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The holism versus reductionism debate

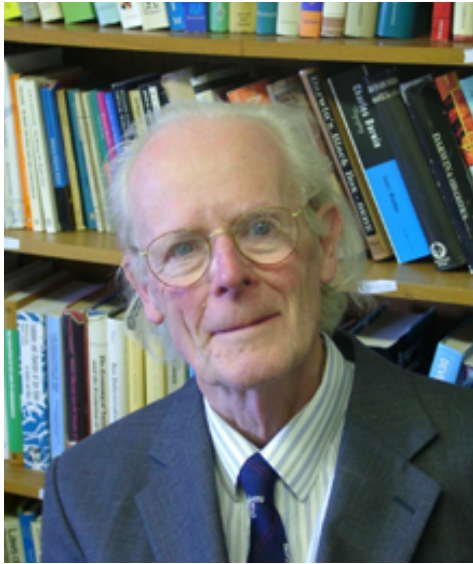
The holistic approach

Macroscopic biologists aim at a top-down approach to describe the phenomena observed in biology.



The reductionists' program

Molecular biologist perform a bottom-up approach to interpret biological phenomena by the methods of chemistry and physics.



What should be the attitude of a biologist working on whole organisms to molecular biology? It is, I think, foolish to argue that we (the macroscopic biologists) are discovering things that disprove molecular biology. It would be more sensible to say to molecular biologists that there are phenomena that they will one day have to interpret in their terms.

John Maynard Smith, *The problems of biology*.
Oxford University Press, 1986.

Genomics and proteomics

Large scale data processing,
sequence comparison, ...

Evolutionary biology

Optimization through variation and
selection, relation between genotype,
phenotype, and function, ...

Structural biology

Protein structures, nucleic
acid structures, supramolecular
complexes, molecular machines, ...

Complexity in 21st Century's Life Sciences

Neurobiology

Neural networks, nonlinear
dynamics, collective properties,
signalling, ...

Systems biology, cell biology

Gene regulation, cell cycle,
metabolic networks, reaction
kinetics, homeostasis, ...

Immunology

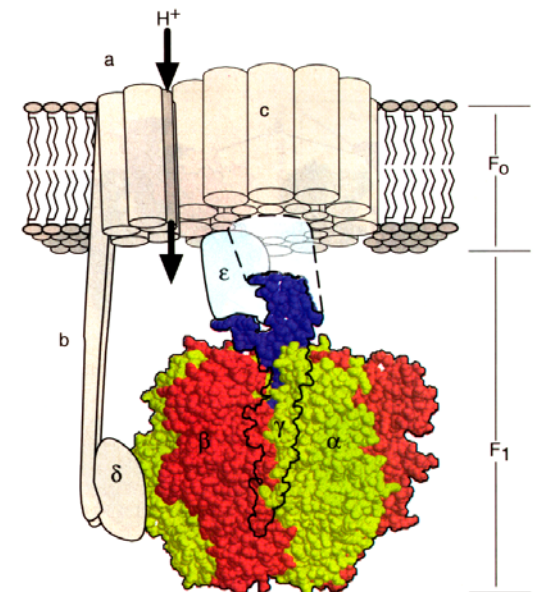
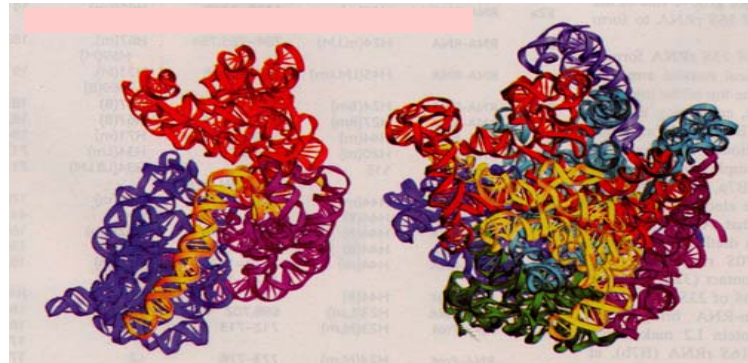
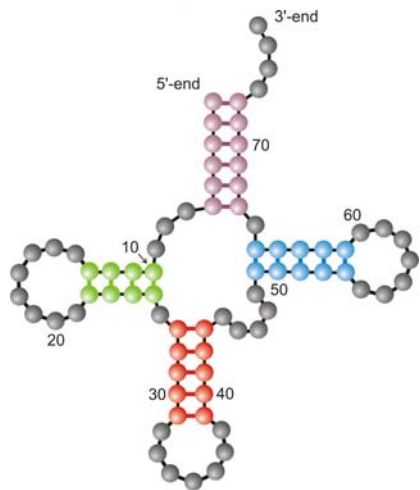
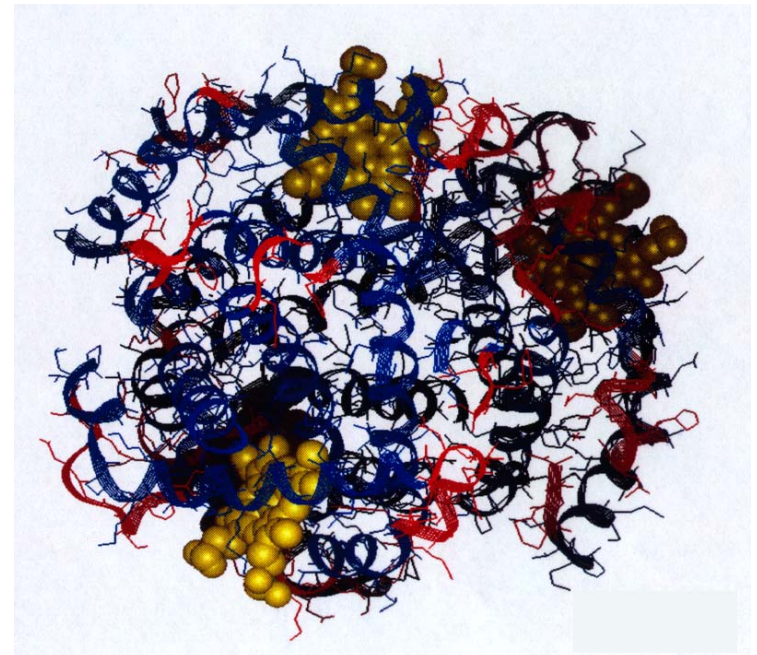
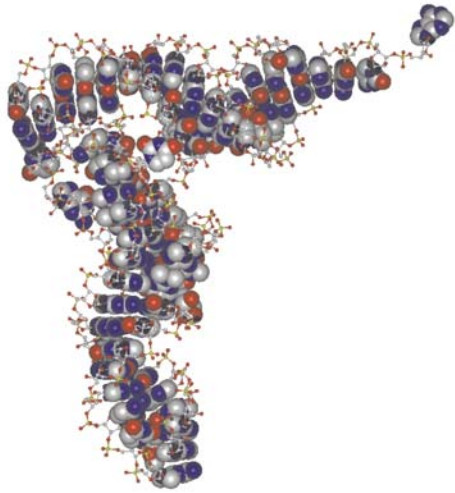
Network theory, immunological
synapse, dynamical systems,
mutation, selection, ...

Developmental biology

Gene regulation networks,
signal propagation, pattern
formation, robustness, ...

Structural biology

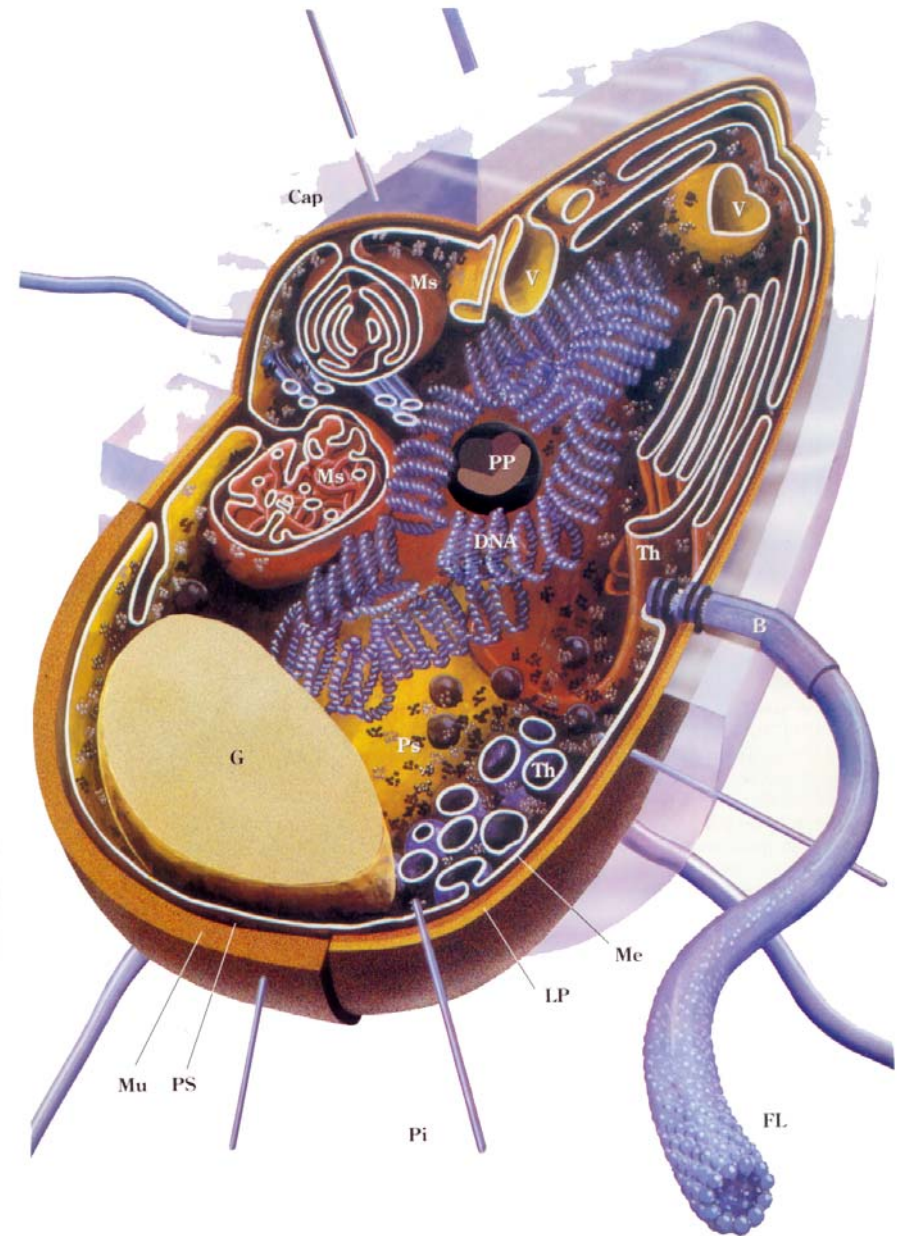
Protein structures, nucleic acid structures, supramolecular complexes, molecular machines, ...



