

Tuesday: 14.15 -16.00, room P016

Jaap Kaandorp

[illegible]

- Bioinformatics or computational biology is the use of techniques from applied mathematics, informatics, statistics, and computer science to solve biological problems. Research in computational biology often overlaps with systems biology. Major research efforts in the field include sequence alignment, gene finding, genome assembly, protein structure alignment, protein structure prediction, prediction of gene expression and protein-protein interactions, and the modeling of evolution. The terms bioinformatics and computational biology are often used interchangeably, although the latter typically focuses on algorithm development and specific computational methods

The diagram illustrates the hierarchy of biological organization, showing the progression from the smallest unit of life to the largest. It is organized into two main columns: **CELLULAR** and **ORGANISMS**.

CELLULAR (Left Column):

- 1. Atom**: The smallest unit of matter.
- 2. Molecule**: A group of atoms bonded together.
- 3. Macromolecule**: A large molecule, such as a protein or nucleic acid.
- 4. Organelle**: A specialized structure within a cell, such as a mitochondrion or nucleus.
- 5. Cell**: The basic unit of life.

ORGANISMS (Right Column):

- 6. Tissue**: A group of similar cells working together.
- 7. Organ**: A structure made of different tissues that performs a specific function.
- 8. Organ system**: A group of organs that work together to perform a specific function.
- 9. Organism**: An individual living entity.
- 10. Population**: A group of individuals of the same species living in the same area.
- 11. Community**: A group of different populations living in the same area.
- 12. Ecosystem**: A community of organisms and their physical environment.
- 13. Biosphere**: The part of Earth where life exists.

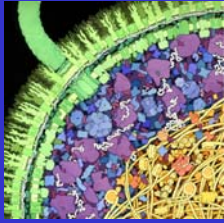
Arrows indicate the flow of organization from the smallest unit (Atom) to the largest (Biosphere). A box at the bottom left specifies the scale of the units:

- 1 cm = 10^7 nm
- 1 μ m = 10^3 nm
- 1 m = 10^9 nm

The diagram consists of a central yellow oval with the text "From Genes to Cell-Function based on Reality". Surrounding this central oval is a larger yellow circle. Four yellow rectangular boxes are positioned around the circle, each containing a challenge:

- Top-left: "too small"
- Top-right: "timescales from nanoseconds to Megaseconds"
- Bottom-right: "too complex"
- Bottom-left: "data - incomplete - wrong - quality?"

Major challenge how to model the cell (for example E.coli) and biochemical pathways?

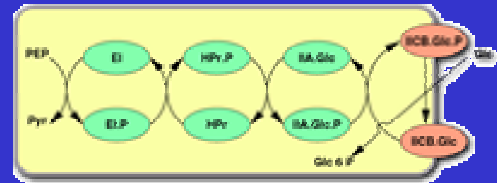


Detail of the E. coli cell



Overview of relevant biochemical pathways

Case study: the Phosphoenolpyruvate dependent phosphotransferase system (PTS) responsible for the uptake of glucose in E. coli



Diagrammatic view of the PTS pathway in E. coli
Problem: part of the pathway is membrane-bound

Computational Biology

Lab sessions
Yves Fomekong Nanfack

Location and time

Monday: 13.00 - 16.00, room P127

Wednesday: 11.00 -14.00, room P126

Lab sessions
start on 5th of February (today!)

Computational Biology

Overview I

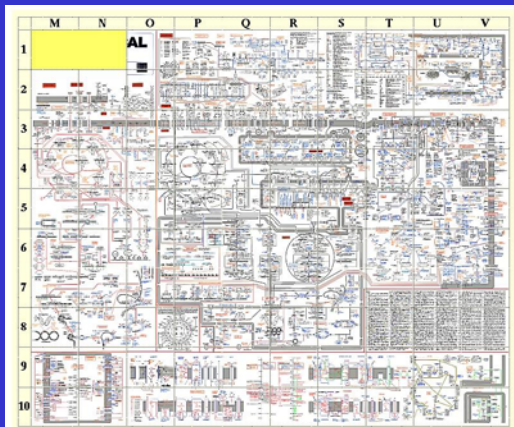
- Monday 5-2-07 Blueprint of life
- Tuesday 6-2-07 cancelled!
- Monday 12-2-07 Blueprint of life, molecular biology databases
- Tuesday 13-2-07 Sequence analysis I
- Monday 19-2-07 Sequence analysis II
- Tuesday 20-2-07 Sequence analysis III
- Monday 26-2-07 Biochemical networks



