation** (Turing machine, algorithm, program) is **hardware independent**

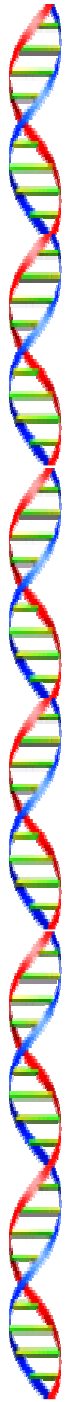


History of Hardware

- Abacus
- Pascal
- Jacquard
- Babbage
- Hollerith
- ENIAC
- Chip

Abacus

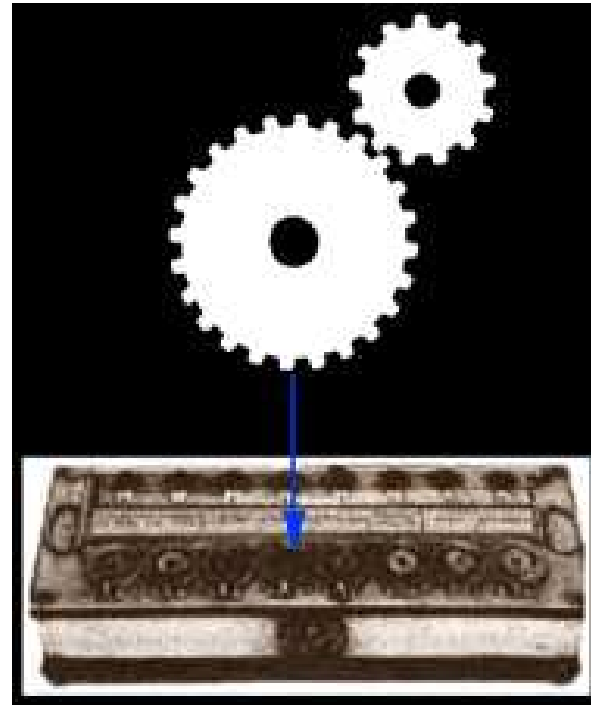


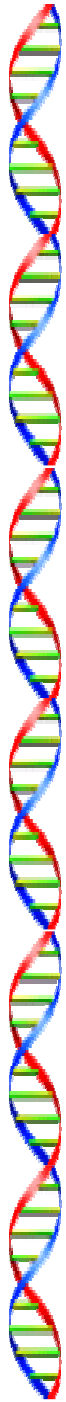


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Mechanical adding machine (1642)



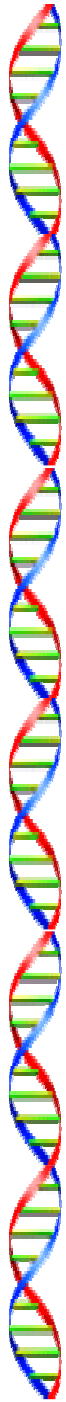


History of Hardware

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Jacquard's punch card loom (1801)

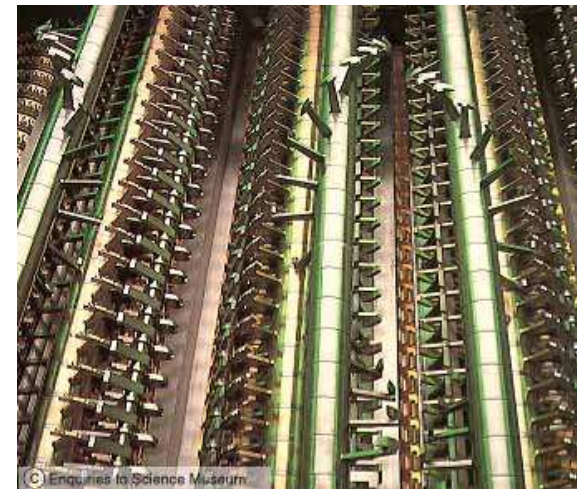
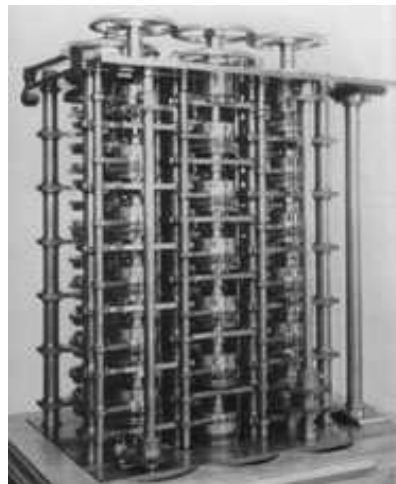


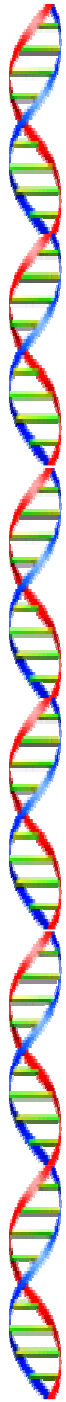


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- Babbage
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- Chip

Babbage's difference engine (1833)

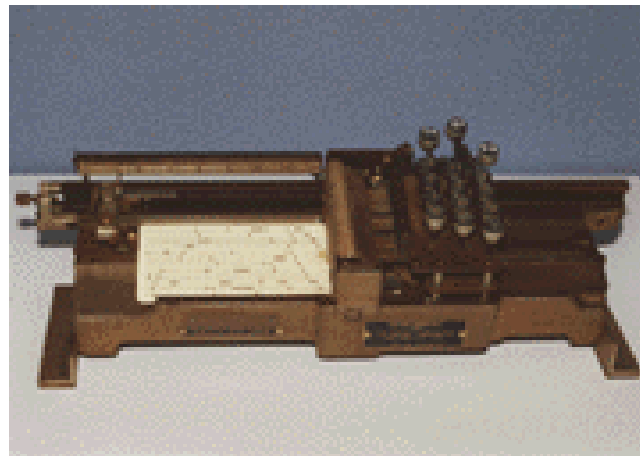


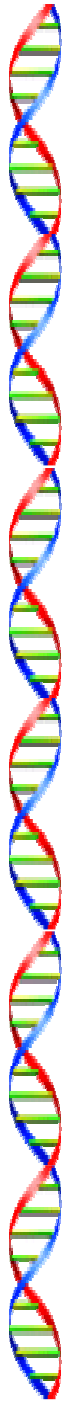


History of Hardware

- Abacus
- Pascal
- Jacquard
- Babbage
- **Hollerith**
- ENIAC
- Chip

Hollerith punch card system (1890)