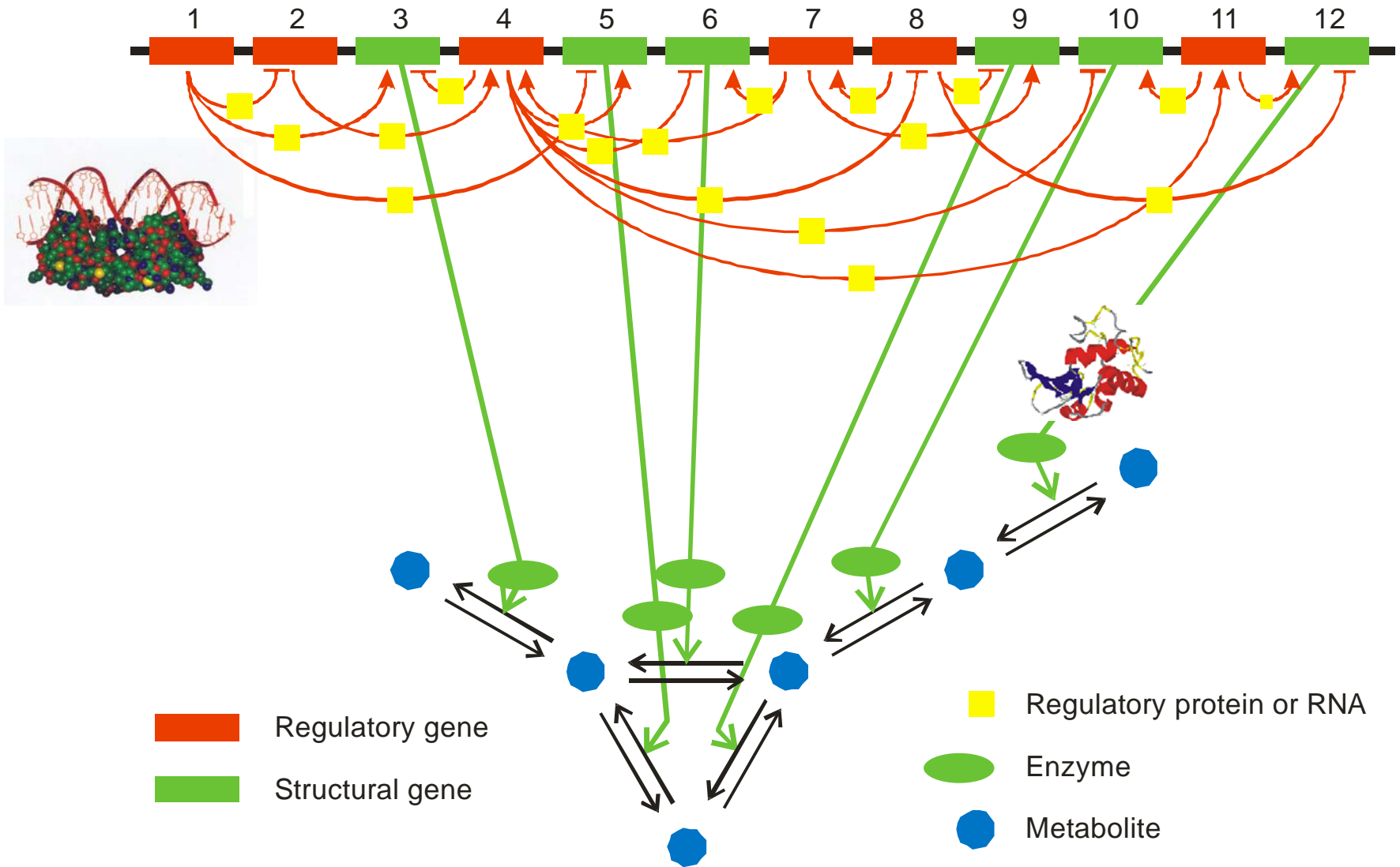
Systems biology, cell biology

Regulation of the cell cycle,
genetic and metabolic networks,
reaction kinetics, homeostasis, ...

The bacterial cell as an example
for the simplest form of
autonomous life



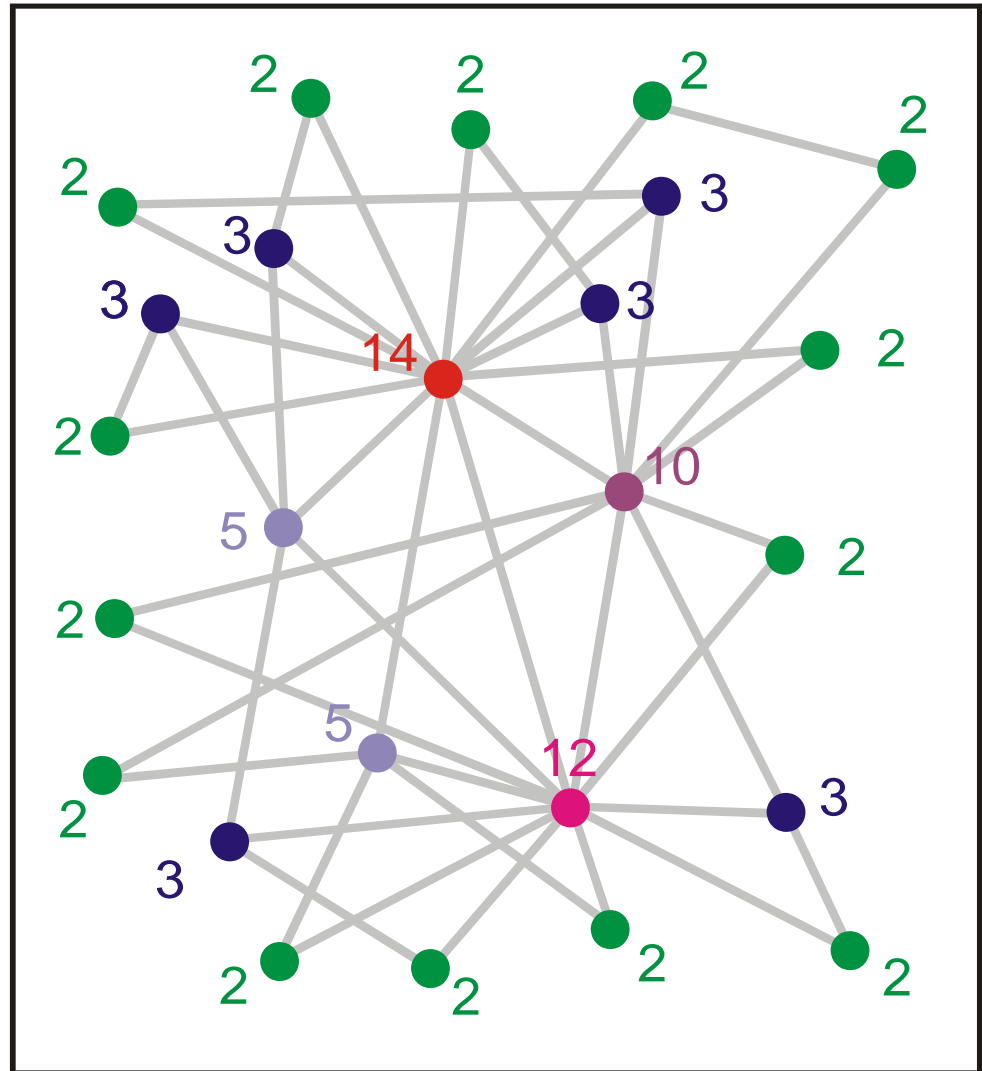
A model genome with 12 genes



Sketch of a genetic and metabolic network

links # nodes

2	14
3	6
5	2
10	1
12	1
14	1



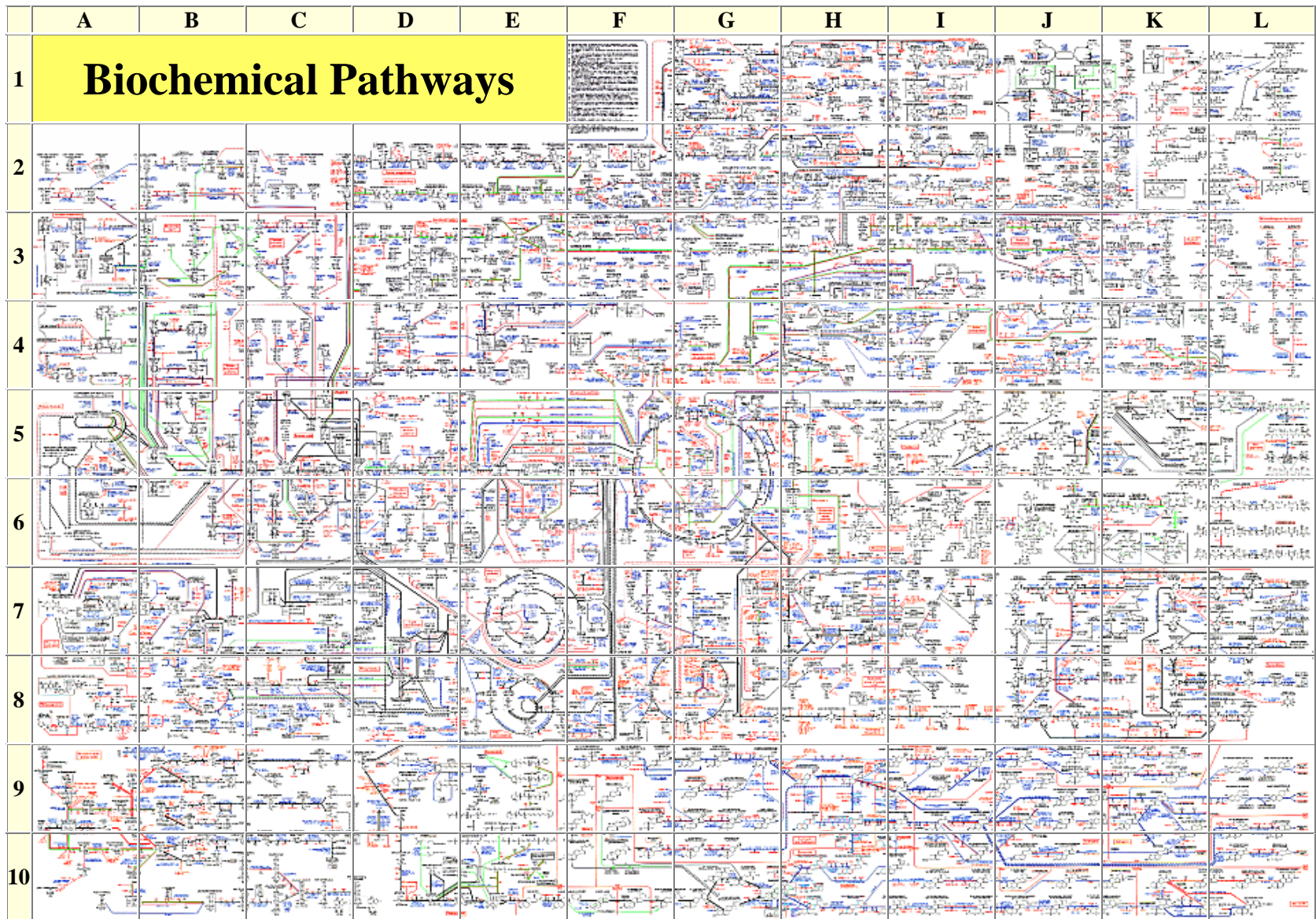
Analysis of nodes and links in a step by step evolved network

E. coli:	Length of the Genome	4×10^6 Nucleotides
	Number of Cell Types	1
	Number of Genes	4 000
Man:	Length of the Genome	3×10^9 Nucleotides
	Number of Cell Types	200
	Number of Genes	30 000 - 60 000