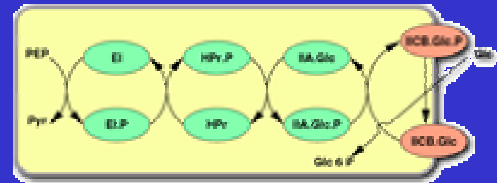
Computational Biology

Overview II

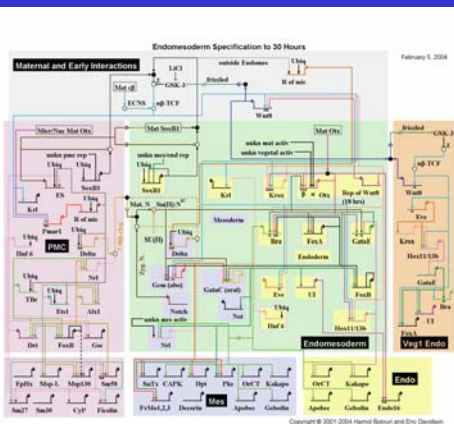
- Tuesday 27-2-07 metabolic networks I
- Monday 5-3-07 metabolic networks II
- Tuesday 6-3-07 Gene regulatory networks I
- Monday 12-3-07 Gene regulatory networks II
- Tuesday 13-3-07 Systems Biology I
- Monday 19-3-07 Systems Biology II
- Tuesday 20-3-07 Systems Biology III



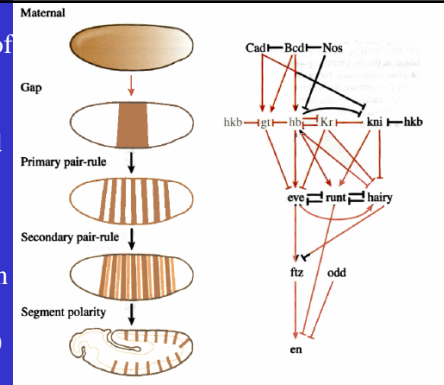
Case study: the Phosphoenolpyruvate dependent phosphotransferase system (PTS) responsible for the uptake of glucose in *E. coli*



Diagrammatic view of the PTS pathway in *E. coli*
Problem: part of the pathway is membrane-bound



Development of morphogen gradients in time (left) and regulatory network for segment specification in *Drosophila* (Carroll, 2001)



Lab assignments

- Introduction solving simple biological problems
- Sequence analysis
- Enzyme kinetics
- Solving non-linear ODE in biological problems
- Compartmental modelling

Lab assignments, slides lectures etc
Computational Biology
Blackboard

Questions (in English) regarding the practical assignments:

Yves Fomekong Nanfack
(yvesf@science.uva.nl)

Computational Biology course material I

- All slides etc on blackboard

Computational Biology course material II (optional) books (in order of being advised)

- Systems Biology in Practice, E Klip et al., Wiley-VCH, Berlin, 2005
- Computational modeling of genetic and biochemical networks, J.M. Bower & H. Bolouri, MIT Press, Cambridge, 2001
- Post-genome informatics, M. Kanehisa, Oxford University Press, 2000
- (Introduction to Bioinformatics, A.M. Lesk, Oxford University Press, 2005 not advised)

Computational Biology, to do:

- Lab assignments
- exam

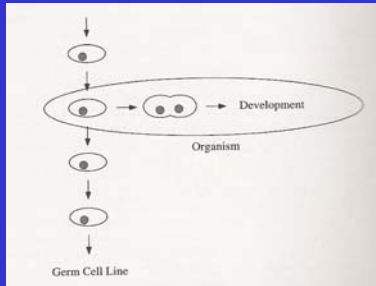
Computational Biology

- Final grade = $\frac{1}{2}$ (lab_assignments + exam)

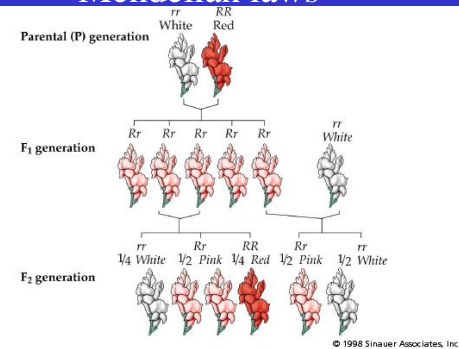
Life is a complex system for information storage and processing

- Vertical information transfer: generation to generation
- Horizontal information transfer: ontogenesis

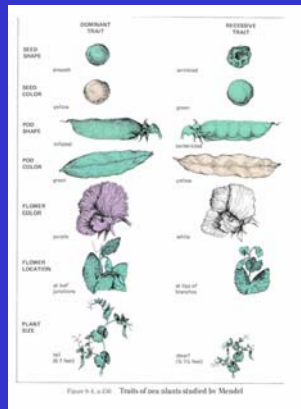
Life is a complex system for information storage and processing



