

# DNA Computation

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## Outline

- **Introduction to DNA**
- Adleman's experiment
- Cutting Edge Technologies
- Pros and Cons
- Conclusion

## What is DNA?

- DNA stands for Deoxyribonucleic Acid
- DNA represents the genetic blueprint of living creatures
- DNA contains "instructions" for assembling cells
- Every cell in human body has a complete set of DNA
- DNA is unique for each individual

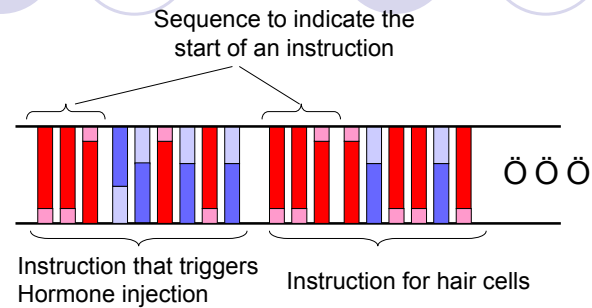
## Double Helix



- "Sides" Sugar-phosphate backbones
- "ladders" complementary base pairs  
Adenine & Thymine  
Guanine & Cytosine
- Two strands are held together by weak hydrogen bonds between the complementary base pairs

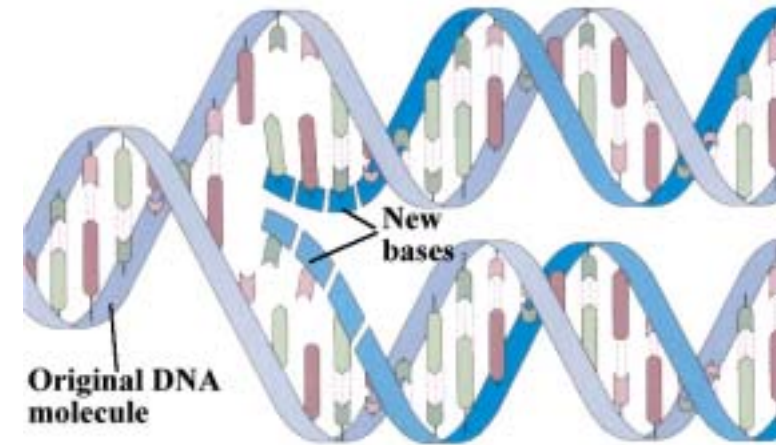
Source: "Human Physiology: From Cells to System"  
4th Ed., L. Sherwood, Brooks/Cole, 2001, C-3

## Instructions in DNA



- Instructions are *coded* in a sequence of the DNA bases
- A segment of DNA is exposed, transcribed and translated to carry out instructions

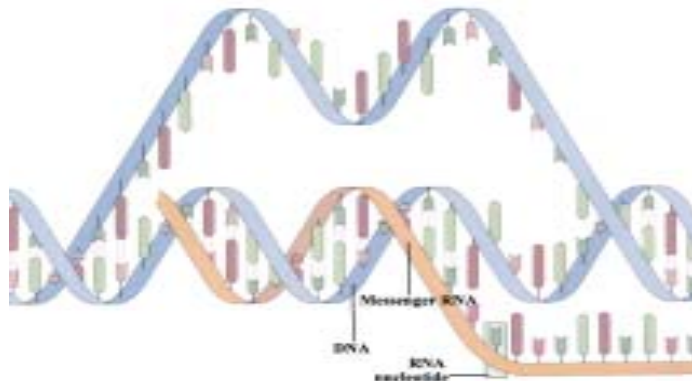
## DNA Duplication



Source: iHuman Physiology: From Cells to System 4th Ed.1, L. Sherwood, Brooks/Cole, 2001, C-5

## Protein Synthesis

- DNA → RNA → Proteins → *actions*



Source: iHuman Physiology: From Cells to System 4th Ed.1, L. Sherwood, Brooks/Cole, 2001, C-6