The human body

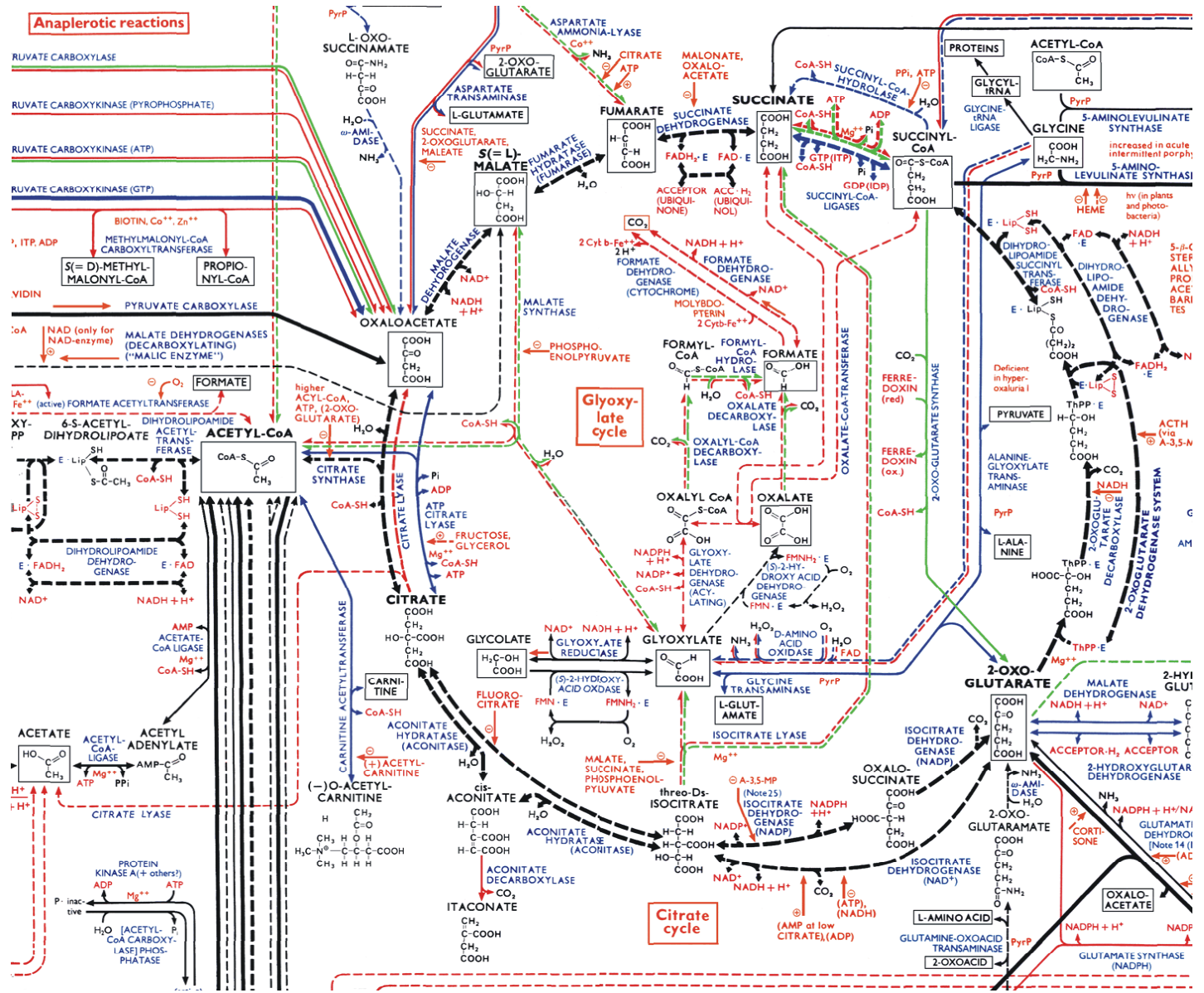
10^{14} cells = 10^{13} eukaryotic cells + 9×10^{13} bacterial (prokaryotic) cells

100 kg = 99.1 kg + 0.9 kg



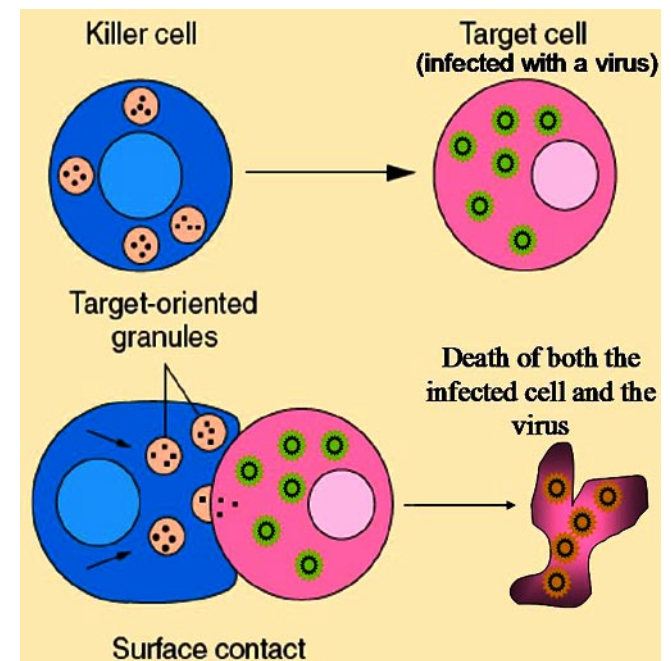
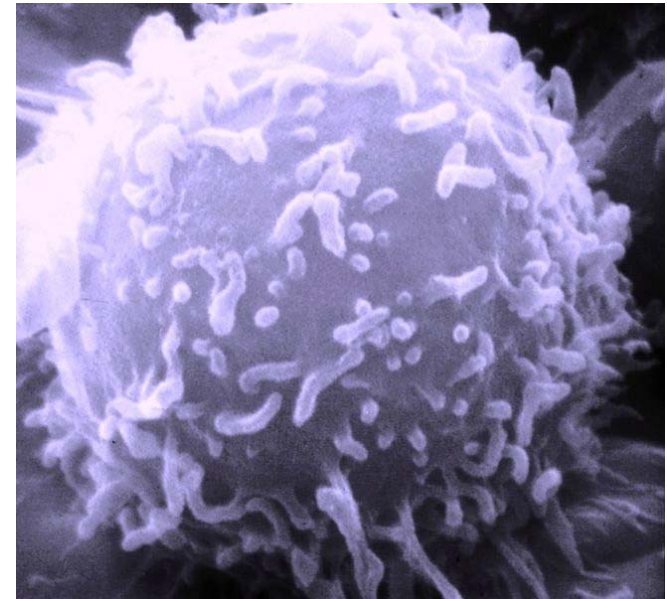
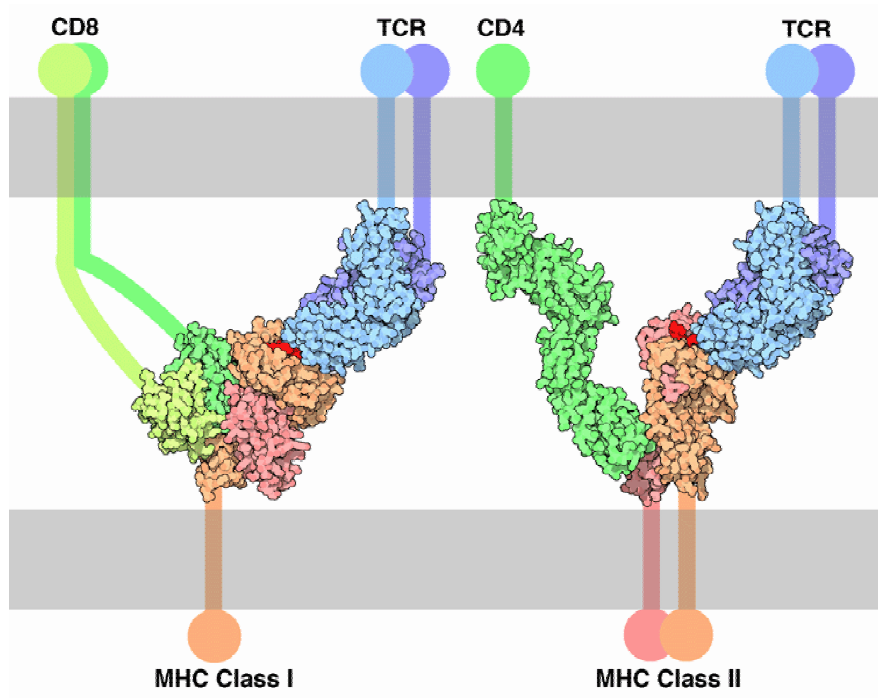
The reaction network of cellular metabolism published by Boehringer-Ingelheim.

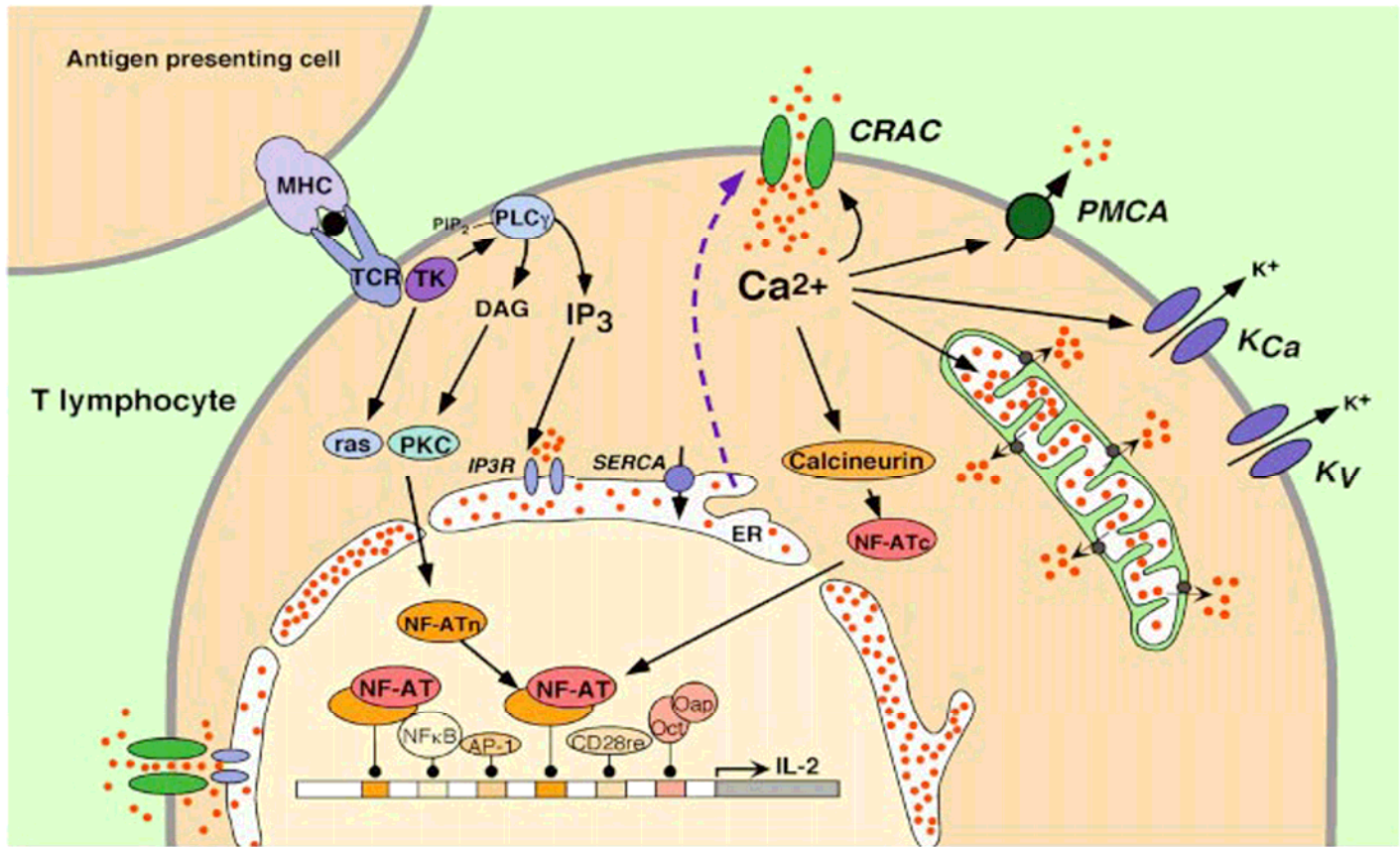
The citric acid or Krebs cycle (enlarged from previous slide).



Immunology

Network theory, immunological synapse, dynamical systems, mutation, selection, ...