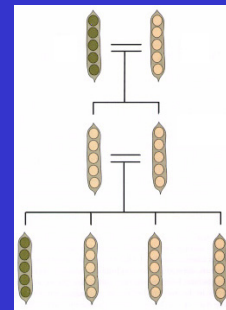
Mendelian laws



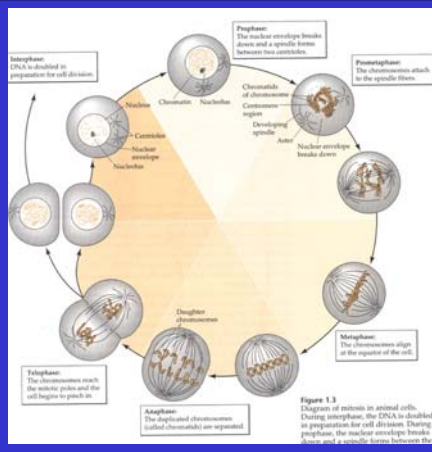
Mendelian laws II



Mendelian laws II



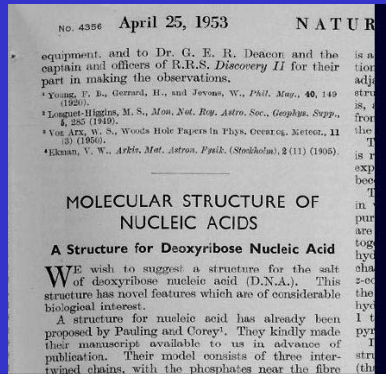
Chromosomes and cell division in animals



Genome and gene

- Genome: unit of information transmission by DNA replication
- Gene : unit of information expression by transcription to RNA or translation to protein

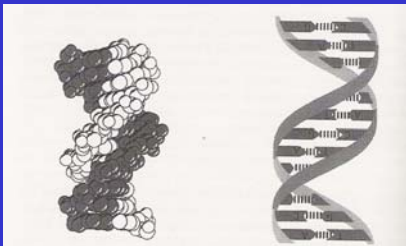
Double helix model of DNA, Watson Crick 1953



Nucleotides

- A Adenine
- G Guanine
- C Cytosine
- T Thymine

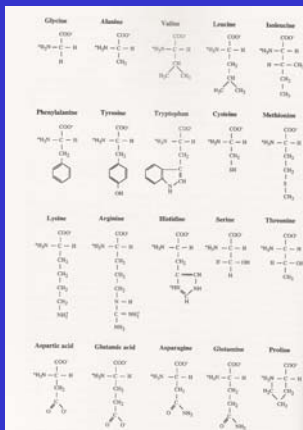
DNA double helix



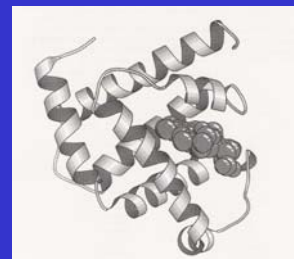
Amino acids

Ala	A	Alanine
Arg	R	Arginine
Asn	N	Asparagine
Asp	D	Aspartic acid
Cys	C	Cysteine
Gln	Q	Glutamine
Glu	E	Glutamic acid
Gly	G	Glycine
His	H	Histidine
Ile	I	Isoleucine
Leu	L	Leucine
Lys	K	Lysine
Met	M	Methionine
Phe	F	Phenylalanine
Pro	P	Proline
Ser	S	Serine
Thr	T	Threonine
Trp	W	Tryptophan
Tyr	Y	Tyrosine
Val	V	Valine
Aax	B	Asn or Asp
Glx	Z	Gln or Glu
Sec	U	Selenocysteine
Unk	X	Unknown

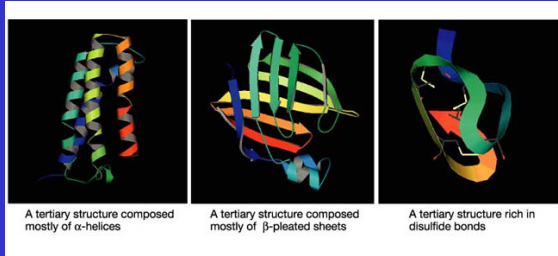
Amino acids



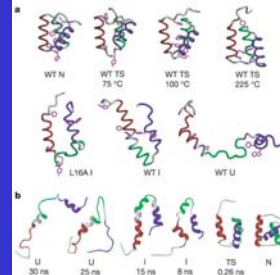
3D protein structure protein: example myoglobin