## Can DNA Compute?

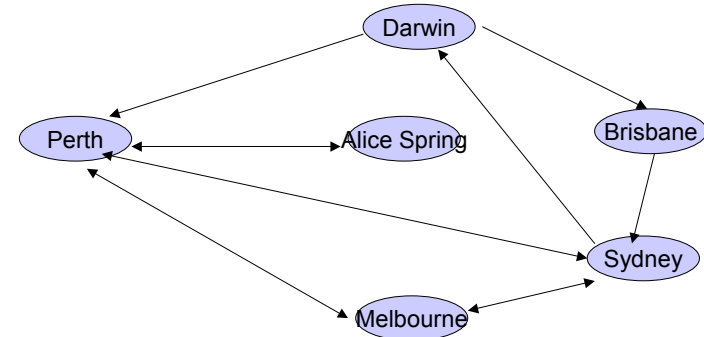
- DNA itself does not carry out any computation. It rather acts as a massive memory.
- BUT, the way complementary bases react with each other can be used to compute things.
- Proposed by Adelman in 1994

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## Adleman's Experiment

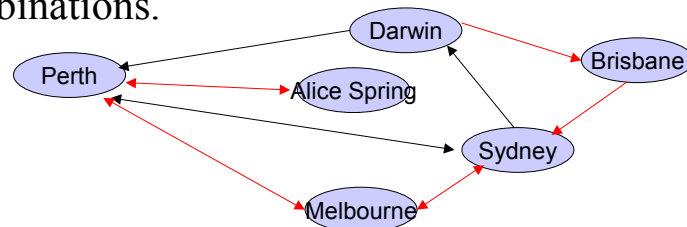
- Hamilton Path Problem  
(also known as the travelling salesperson problem)



Is there any Hamiltonian path from Darwin to Alice Spring?

## Adleman's Experiment (Cont'd)

- Solution by inspection is:  
Darwin → Brisbane → Sydney → Melbourne → Perth → Alice Spring
- BUT, there is no deterministic solution to this problem, i.e. we must check all possible combinations.



## Adleman's Experiment (Cont'd)

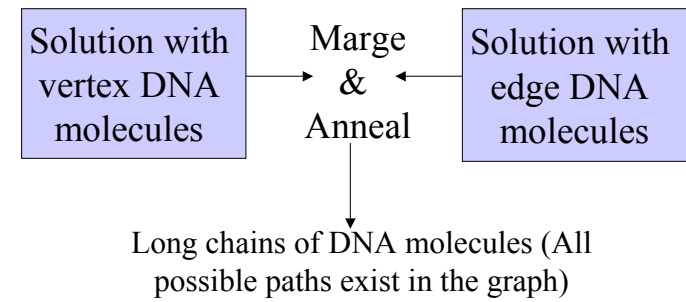
1. Encode each city with complementary base - *vertex molecules*  
Sydney - TTAAGG  
Perth - AAAGGG  
Melbourne - GATACT  
Brisbane - CGGTGC  
Alice Spring - CGTCCA  
Darwin - CCGATG

## Adleman's Experiment (Cont'd)

- Encode all possible paths using the complementary base  $\bar{n}$  *edge molecules*  
 Sydney  $\rightarrow$  Melbourne  $\bar{n}$  AGGGAT  
 Melbourne  $\rightarrow$  Sydney  $\bar{n}$  ACTTTA  
 Melbourne  $\rightarrow$  Perth  $\bar{n}$  ACTGGG  
 etc.