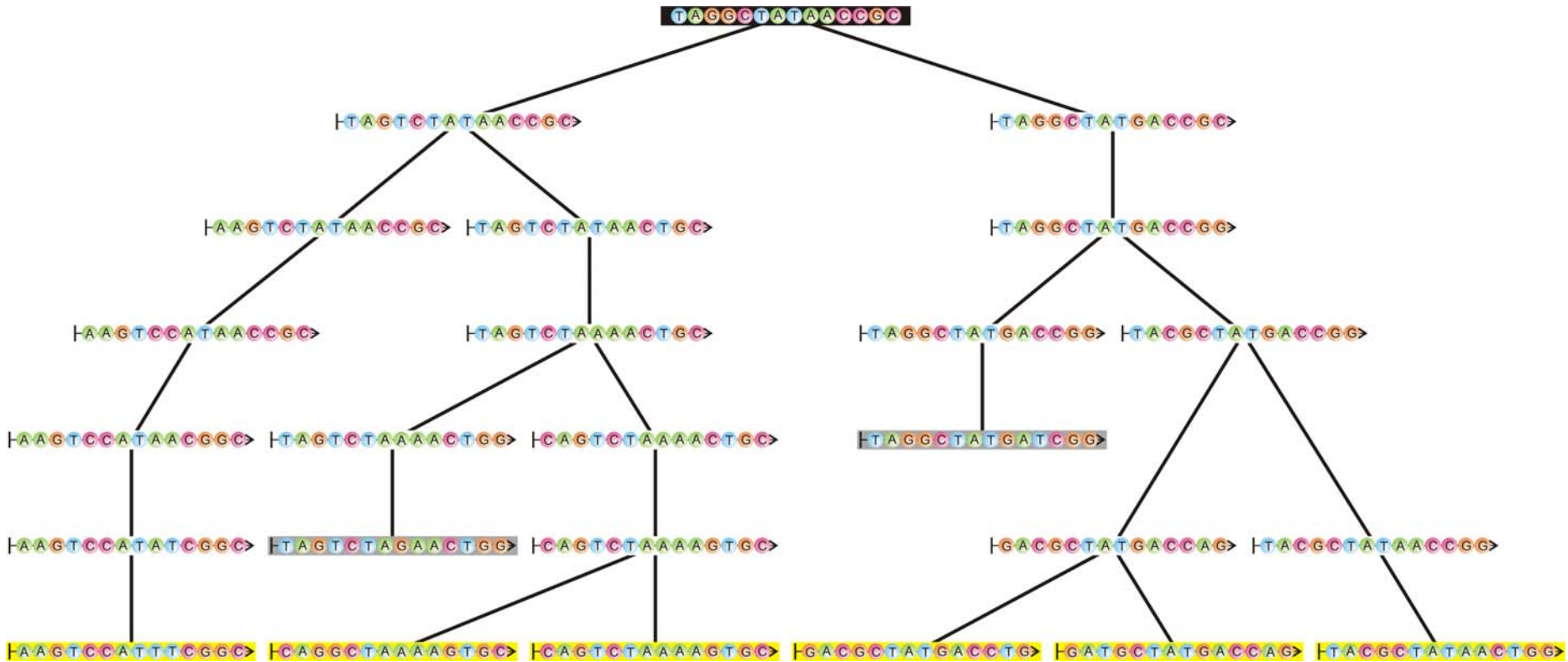
Ca²⁺-signalling at the immune synapse

Evolutionary biology

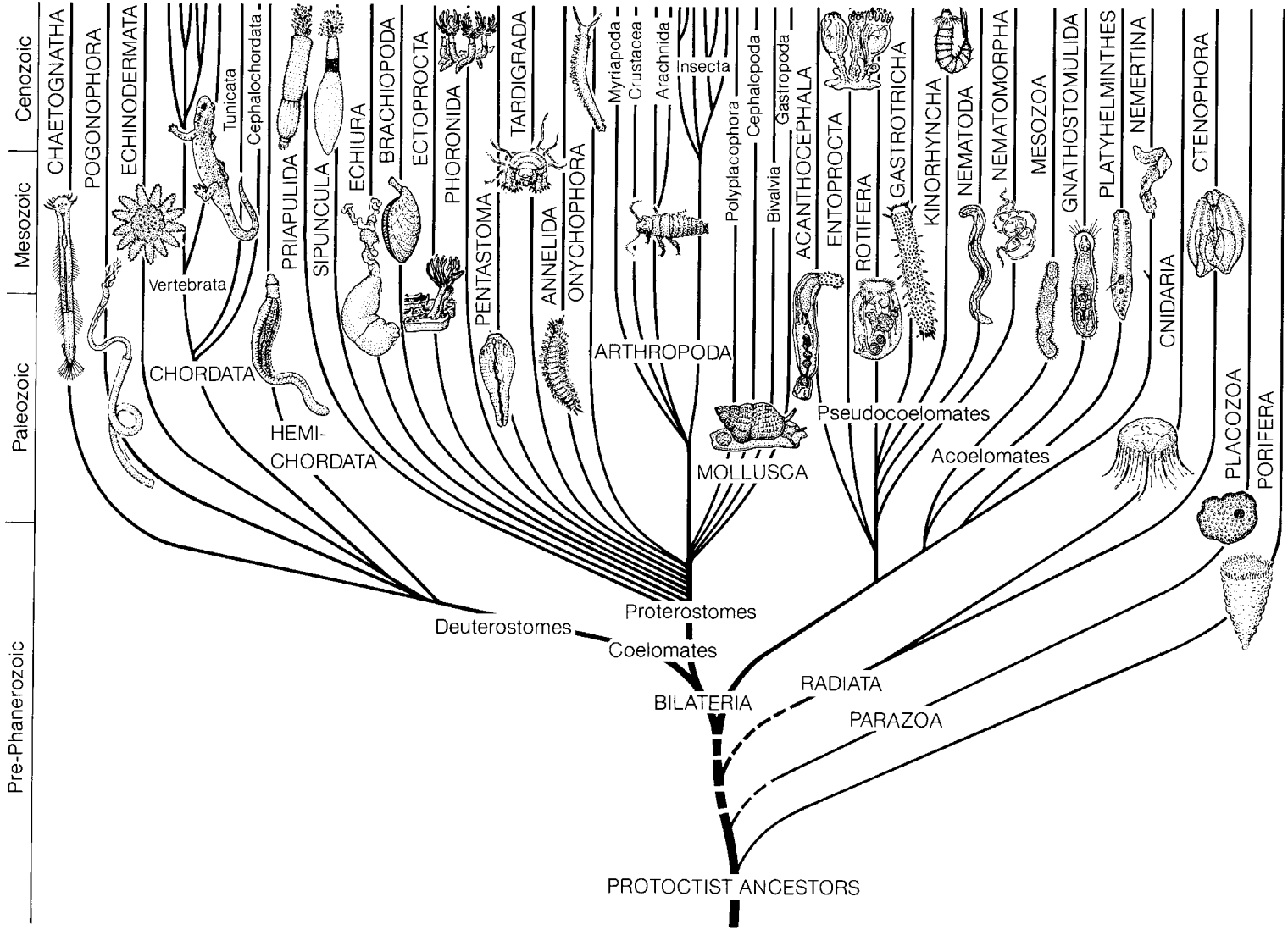
Optimization through variation and selection, relation between genotype, phenotype, and function, ...

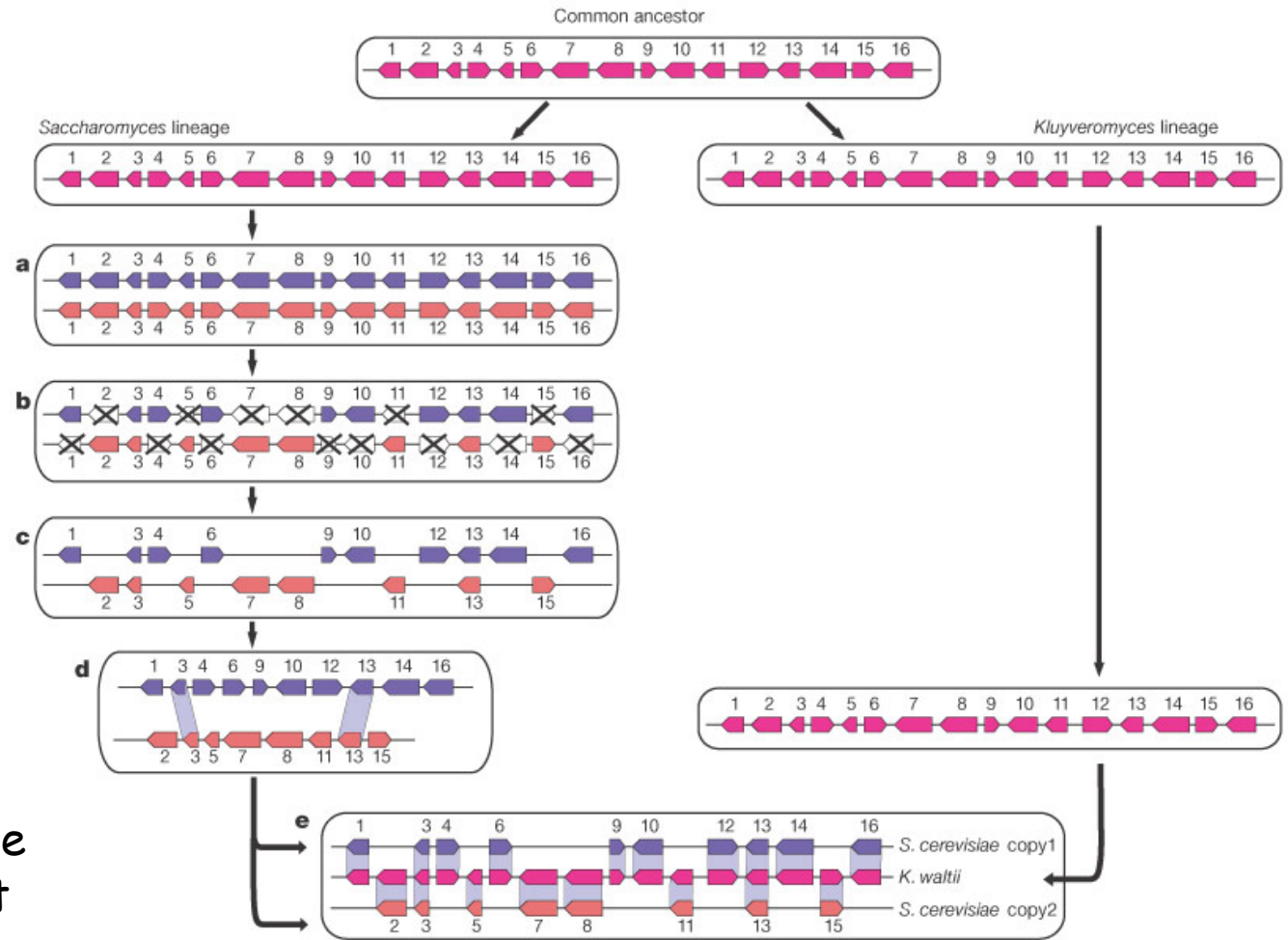
	Generation time	Selection and adaptation 10 000 generations	Genetic drift in small populations 10^6 generations	Genetic drift in large populations 10^7 generations
RNA molecules	10 sec 1 min	27.8 h = 1.16 d 6.94 d	115.7 d 1.90 a	3.17 a 19.01 a
Bacteria	20 min 10 h	138.9 d 11.40 a	38.03 a 1 140 a	380 a 11 408 a
Multicellular organisms	10 d 20 a	274 a 200 000 a	27 380 a 2×10^7 a	273 800 a 2×10^8 a