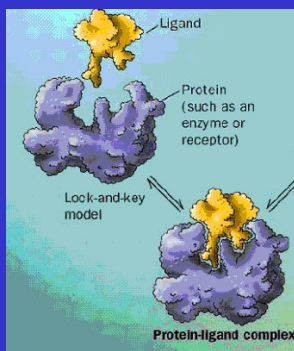
3D protein structure protein: alpha and beta helices



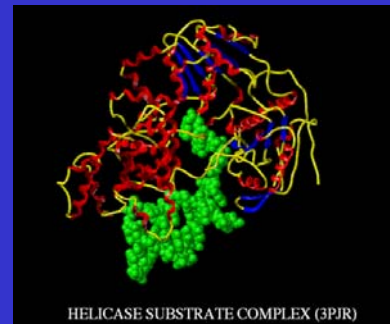
Unfolding and folding of the engrailed homeodomain protein (Molecular Dynamics simulations, Nature 421:863-7, 2003)



Enzyme substrate



Enzyme substrate recognition



Standard genetic code: codon \rightarrow amino acid

	U	C	A	G
U	UUU Phe UUC Phe UUA Leu UUG Leu	UCU Ser UCC Ser UCA Ser UCG Ser	UAU Tyr UAC Tyr UAA Stop UAG Stop	UGU Cys UGC Cys UGA Stop UGG Trp
C	CUU Leu CUC Leu CUA Leu CUG Leu	CCU Pro CCC Pro CCA Pro CCG Pro	CAU His CAC His CAA Gln CAG Gln	CGU Arg CGC Arg CGA Arg CGG Arg
A	AUU Ile AUC Ile AUA Ile AUG Met, Start	ACU Thr ACC Thr ACA Thr ACG Thr	AAU Asn AAC Asn AAA Lys AAG Lys	AGU Ser AGC Ser AGA Arg AGG Arg
G	GUU Val GUC Val GUA Val GUG Val	GCU Ala GCC Ala GCA Ala GCG Ala	GAU Asp GAC Asp GAA Glu GAG Glu	GGU Gly GGC Gly GGA Gly GGG Gly

Nucleotides:

DNA

- A Adenine
- G guanine
- C Cytosine
- T Thymine

RNA

- A Adenine
- G Guanine
- C Cytosine
- U Uracil

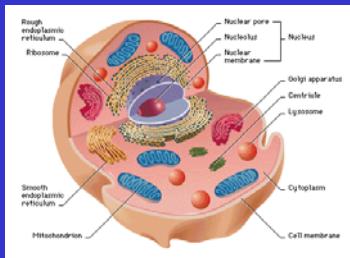
Variations on the standard genetic code: codon>amino acid I

	T1	T2	T3	T4	T5	T6	T9	T10	T12	T13	T14	T15
CUU	Leu	-	Thr	-	-	-	-	-	-	-	-	-
CUC	Leu	-	Thr	-	-	-	-	-	-	-	-	-
CUA	Leu	-	Thr	-	-	-	-	-	Ser	-	-	-
CUG	Leu	-	Thr	-	-	-	-	-	-	-	-	-
AUU	Ile	-	-	-	-	-	-	-	-	-	-	-
AUC	Ile	-	-	-	-	-	-	-	-	-	-	-
AUA	Ile	Met	Met	-	Met	-	-	-	Met	-	-	-
AUG	Met	-	-	-	-	-	-	-	-	-	-	-
UAU	Tyr	-	-	-	-	-	-	-	-	-	-	-
UAC	Tyr	-	-	-	-	-	-	-	-	-	-	-
UAA	Stop	-	-	-	-	Gln	-	-	-	-	Tyr	-
UAG	Stop	-	-	-	-	Gln	-	-	-	-	-	Gln
AAU	Asn	-	-	-	-	-	-	-	-	-	-	-
AAC	Asn	-	-	-	-	-	-	-	-	-	-	-
AAA	Lys	-	-	-	-	-	Asn	-	-	-	Asn	-
AAG	Lys	-	-	-	-	-	-	-	-	-	-	-
UCU	Cys	-	-	-	-	-	-	-	-	-	-	-
UGC	Cys	-	-	-	-	-	-	-	-	-	-	-
UGA	Stop	Trp	Trp	Trp	Trp	-	Trp	Cys	-	Trp	Trp	-
UGG	Trp	-	-	-	-	-	-	-	-	-	-	-
AGU	Ser	-	-	-	-	-	-	-	-	-	-	-
AGC	Ser	-	-	-	-	-	-	-	-	-	-	-
AGA	Arg	Stop	-	-	Ser	-	Ser	-	-	Gly	Ser	-
AGG	Arg	Stop	-	-	Ser	-	Ser	-	-	Gly	Ser	-