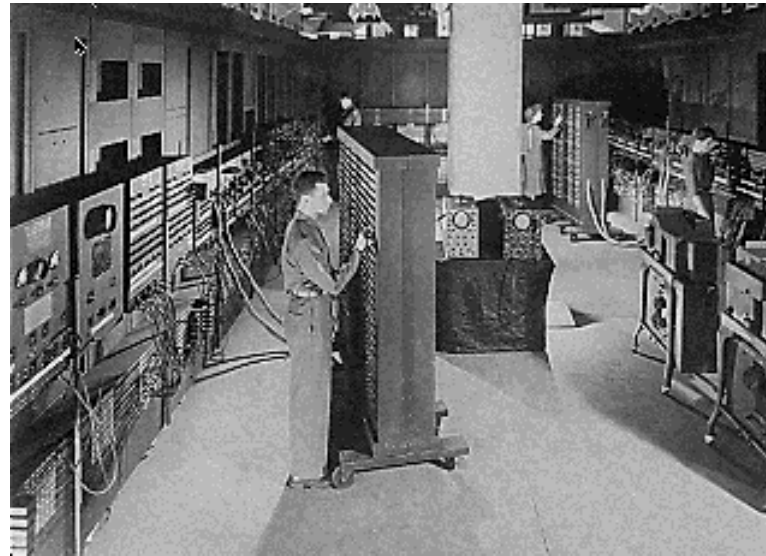


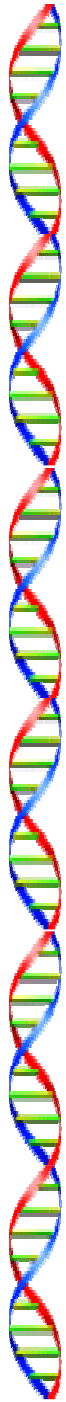
History of Hardware

- Abacus
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ENIAC (1939-45) -167 sq.m.

- 18,000 vacuum tubes
- not programmable



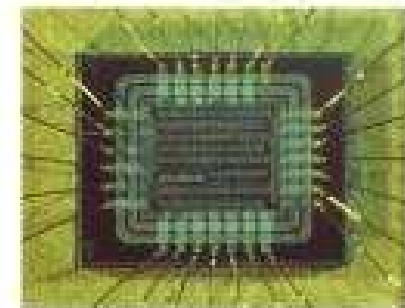
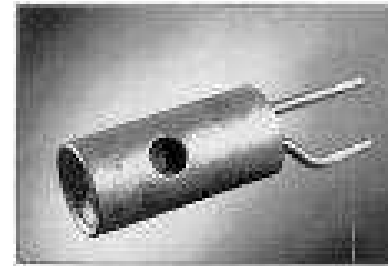


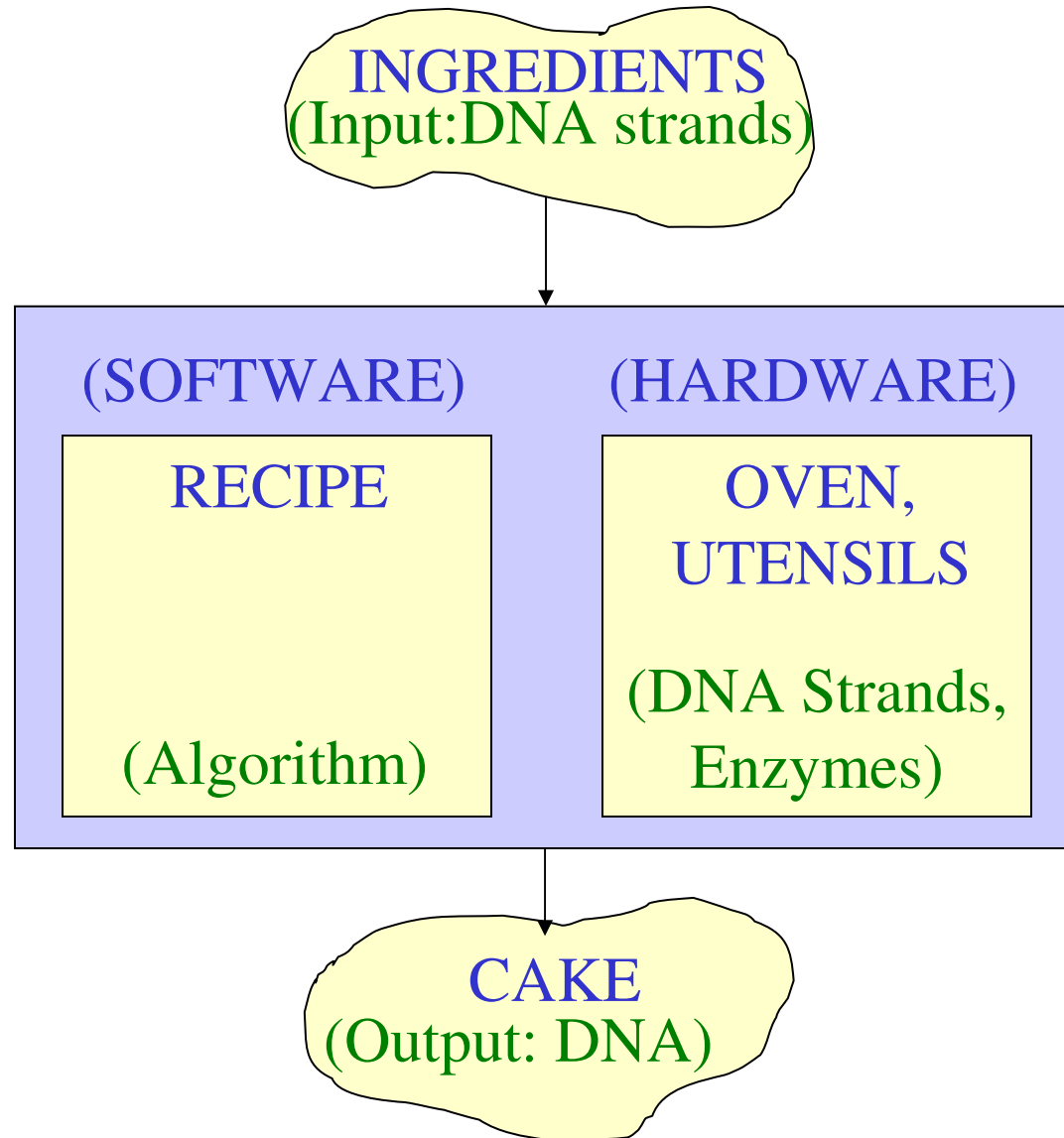
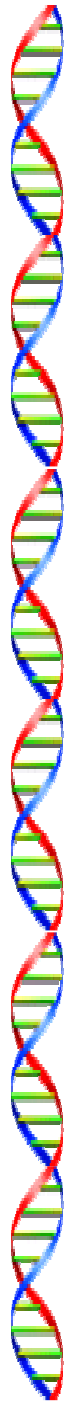
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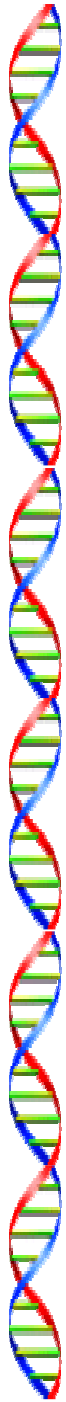
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Modern computer chip