Time scales of evolutionary change



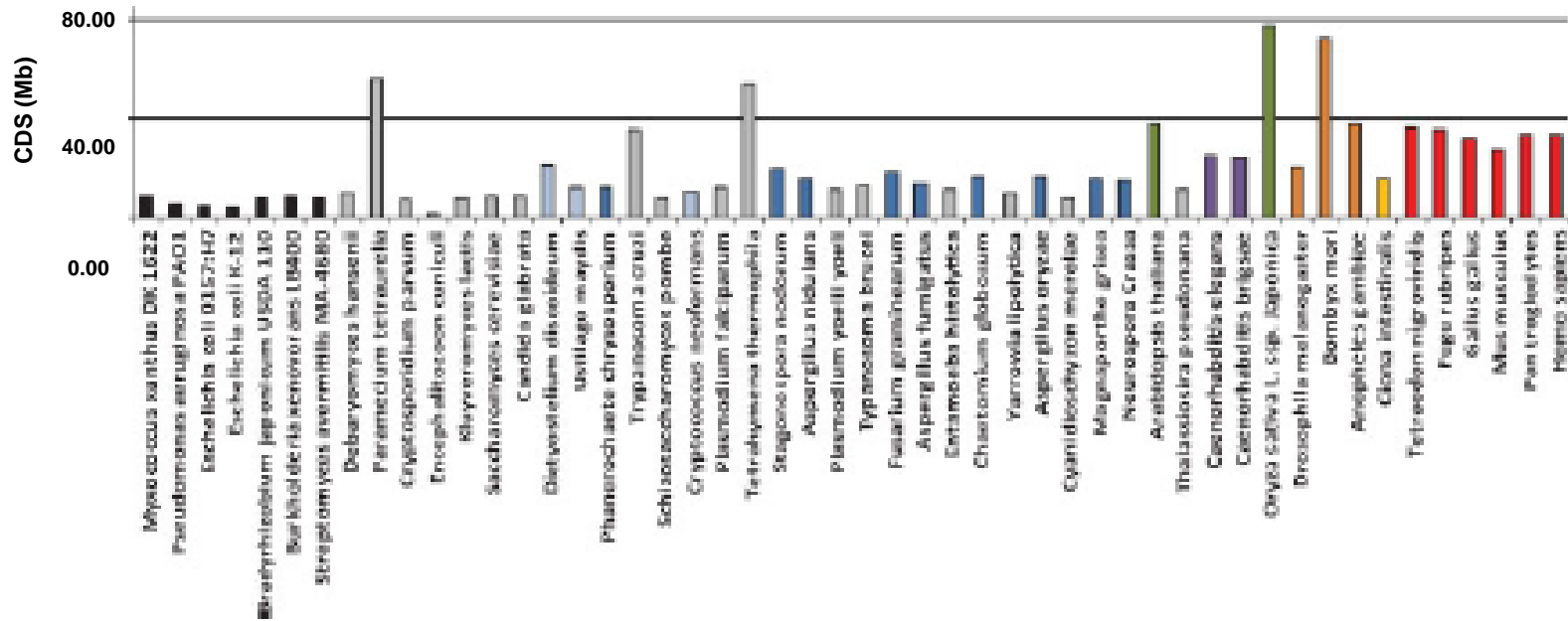
Reconstruction of a phylogenetic tree from present day sequences





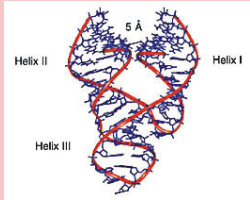
A model for genome duplication in yeast $\approx 1 \times 10^8$ years ago

Manolis Kellis, Bruce W. Birren, and Eric S. Lander. Proof and evolutionary analysis of ancient genome duplication in the yeast *Saccharomyces cerevisiae*. *Nature* **428**: 617-624, 2004



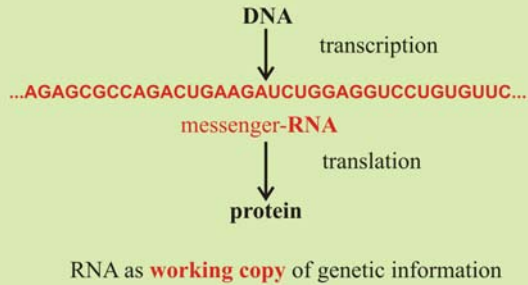
R.J.Taft, M.Pheasant, J.S.Mattick. The relationship between non-protein-coding DNA and eukaryotic complexity. *BioEssays* 29:288-297, 2007.

RNA as catalyst

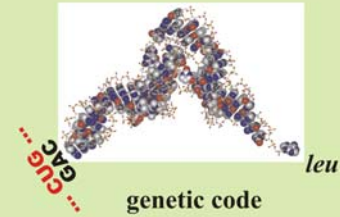


Ribozyme

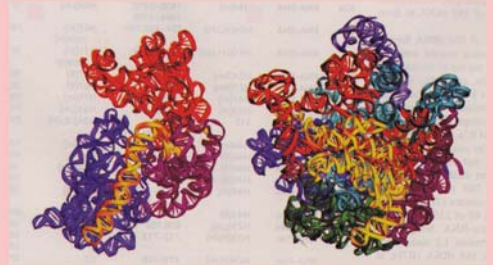
RNA as transmitter of genetic information



RNA as adapter molecule



RNA is the catalytic subunit in supramolecular complexes

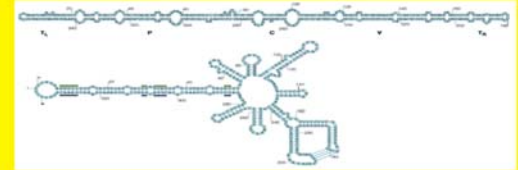


The **ribosome** is a **ribozyme** !