Variations on the standard genetic code: codon>amino acid II

- T1 standard code
- T2 vertebrate mitochondrial code
- T3 yeast mitochondrial code
- T4 mould, protozoan and coelenterate mitochondrial code; mycoplasma and spiroplasma code
- T5 invertebrate mitochondrial code
- T6 ciliate, dasycladacean and hexamita nuclear code
- T9 echinoderm mitochondrial code
- T10 euplotid code
- T12 alternative yeast code
- T13 ascidian mitochondrial code
- T14 flatworm mitochondrial code
- T15 blepharisma nuclear code

Animal cell with mitochondria



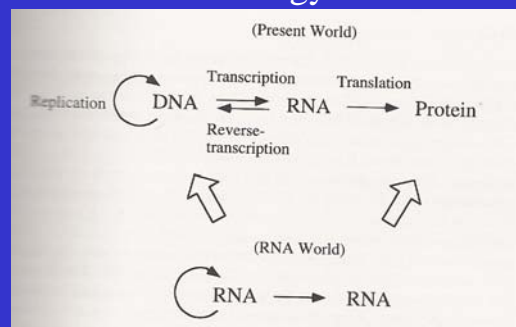
3 types of RNA

- Messenger RNA: mRNA
- Transfer RNA: tRNA
- Ribosomal RNA: rRNA

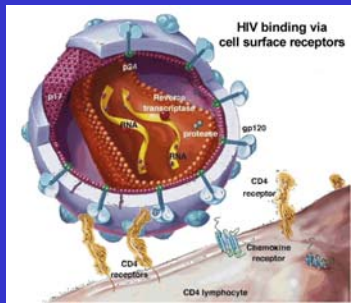
Nucleotides: DNA (deoxyribose backbone) RNA (ribose)

- | | |
|--------------|------------|
| • A Adenine | A Adenine |
| • G guanine | G Guanine |
| • C Cytosine | C Cytosine |
| • T Thymine | U Uracil |

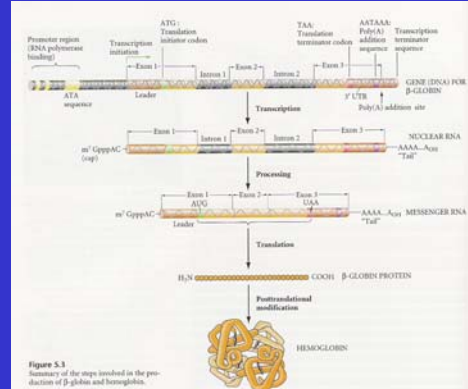
Central dogma of molecular biology



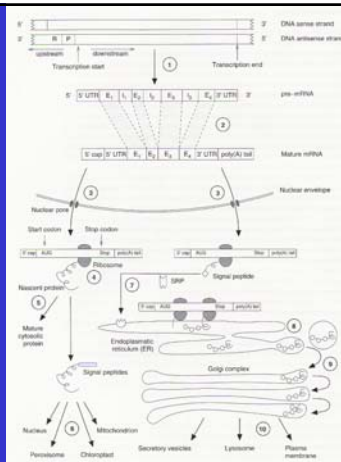
RNA virus: HIV



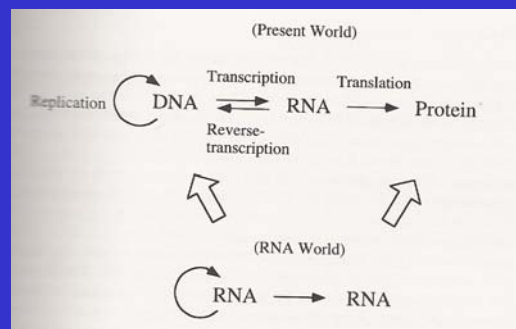
mRNA splicing (in animals)



Transcription and processing of mRNA (in eukaryotes, for example animals)