- Transistors
- Integrated circuit

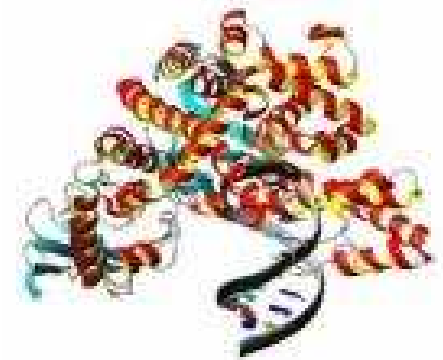
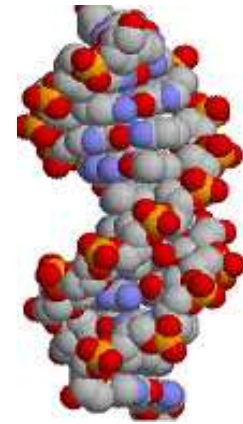


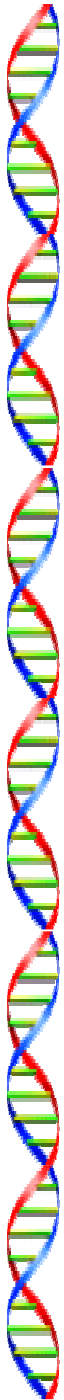




DNA Computer

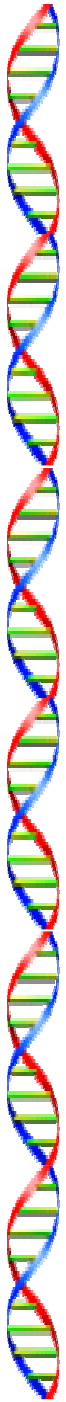
- **Input / Output (DNA)**
 - § Data encoded using the DNA alphabet {A, C, G, T} and synthesized as DNA strands
- **Bio-operations**
 - § Cut
 - § Paste
 - § Copy
 - § Anneal
 - § Recombination





Biomolecular (DNA) Computing

- Hamiltonian Path Problem [Adleman, Science, 1994]
- DNA-based addition [Guarnieri et al, Science, 1996]
- Maximal Clique Problem [Ouyang et al, Science, 1997]
- DNA computing by self-assembly [Winfree et al, Nature 1998]
- Computations by circular insertions, deletions [Daley et al, 1999]
- DNA computing on surfaces [Liu et al, Nature, 2000]
- Molecular computation by DNA hairpin formation
[Sakamoto et al, Science, 2000]
- Programmable and autonomous computing machines made of biomolecules [Benenson et al, Nature, 2001]
- 20-variable Satisfiability [Braich et al., Science 2002]



How Does Nature Compute?

- Technical difficulties encountered in experimental DNA computations (error-detection, error-correction) are routinely solved by **biological systems in nature**
- Idea: study and utilize the **computational abilities of unicellular organisms**

Ciliates: Unicellular Protozoa

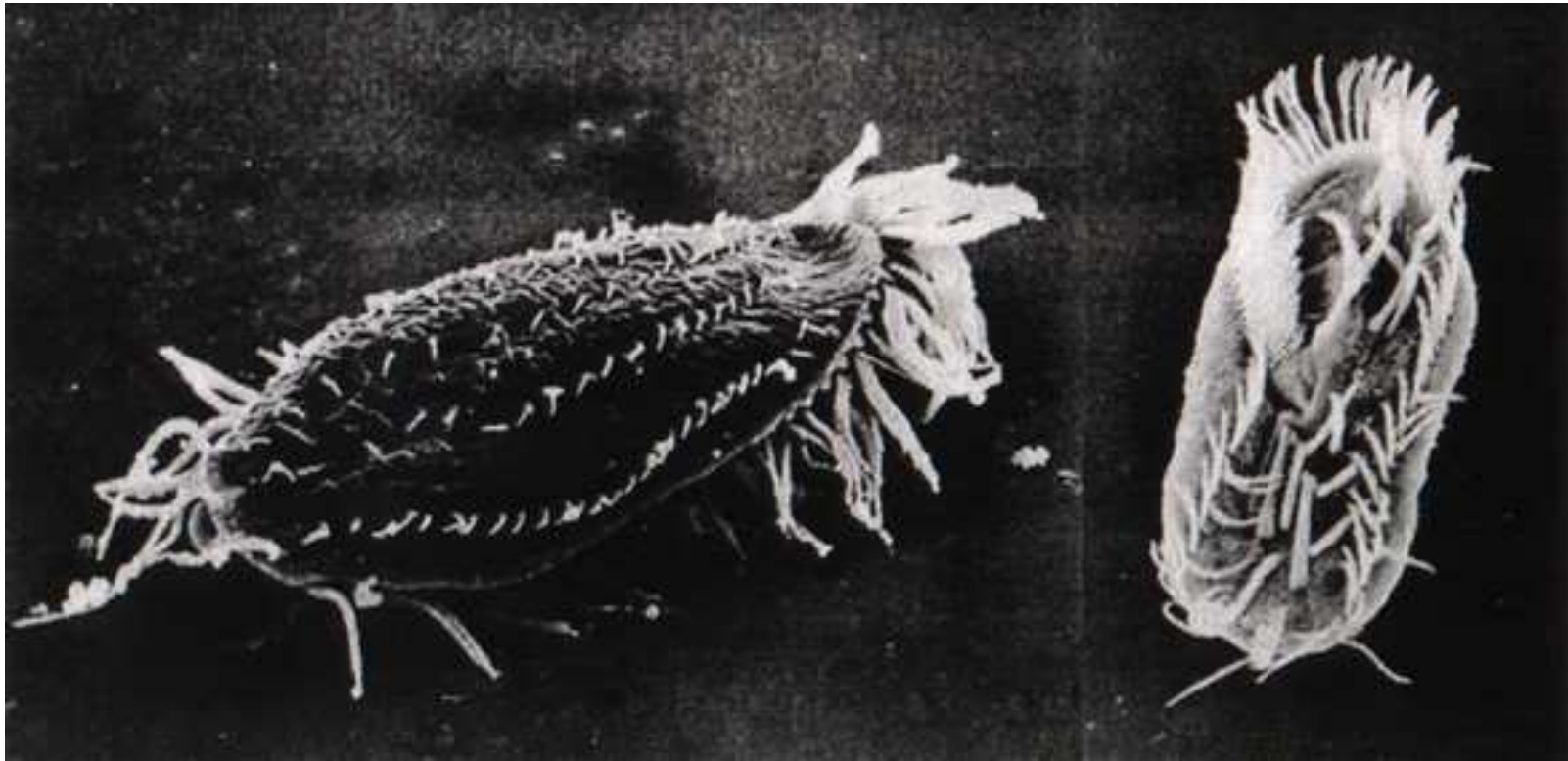
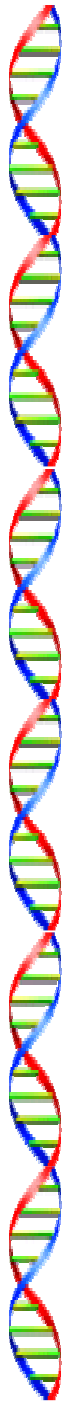