RNA

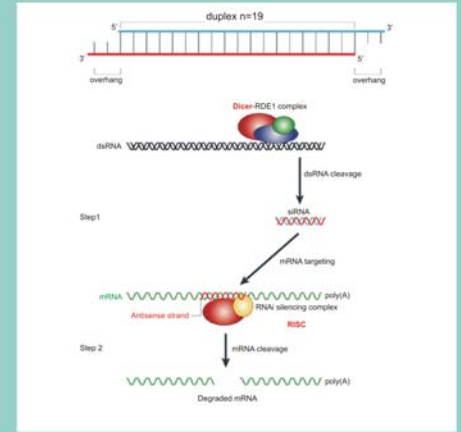
RNA is modified by epigenetic control

RNA editing, alternative splicing



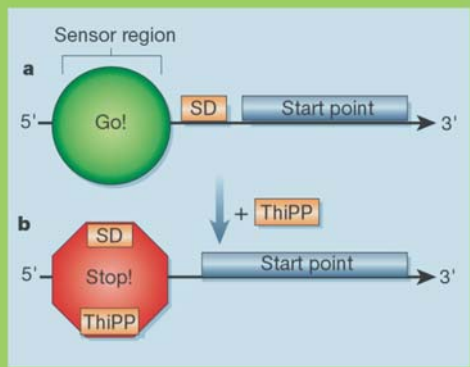
Viroids

RNA as regulator of gene expression



Gene silencing by siRNA

Allosteric control of transcribed RNA



Riboswitches controlled by metabolites

The RNA world as a precursor of the current DNA + protein biology

RNA as carrier of genetic information

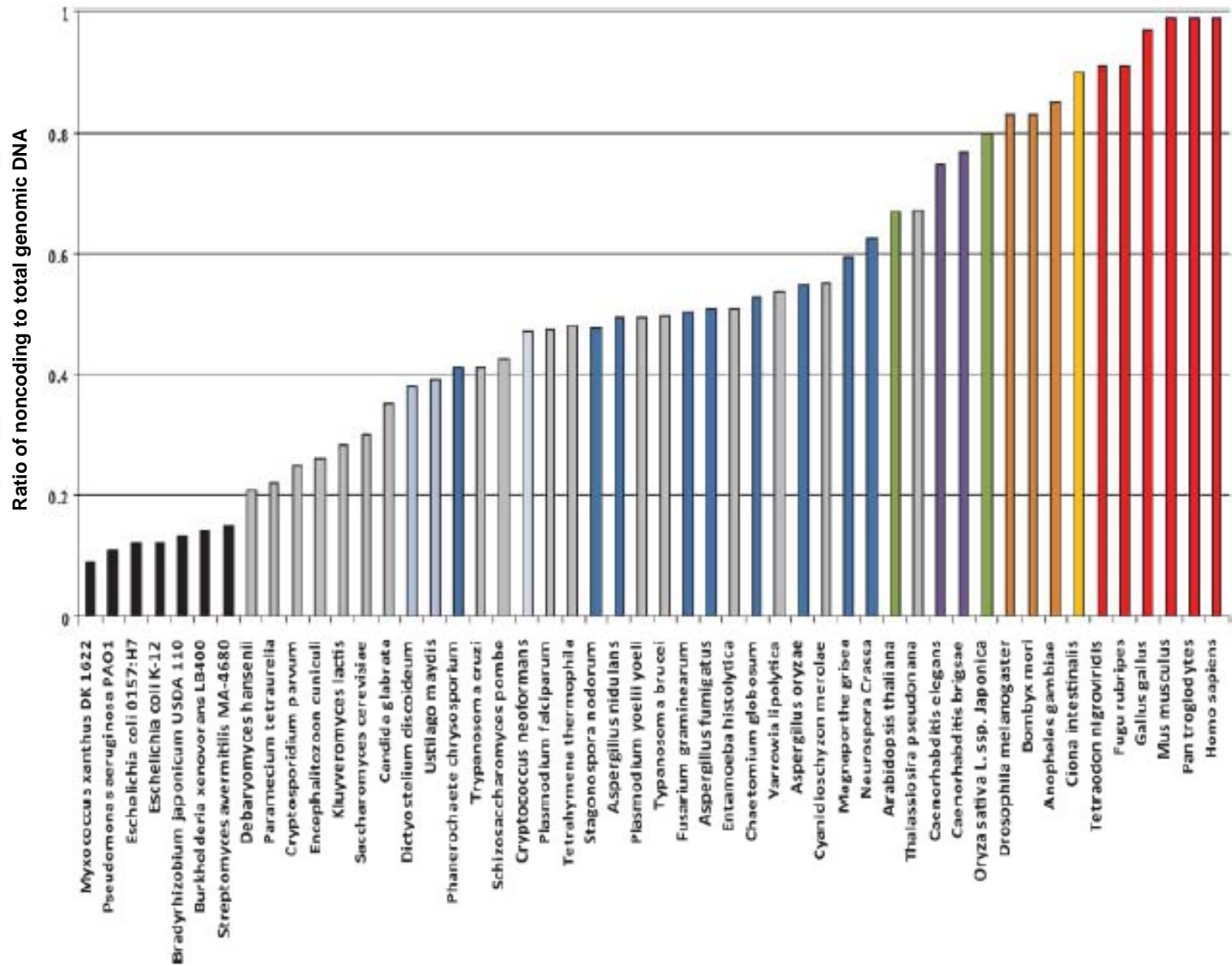
RNA viruses and retroviruses

RNA evolution *in vitro*

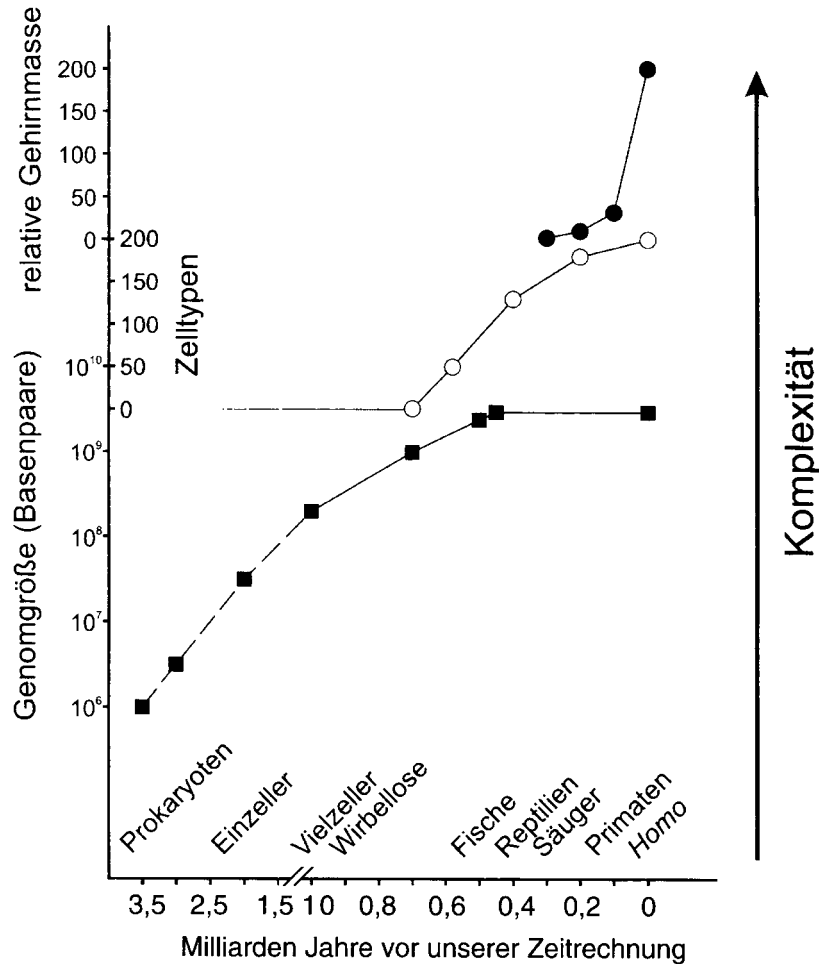
Evolutionary biotechnology

RNA aptamers, artificial ribozymes, allosteric ribozymes

Functions of RNA molecules



R.J.Taft, M.Pheasant, J.S.Mattick. The relationship between non-protein-coding DNA and eukayotic complexity. *BioEssays* 29:288-297, 2007.



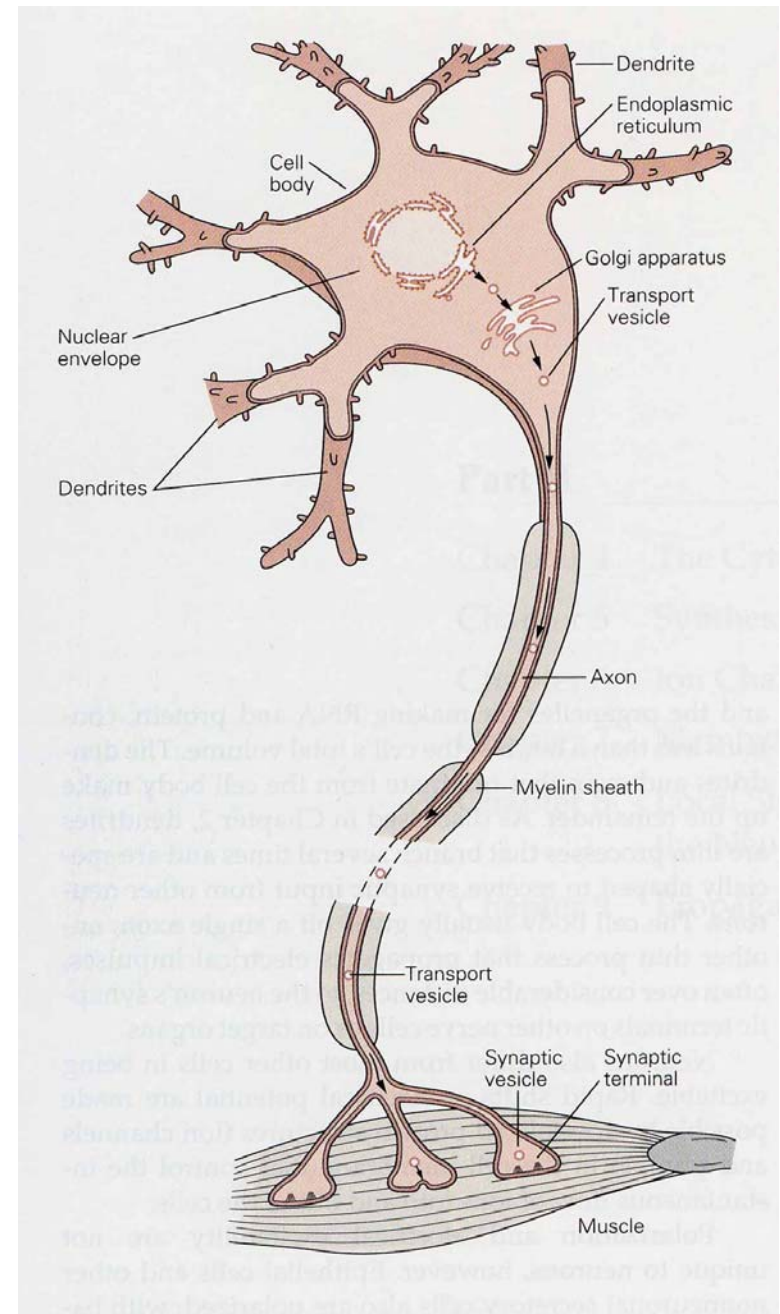
4.10 Die Zunahme der Komplexität ist ein wesentlicher Aspekt der biologischen Evolution, wobei höhere Komplexität sowohl durch Vergrößerung der Zahl von miteinander in Wechselwirkung stehenden Elementen als auch durch Differenzierung der Funktionen dieser Elemente entstehen kann. In dieser Abbildung wird zwischen drei Phasen oder Strategien der Evolution von Komplexität unterschieden. *Untere Kurve*: Zunahme der Genomgröße; logarithmische Auftragung der Zahl der Basenpaare im Genom von Zellen seit Beginn der biologischen Evolution (Daten aus Abbildung 2.3). *Mittlere Kurve*: Zunahme der Zahl der Zelltypen in der Evolution der Metazoa (Daten aus Abbildung 4.8). *Obere Kurve*: Zunahme des relativen Gehirngewichts (bezogen auf die Körperoberfläche) bei Säugetieren (Daten aus Wilson 1985). Für die Abszisse wurden zwei Skaleneinteilungen verwendet, eine für den Zeitraum >10⁹ Jahre, eine andere für den Zeitraum <10⁹ Jahre vor der Gegenwart. Oberhalb der Abszisse sind die Namen einiger wichtiger taxonomischer Einheiten angeführt, deren Evolution in etwa beim jeweiligen Wortbeginn einsetzt.

Wolfgang Wieser. *Die Erfindung der Individualität oder die zwei Gesichter der Evolution*. Spektrum Akademischer Verlag, Heidelberg 1998.

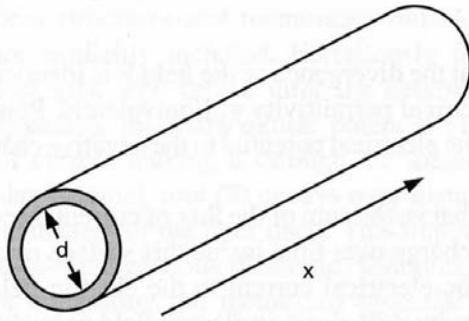
A.C.Wilson. *The Molecular Basis of Evolution*. Scientific American, Oct.1985, 164-173.

Neurobiology

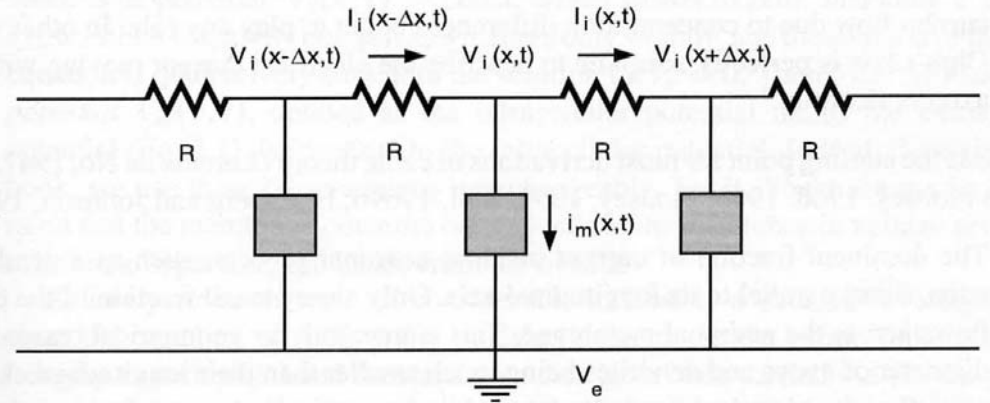
Neural networks, collective properties, nonlinear dynamics, signalling, ...



A single neuron signaling to a muscle fiber



A



B

Fig. 2.2 ELECTRICAL STRUCTURE OF A CABLE (A) Idealized cylindrical axon or dendrite at the heart of one-dimensional cable theory. Almost all of the current inside the cylinder is longitudinal due to geometrical (the radius is much smaller than the length of the cable) and electrical factors (the membrane covering the axon or dendrite possesses a very high resistivity compared to the intracellular cytoplasm). As a consequence, the radial and angular components of the current can be neglected, and the problem of determining the potential in these structures can be reduced from three spatial dimensions to a single one. On the basis of the bidomain approximation, gradients in the extracellular potentials are neglected and the cable problem is expressed in terms of the transmembrane potential $V_m(x, t) = V_i(x, t) - V_e$. (B) Equivalent electrical structure of an arbitrary neuronal process. The intracellular cytoplasm is modeled by the purely ohmic resistance R . This tacitly assumes that movement of carriers is exclusively due to drift along the voltage gradient and not to diffusion. Here and in the following the extracellular resistance is assumed to be negligible and V_e is set to zero. The current per unit length across the membrane, whether it is passive or contains voltage-dependent elements, is described by i_m and the system is characterized by the second-order differential equation, Eq. 2.5.

$$\frac{1}{R} \frac{\partial^2 V}{\partial \xi^2} = C_M \theta \frac{\partial V}{\partial \xi} + [g_{Na} m^3 h (V - V_{Na}) + g_K n^4 (V - V_K) + g_l (V - V_l)] 2\pi r L$$