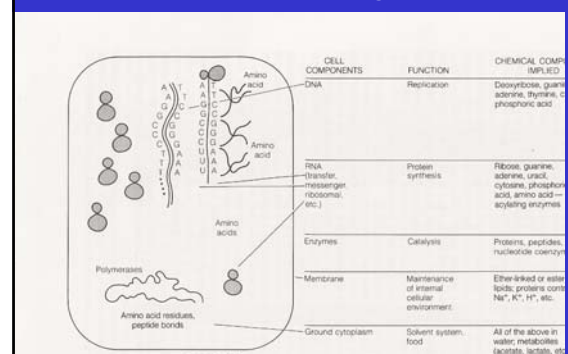
RNA world



Definition of life

- Self-maintaining metabolic system capable of reproduction

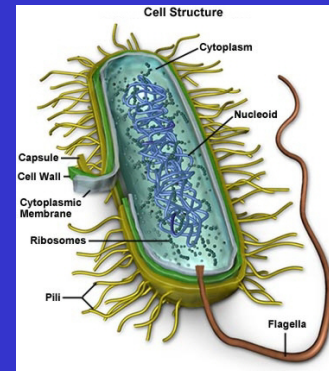
Minimal cell (Margulis, 1993)



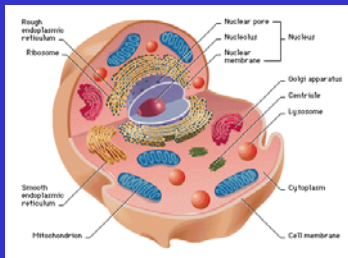
Minimal cell II (Margulis, 1993)

- Cells, minimal autopoietic systems, are membrane entities, the smallest capable of self-reproduction. Even the tiniest cells today contain genes in the form of DNA molecules, single or in several copies, a protein-synthesizing machinery composed of several types of RNA, and many proteins. All cells contain ribosomes, bodies about 0.02μ m in diameter composed of at least three kinds of RNA and about 50 different proteins

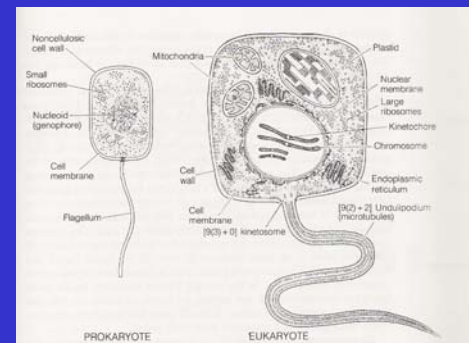
Prokaryotic cell (E. coli)



Eukaryotic cell (animal cell)



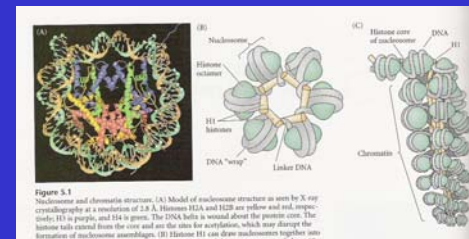
Prokaryotic and Eukaryotic cell



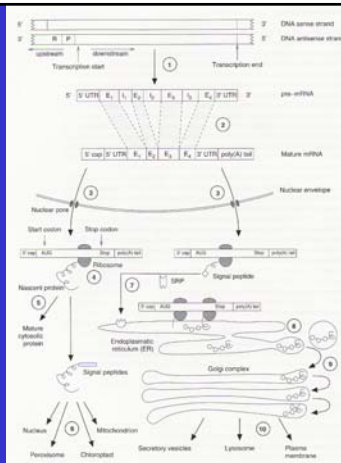
Differences prokaryotes and eukaryotes

- No distinct compartments in prokaryotes: no nucleus, no organelles (chloroplasts, mitochondria, endoplasmic reticulum, Golgi complex, centriole, lysosome, peroxisome, etc.)
- Genome is not packed into a nucleus, no chromatin, no chromosomes, in prokaryotes
- Cell is smaller (in general) in prokaryotes ($1-10\mu$ length; eukaryotes $10-100\mu$ length)
- Prokaryotes mostly single cellular, many eukaryotes are multi-cellular

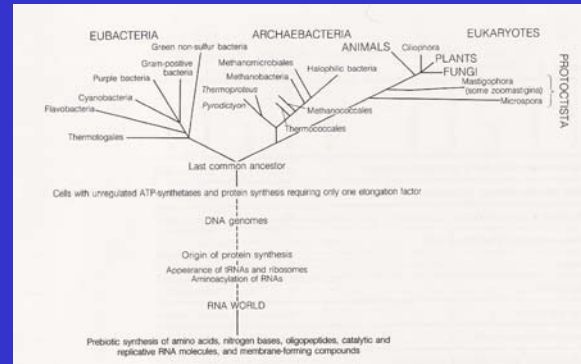
Chromatin in eukaryotes



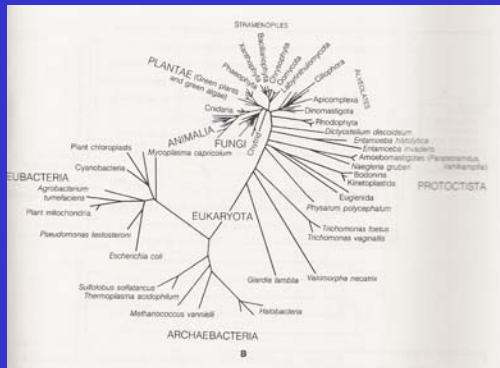
After transcription and processing of mRNA (in eukaryotes, for example animals)