Photo courtesy of L.F. Landweber

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Ciliates: Genetic Info Exchange

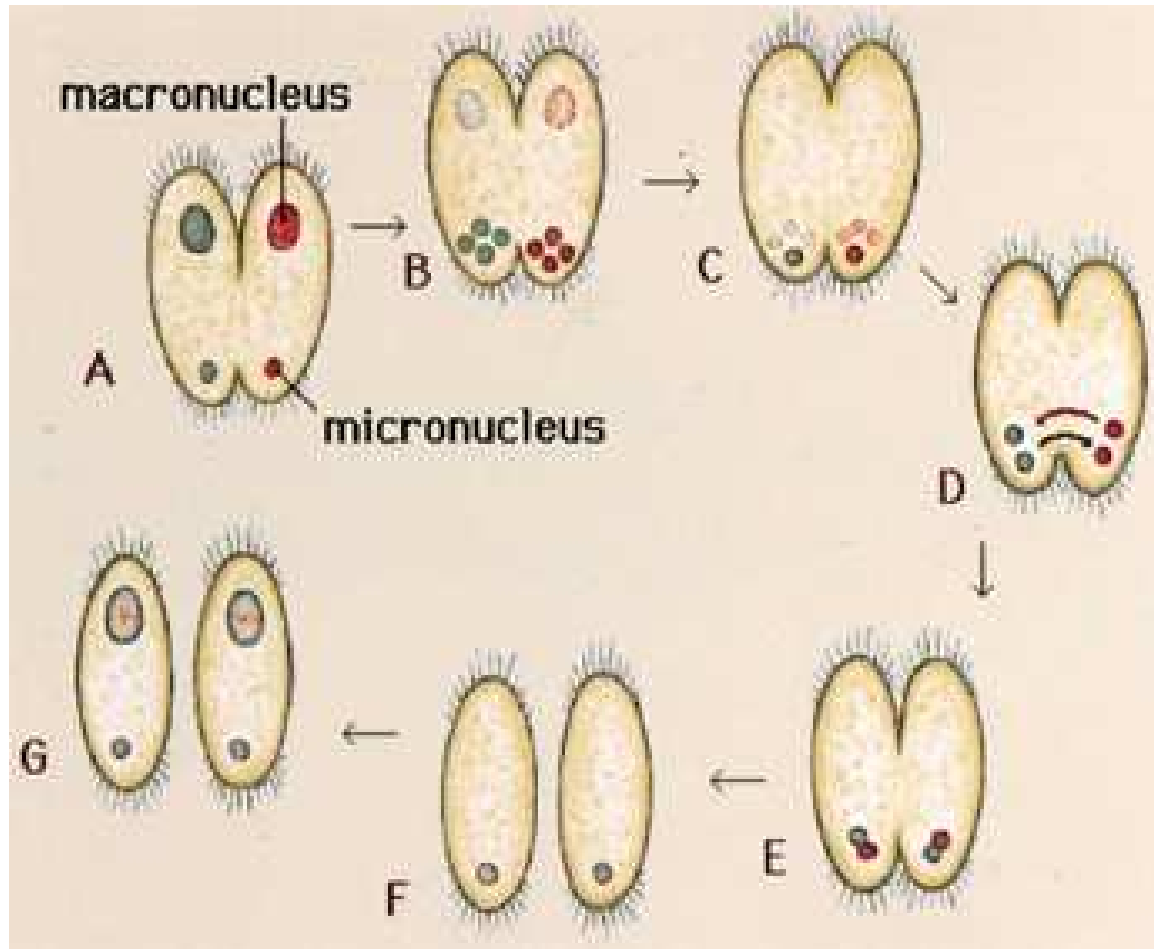
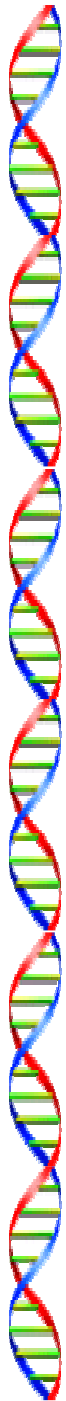


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Ciliates: Gene Rearrangement