## Adleman's Experiment (Cont'd)

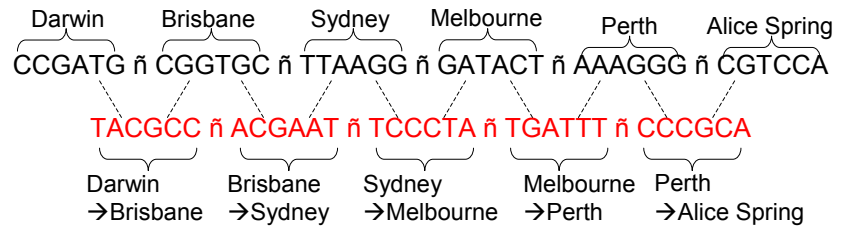
- Merge vertex molecules and edge molecules. All complementary base will adhere to each other to form a long chains of DNA molecules



## Adleman's Experiment (Cont'd)



- The solution is a double helix molecule:



## Operations

- Melting**  
 breaking the weak hydrogen bonds in a double helix to form two DNA strands which are complementary to each other
- Annealing**  
 reconnecting the hydrogen bonds between complementary DNA strands

## Operations (Cont'd)

- Merging  
mixing two test tubes with many DNA molecules
- Amplification  
DNA replication to make many copies of the original DNA molecules
- Selection  
elimination of errors (e.g. mutations) and selection of correct DNA molecules

## Outline

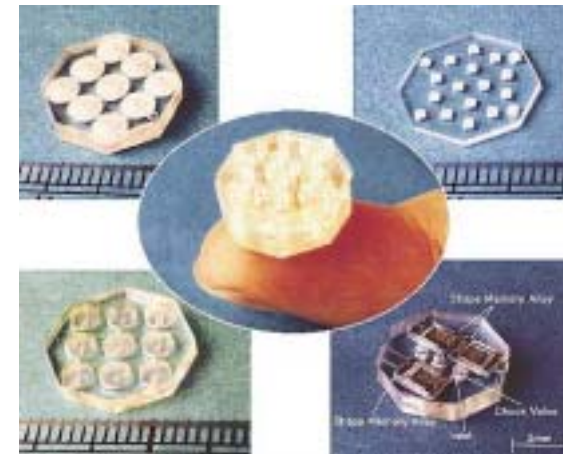
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## DNA Chip



Source: Stanford Medicine Magazine, Vol 19, 3 Nov 2002  
<http://mednews.stanford.edu/stanmed/2002fall/translational-dna.html>

## Chemical IC



Source: Tokyo Techno Forum 21, 21 June 2001  
<http://www.techno-forum21.jp/study/st010627.htm>

## The Smallest Computer

- The smallest programmable DNA computer was developed at Weizmann Institute in Israel by Prof. Ehud Shapiro last year
- It uses enzymes as a program that processes on on the input data (DNA molecules).
- [http://www.weizmann.ac.il/mathusers/lbn/new\\_pages/Research\\_Biological.html](http://www.weizmann.ac.il/mathusers/lbn/new_pages/Research_Biological.html)

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## Pros and Cons

- + Massively parallel processor  
DNA computers are very good to solve Non-deterministic Polynomial problems such as DNA analysis and code cracking.
- + Small in size and power consumption

## Pros and Cons (Cont'd)

- Requires constant supply of proteins and enzymes which are expensive
- Errors occur frequently  
a complex selection mechanism is required and errors increase the amount of DNA solutions needed to compute
- Application specific
- Manual intervention by human is required

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## Conclusion

- Many issues to be overcome to produce a useful DNA computer.
- It will not replace the current computers because it is application specific, but has a potential to replace the high-end research oriented computers in future.
- Nanotechnology?

## References

- **Molecular Computation of Solutions to Combinatorial Problems**, *L.M. Adleman*, Science Vol.266 pp1021-1024, 11 Nov 1994
- **Computing With Cells and Atoms** – an introduction to quantum, DNA and membrane computing, *C.S. Calude and G. Paun*, Taylor & Francis, 2001
- **The Cutting Edge Biomedical Technologies in the 21<sup>st</sup> Century**, Newton, 1999
- **Human Physiology: From Cells to Systems** 4<sup>th</sup> Ed., *L. Sherwood*, Brooks/Cole, 2001