Tree of life (Margulis, 1993; based on RNA polymerases)



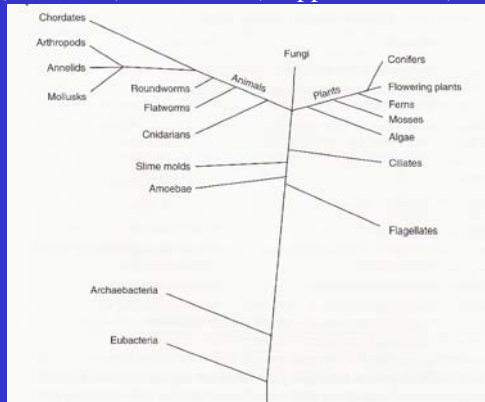
Tree of life (Margulis, 1993; based on ribosomal RNA sequences)



Example
Rhodophyta (red seaweeds)



(Incorrect!) tree of life (Klipp et al, 2005) II



Tree of life
symbiogenesis
(Margulis, 1993)

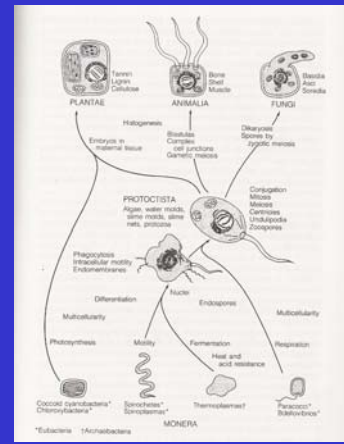
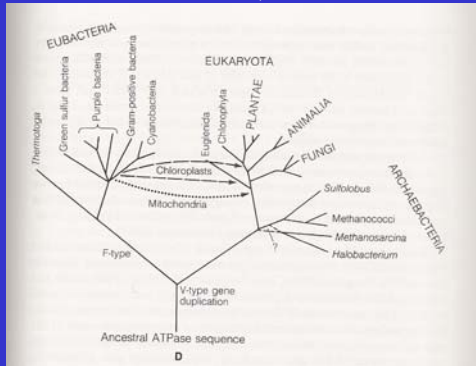


Diagram tree of life symbiogenesis (Margulis, 1993) II



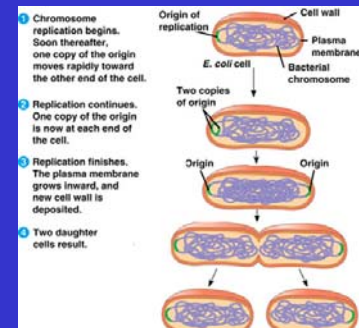
Variations on the standard genetic code: codon>amino acid I

	T1	T2	T3	T4	T5	T6	T9	T10	T12	T13	T14	T15
CUU	Leu	-	Thr	-	-	-	-	-	-	-	-	-
CUC	Leu	-	Thr	-	-	-	-	-	-	-	-	-
CUA	Leu	-	Thr	-	-	-	-	-	Ser	-	-	-
CUG	Leu	-	Thr	-	-	-	-	-	-	-	-	-
AUU	Ile	-	-	-	-	-	-	-	-	-	-	-
AUC	Ile	-	-	-	-	-	-	-	-	-	-	-
AUA	Ile	Met	Met	-	Met	-	-	-	-	Met	-	-
AUG	Met	-	-	-	-	-	-	-	-	-	-	-
UAU	Tyr	-	-	-	-	-	-	-	-	-	-	-
UAC	Tyr	-	-	-	-	-	-	-	-	-	-	-
UAA	Stop	-	-	-	-	Gln	-	-	-	-	Tyr	-
UAG	Stop	-	-	-	-	Gln	-	-	-	-	-	Gln
AAU	Asn	-	-	-	-	-	-	-	-	-	-	-
AAC	Asn	-	-	-	-	-	-	-	-	-	-	-
AAA	Lys	-	-	-	-	-	Asn	-	-	-	Asn	-
AAG	Lys	-	-	-	-	-	-	-	-	-	-	-
UGU	Cys	-	-	-	-	-	-	-	-	-	-	-
UGC	Cys	-	-	-	-	-	-	-	-	-	-	-
UGA	Stop	Trp	Trp	Trp	Trp	-	Trp	Cys	-	Trp	Trp	-
UGG	Trp	-	-	-	-	-	-	-	-	-	-	-
AGU	Ser	-	-	-	-	-	-	-	-	-	-	-
AGC	Ser	-	-	-	-	-	-	-	-	-	-	-
AGA	Arg	Stop	-	-	Ser	-	Ser	-	-	Gly	Ser	-
AGG	Arg	Stop	-	-	Ser	-	Ser	-	-	Gly	Ser	-