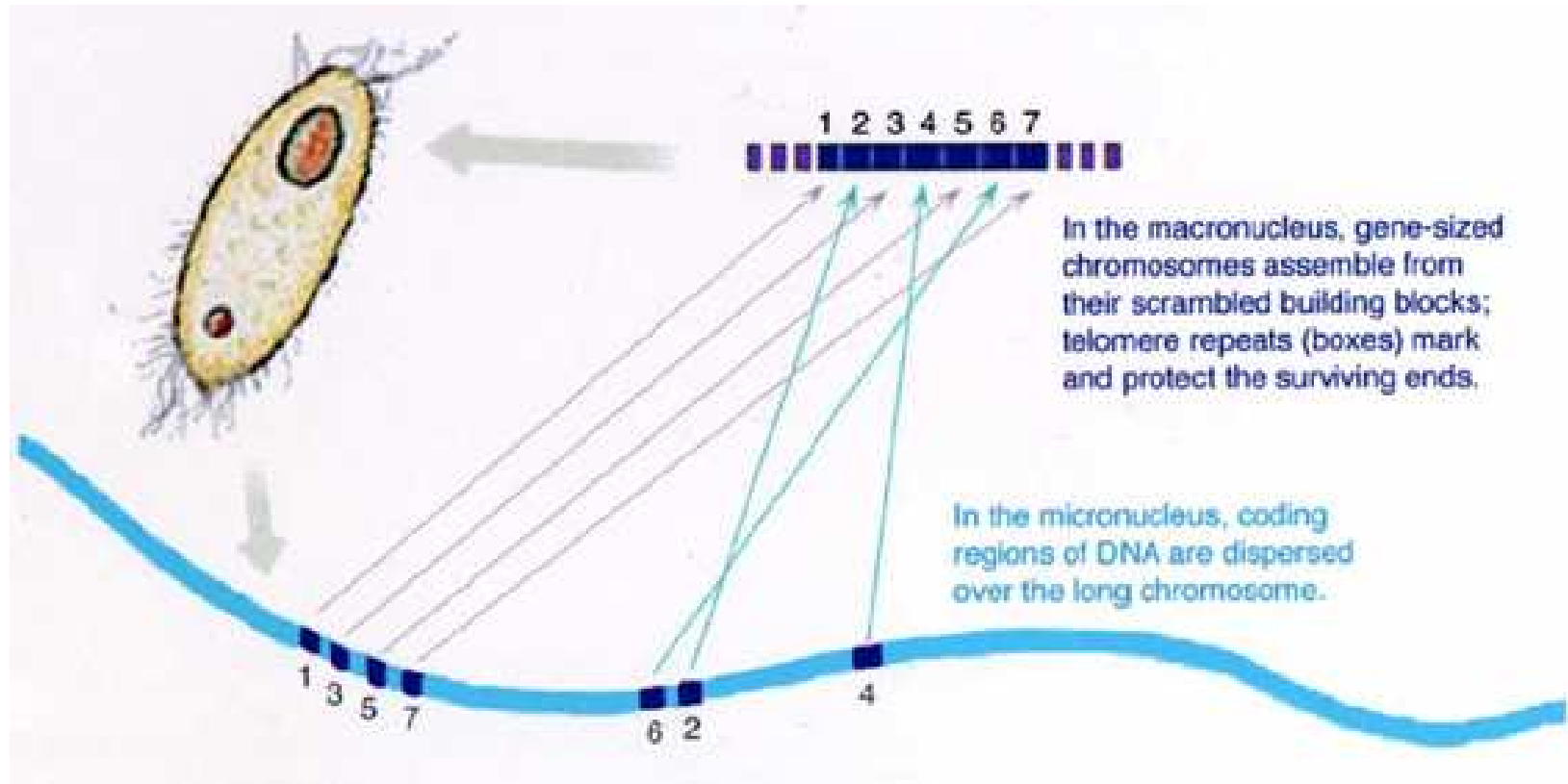
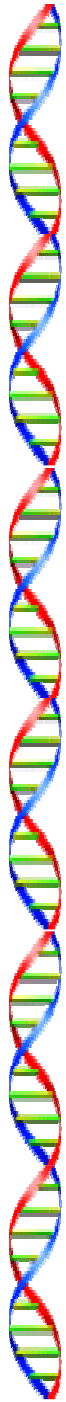
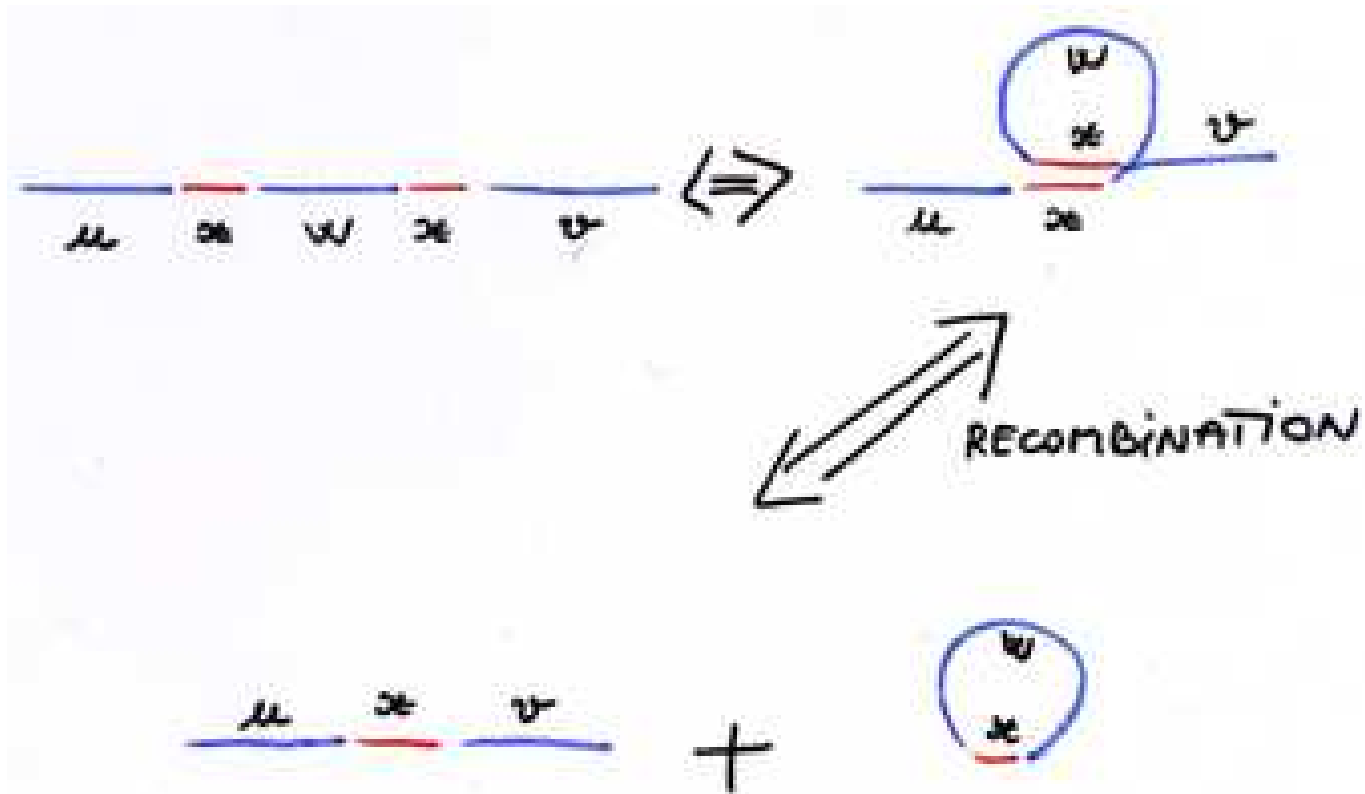


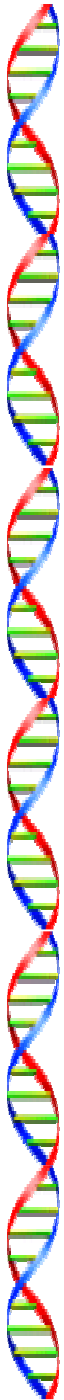
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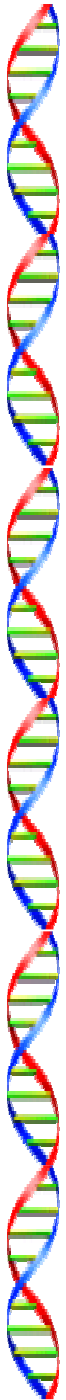
Ciliates: Bio-operations





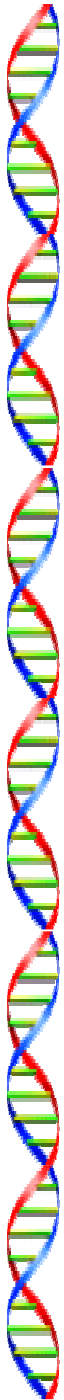
Ciliates: Results

- **Guided Recombination System** = A formal computational model based on **contextual circular insertions and deletions**
- Such systems have the **computational power** of **Turing Machines** (Landweber, Kari, '99)
- The model is consistent with the limited knowledge of this biological process



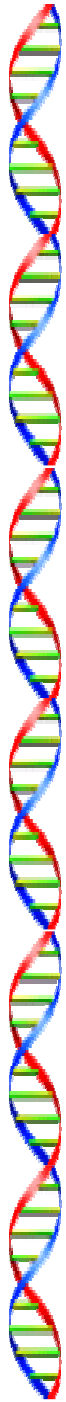
Essential Feature of Biocomputing: Self-Assembly

- Use knowledge of how **simple components** (DNA molecules, enzymes) interact
- Design a setup such that the computation happens essentially **by itself**
- Useful in **nano-technology** where components are too small for existing tools