$$\theta \frac{\partial m}{\partial \xi} = \alpha_m (1 - m) - \beta_m m$$

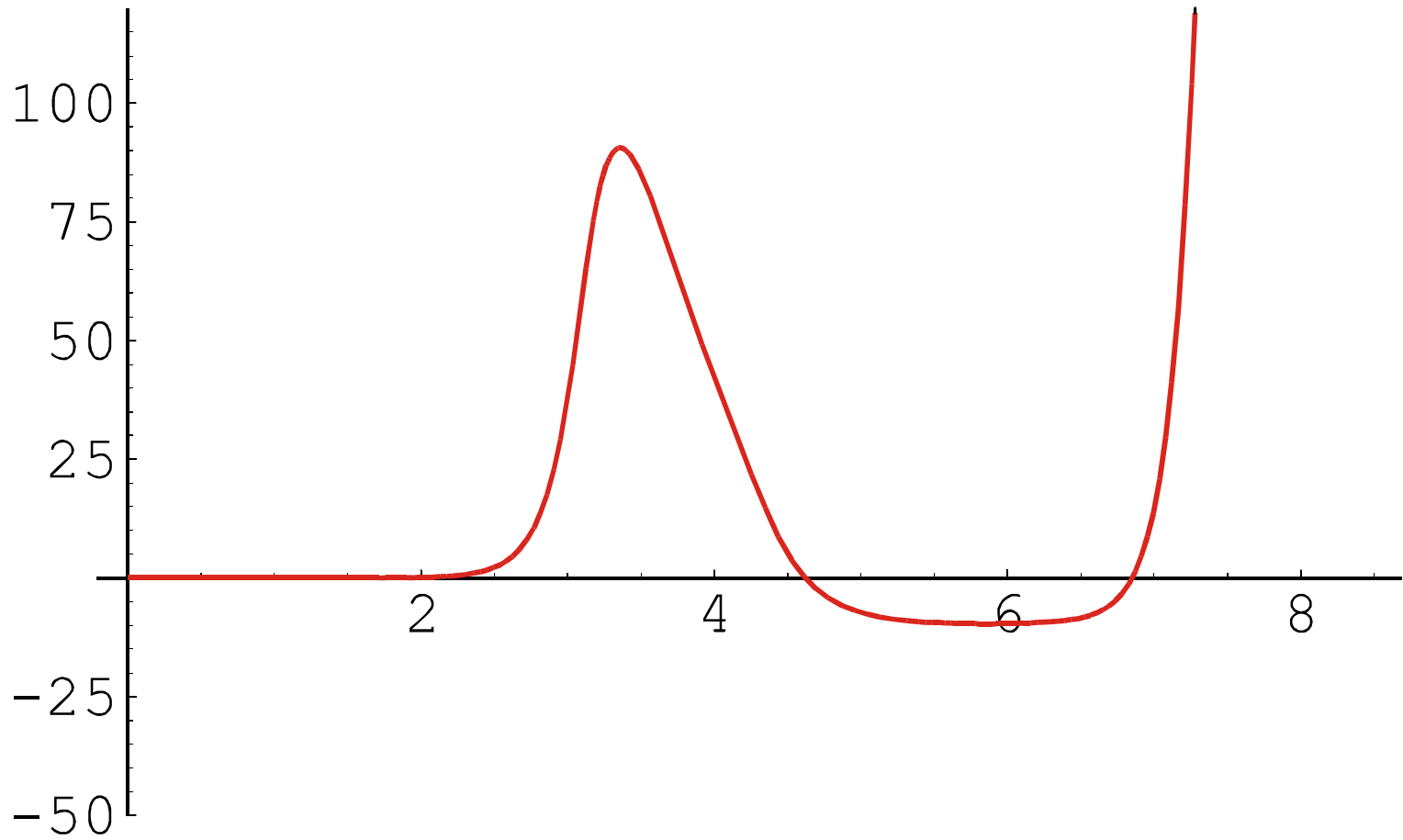
$$\theta \frac{\partial h}{\partial \xi} = \alpha_h (1 - h) - \beta_h h$$

$$\theta \frac{\partial n}{\partial \xi} = \alpha_n (1 - n) - \beta_n n$$

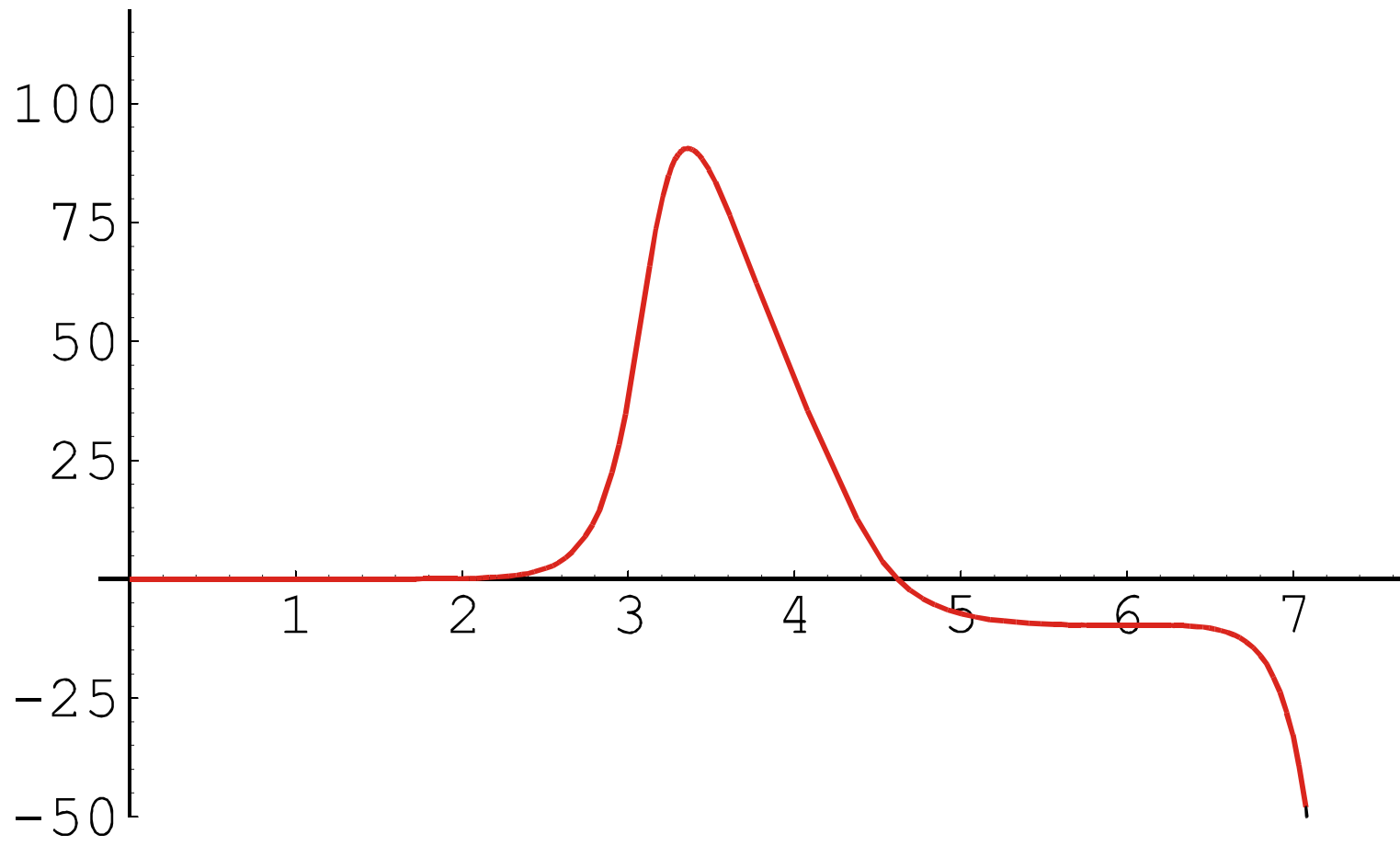
Hodgkin-Huxley ordinary differential equations
(ODE)

Travelling pulse solution: $V(x, t) = V(\xi)$ with
 $\xi = x + \theta t$

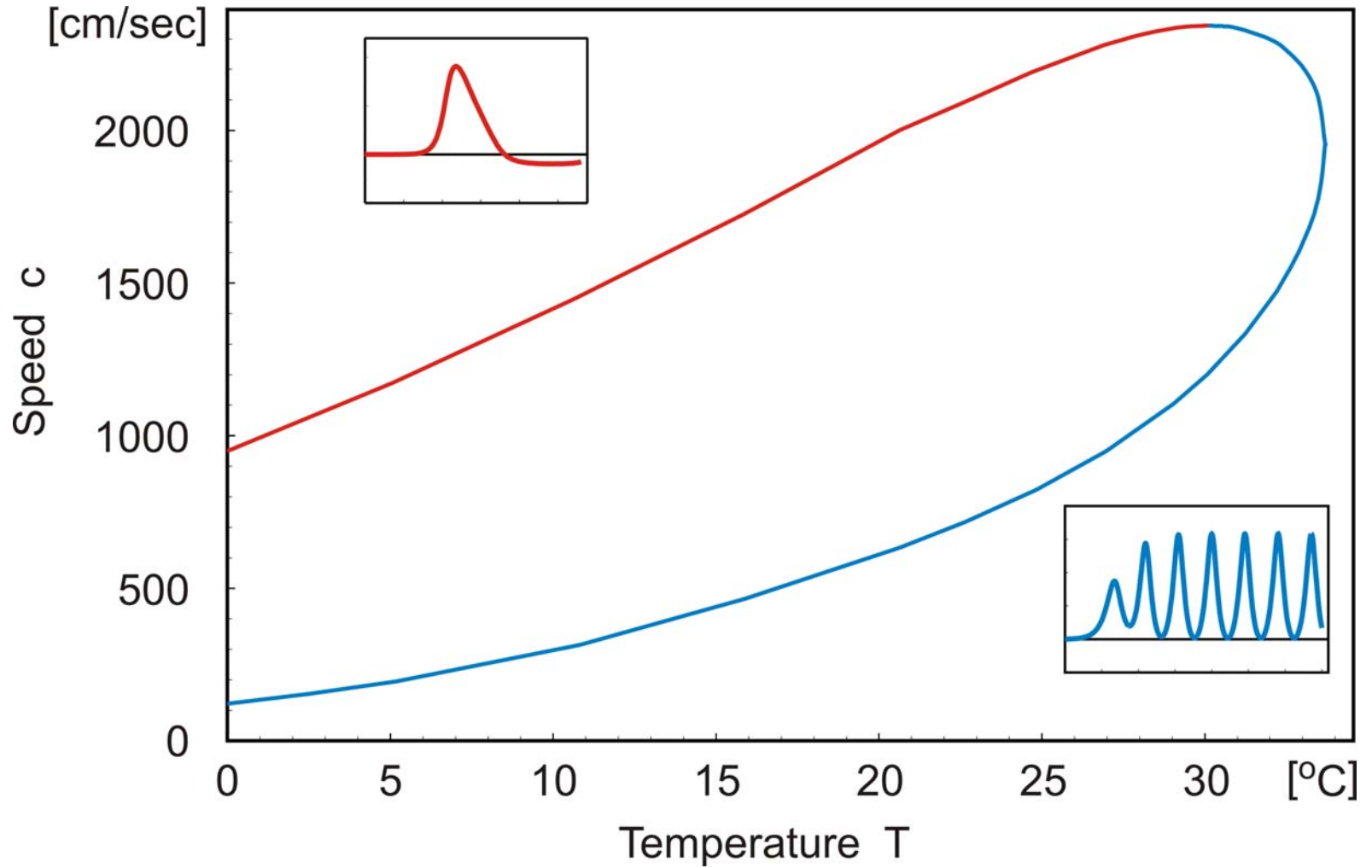
Hodgkin-Huxley equations describing pulse propagation along nerve fibers