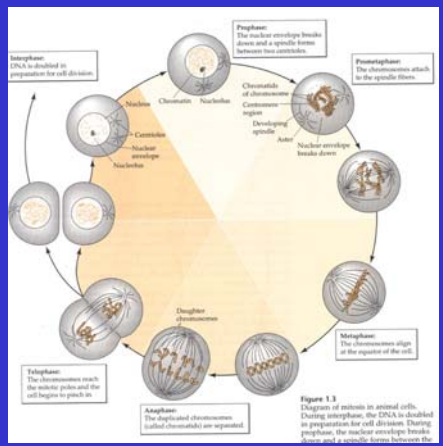
Variations on the standard genetic code: codon>amino acid II

- T1 standard code
- T2 vertebrate mitochondrial code
- T3 yeast mitochondrial code
- T4 mould, protozoan and coelenterate mitochondrial code; mycoplasma and spiroplasma code
- T5 invertebrate mitochondrial code
- T6 ciliate, dasycladacean and hexamita nuclear code
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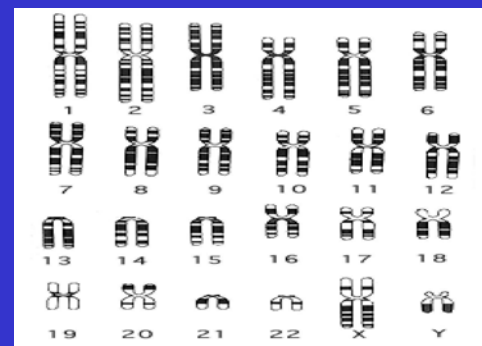
Cell division in bacteria (note there are actually no chromosomes in bacteria!)



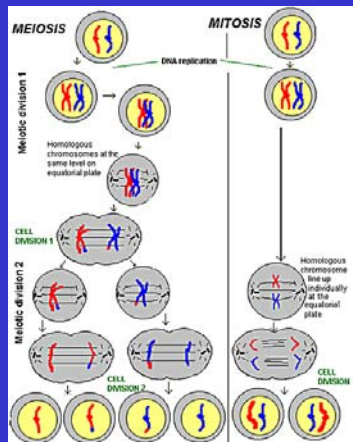
Cell division in animals (eukaryotes) and cell cycle



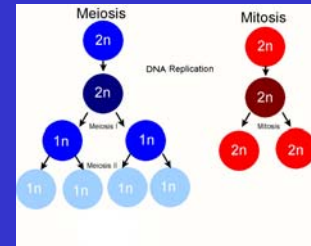
Human chromosomes



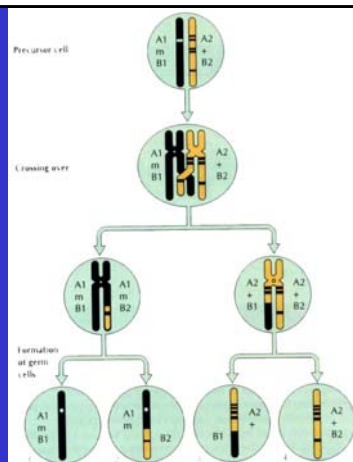
Mitosis and meiosis in animals (eukaryotes) I



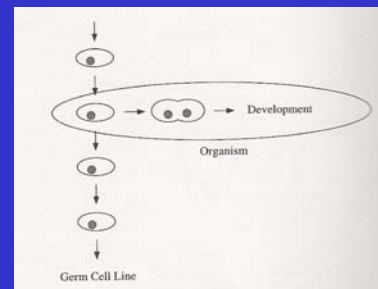
Mitosis and meiosis in animals (eukaryotes) II



Recombination of chromosomes (for example in animals)



Life is a complex system for information storage and processing + cell division



Computational Biology

End Part I blueprint of life
Jaap Kaandorp

12 ECTS course bioinformatics at Academic Medical Center Amsterdam

- Block 2b
- 12 ECTS
- See website programme bioinformatics MGC <http://www.science.uva.nl/bioinformatics> for details
- Registration via email contact Dr A.H. van Kampen (a.h.vankampen@amc.uva.nl)
- Number of students limited