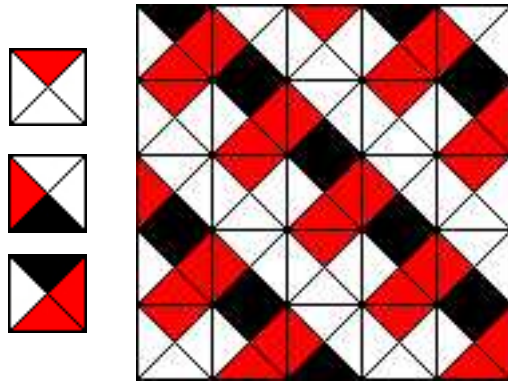
Model of Self-Assembly

- Tile
 - § A 1 x 1 square
 - § Each side is “painted” with a certain kind of glue
 - § Tiles cannot be rotated
 - § Two “adjacent” tiles will “stick” only if they have matching glues at the touching edges
- Tile system
 - § T = A finite number of tile types (as above)
 - § Unlimited supply of each tile type available

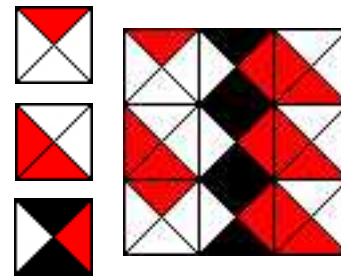


(Classic) Tiling Problem

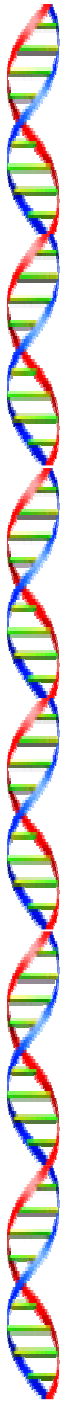
- Can **any** square, of any size, be tiled using only the available tile types, without violating the glue-matching rule?



Yes

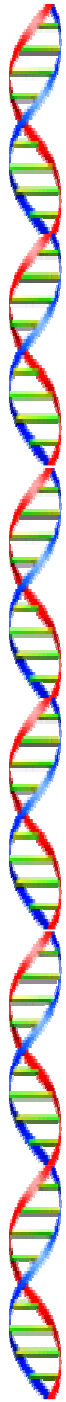


No



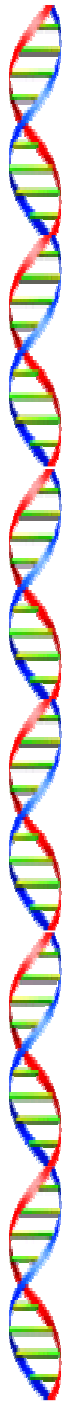
Self-Assembly Problems

- What is the **minimal** number of tile types that can **self-assemble** into a given shape and nothing else?
- What is the **optimal initial concentration** of tile types that ensures fastest **self-assembly**?
- What happens if “bonds” have **different strengths**?

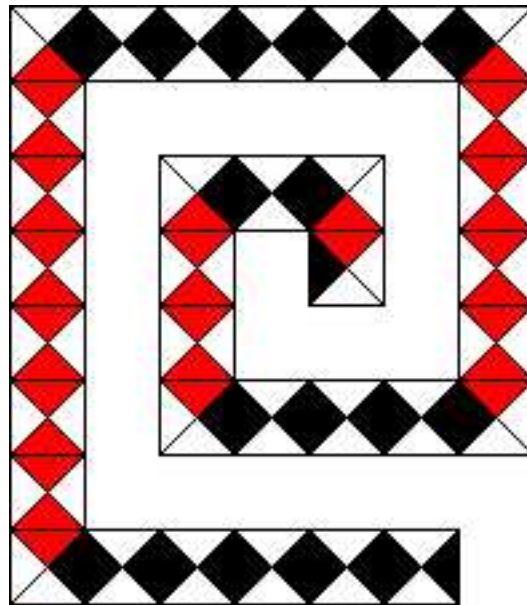
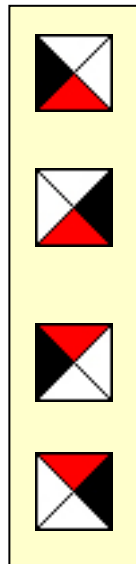


The Ribbon Problem

- Given a tile system, can we determine whether or not it can produce shapes that *grow indefinitely*?
- Can we decide whether or not a set of given tiles can produce *unlimited-size ribbons*?



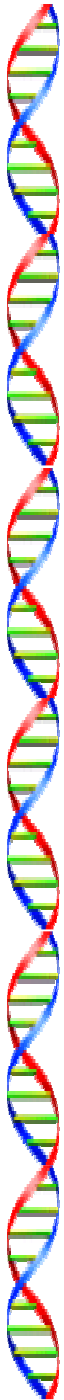
Generating Ribbons





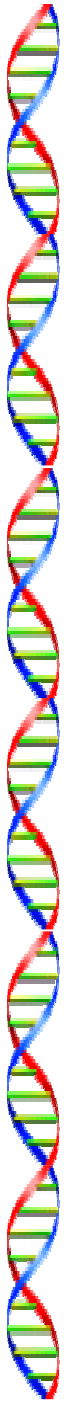
Answer

- There is **no algorithm**, and there never will be, for solving the **Ribbon Problem!**
[Adleman, Kari, Kari, Reishus, 2002]
- You can devise a program that might work quite well, on **some of the inputs**. But there always will be inputs upon which your algorithm will misbehave; it will either run forever, or produce the wrong output.

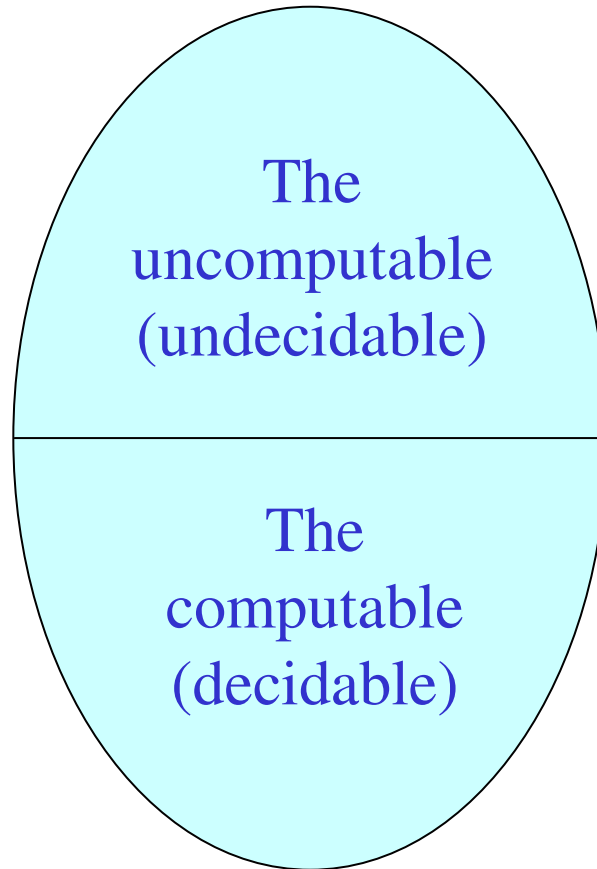


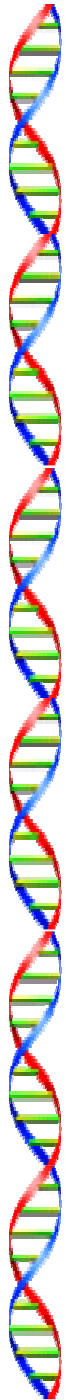
Computable vs. Uncomputable!

- An algorithmic problem that admits no solution is termed **uncomputable**
- If it is a **Yes/No** problem, it is termed undecidable
- The **Ribbon Problem** is **undecidable**



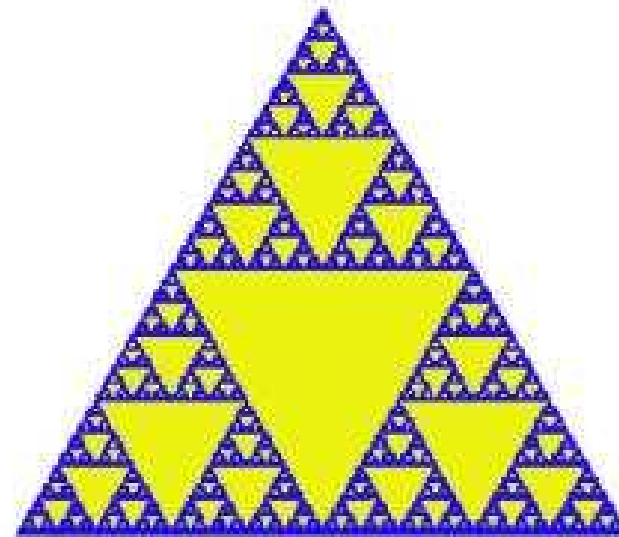
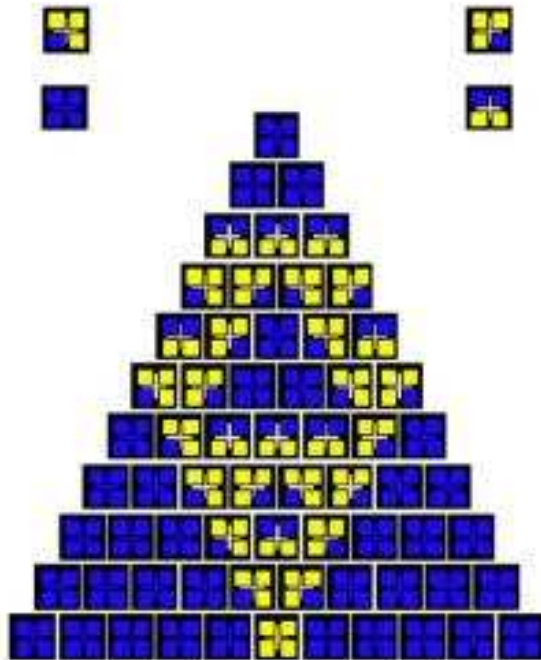
Sometimes we cannot do it!



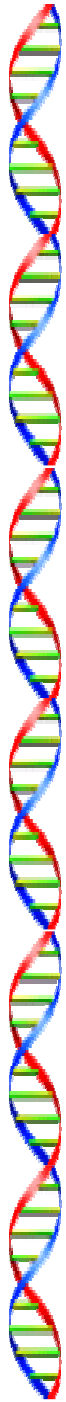


Experimental Self-Assembly

- Self-assembly using capillary forces (Rothemund)
- DNA computing by self-assembly (Winfree, Seeman)

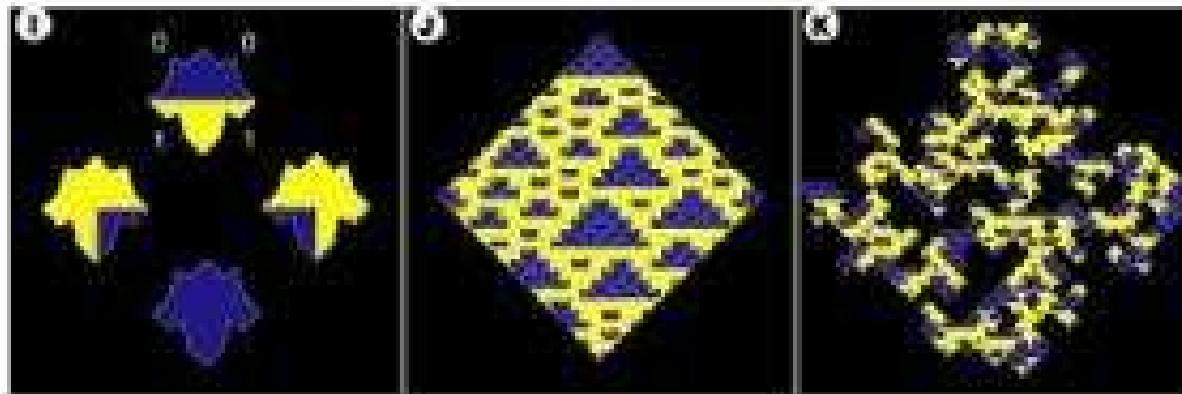


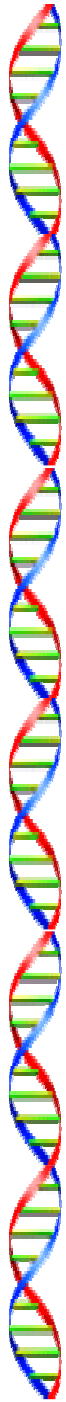
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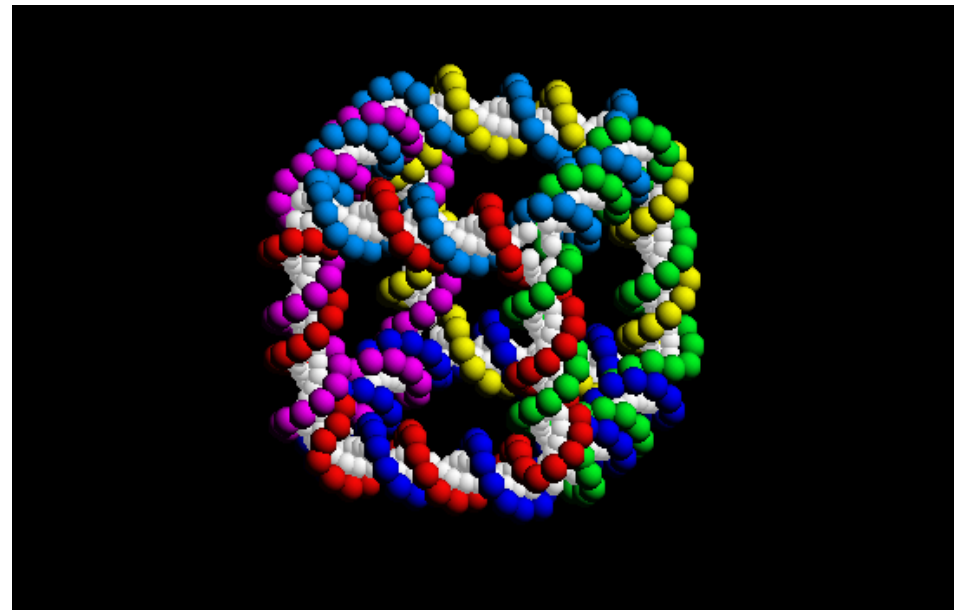
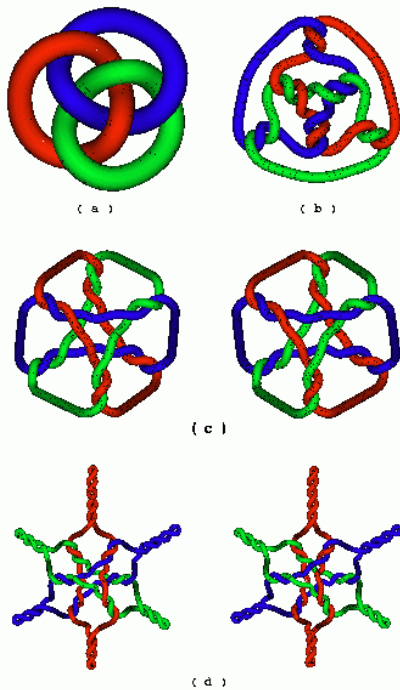
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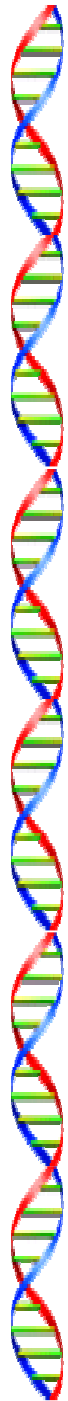


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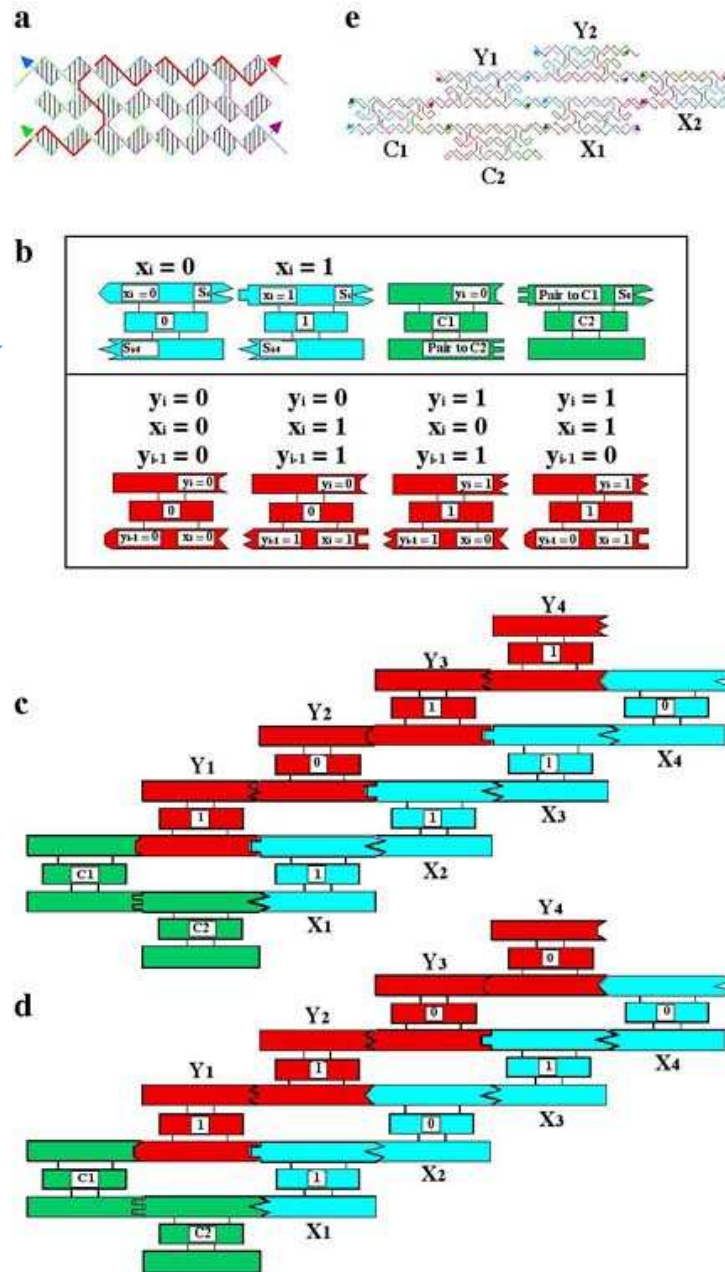
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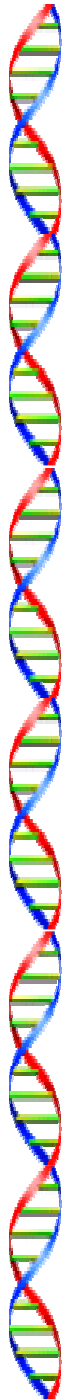


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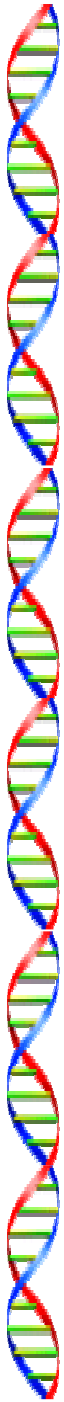
DNA computing by self-assembly
 (Winfree, Seeman, Mao,
 LaBean, Reif)





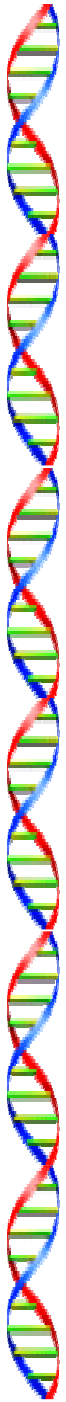
Potential Advantages of Biomolecular (DNA) Computing

- Information density
 - 1 gram of DNA (1 cm³ when dry) = 1 trillion CDs
 - 1 lb DNA – more memory than all computers together.
- Speed
 - Thousand to million times faster than an electronic computer due to massive parallelism
- Energy consumption
 - thousand times more energy efficient



IMPACT OF DNA COMPUTING

- Sheds new light into the nature of computation
- Opens prospects of radically different computers
- Could lend new insights into the information processing abilities of cells
- “Biology and Computer Science – life and computation – are related” (Adleman)



“If we knew what it was that we were doing, it wouldn’t be called research, would it?” (Einstein)