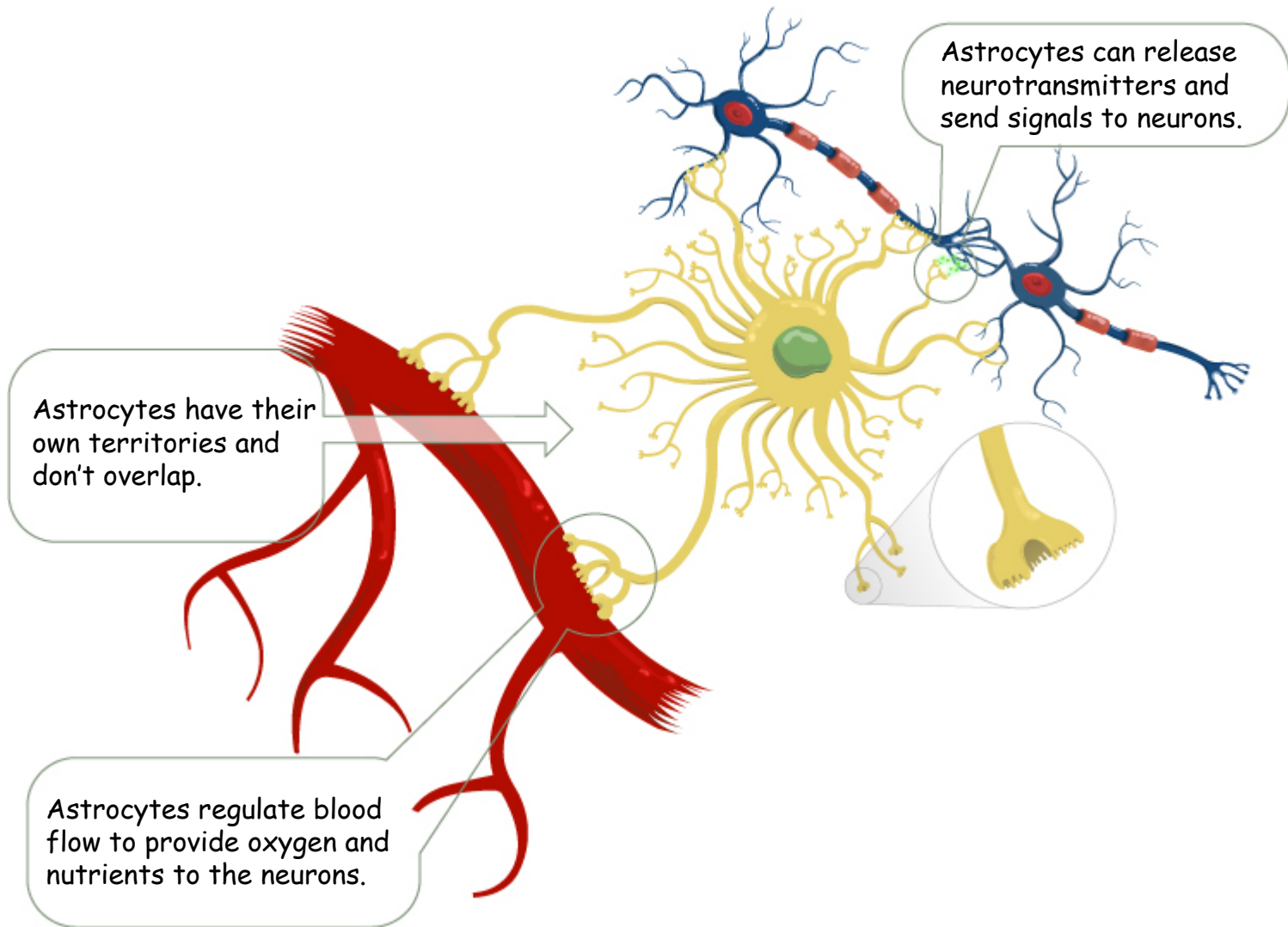
$T = 18.5 \text{ C}; \theta = 1873.3324514717698 \text{ cm / sec}$



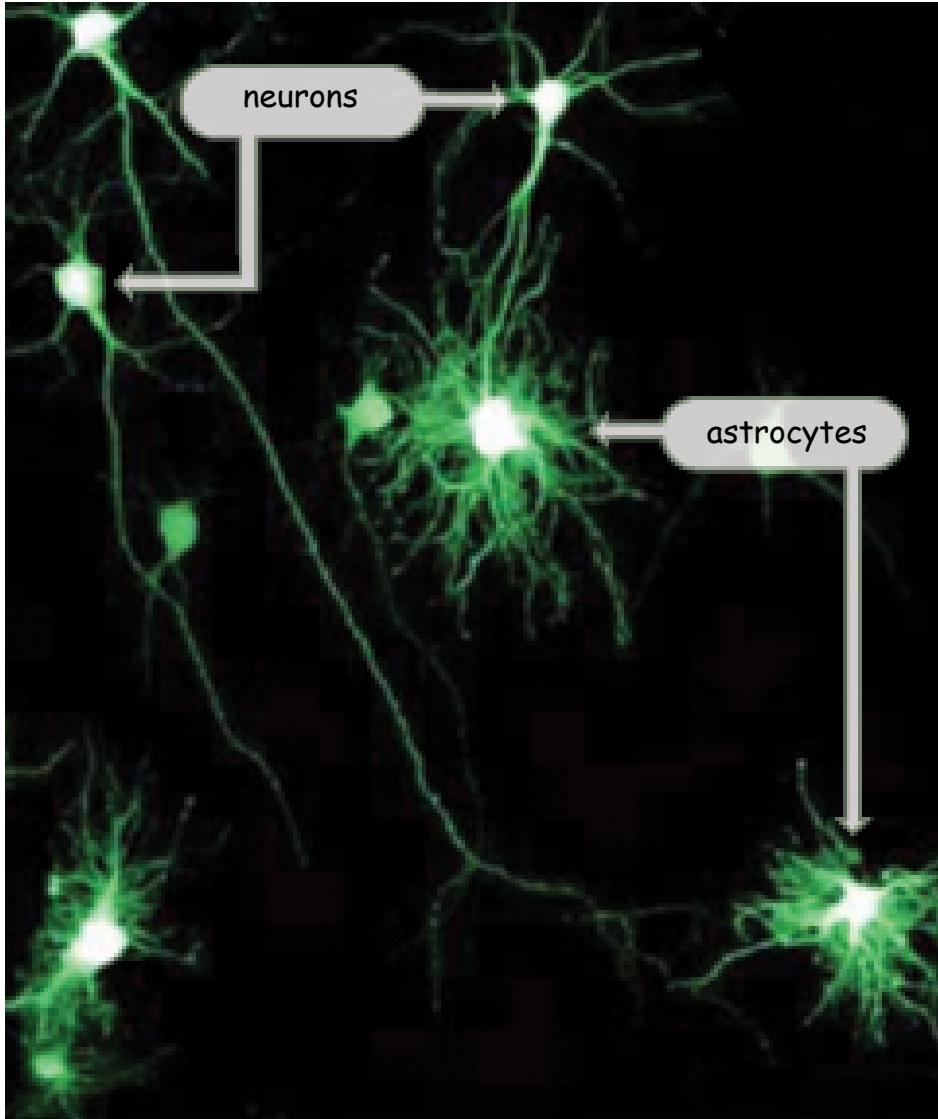
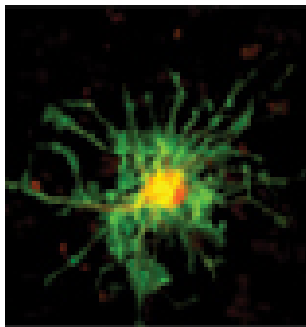
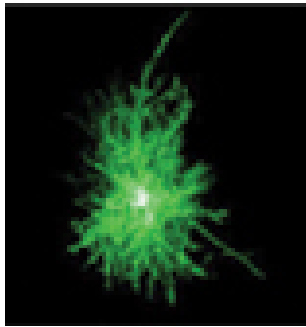
$T = 18.5 \text{ C}; \theta = 1873.3324514717697 \text{ cm / sec}$



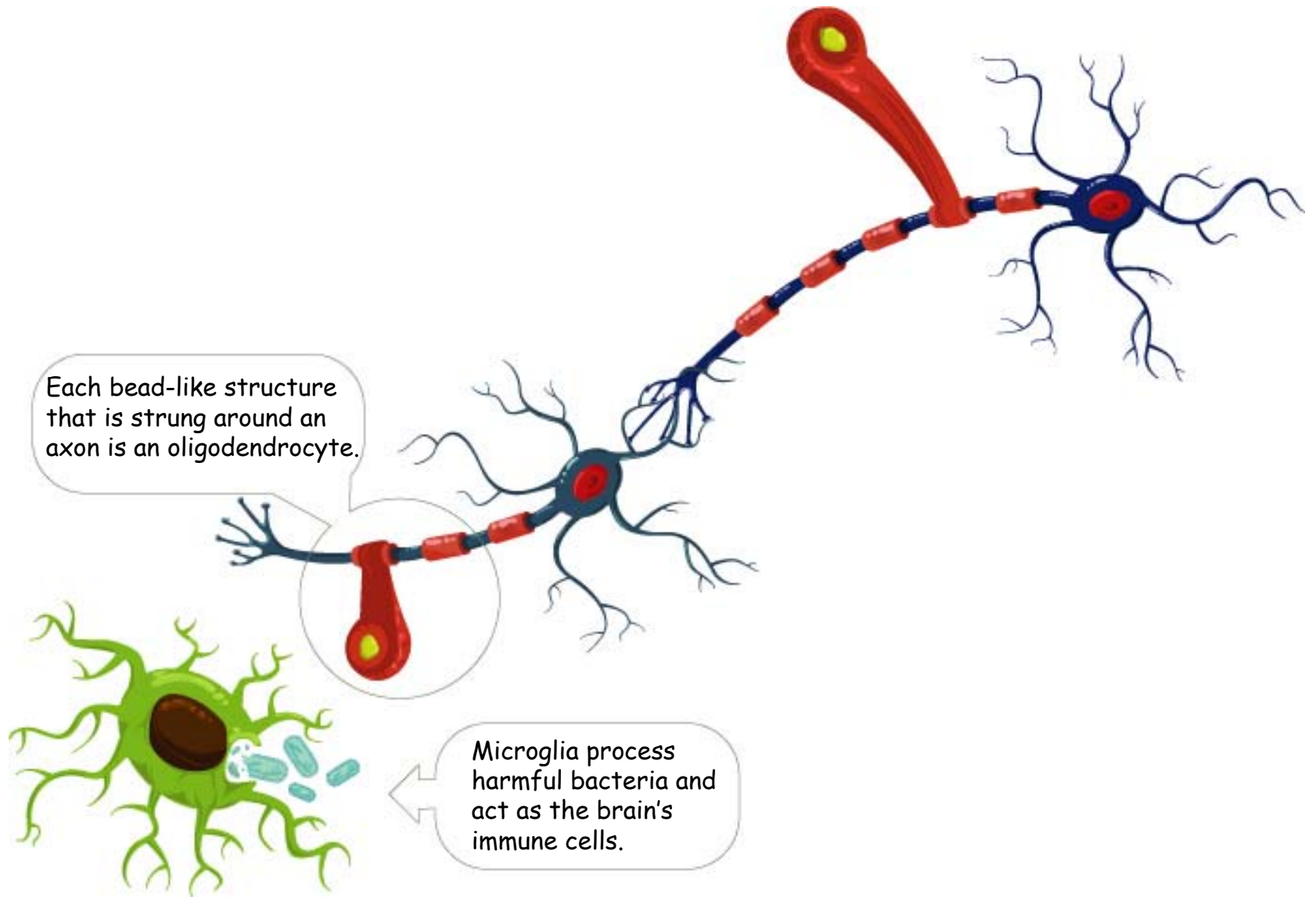
Propagating wave solutions of the Hodgkin-Huxley equations



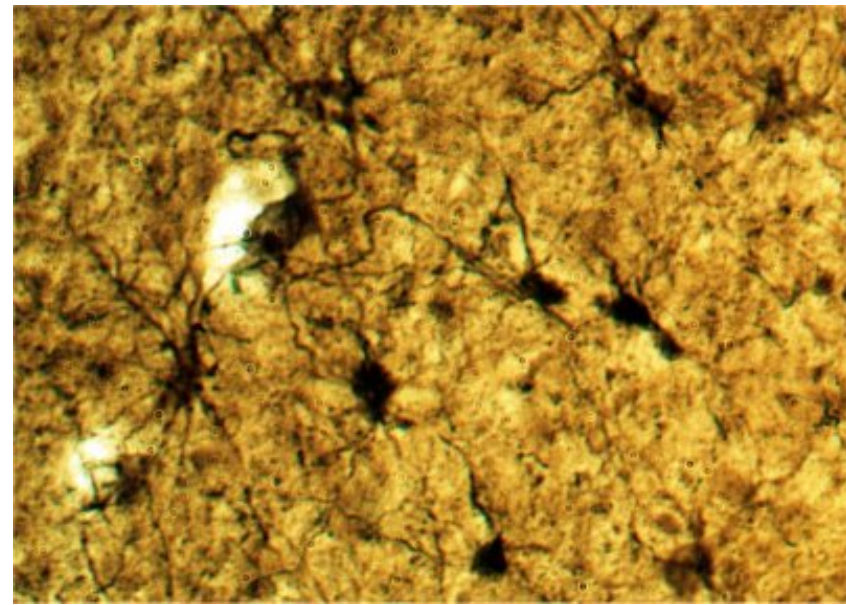
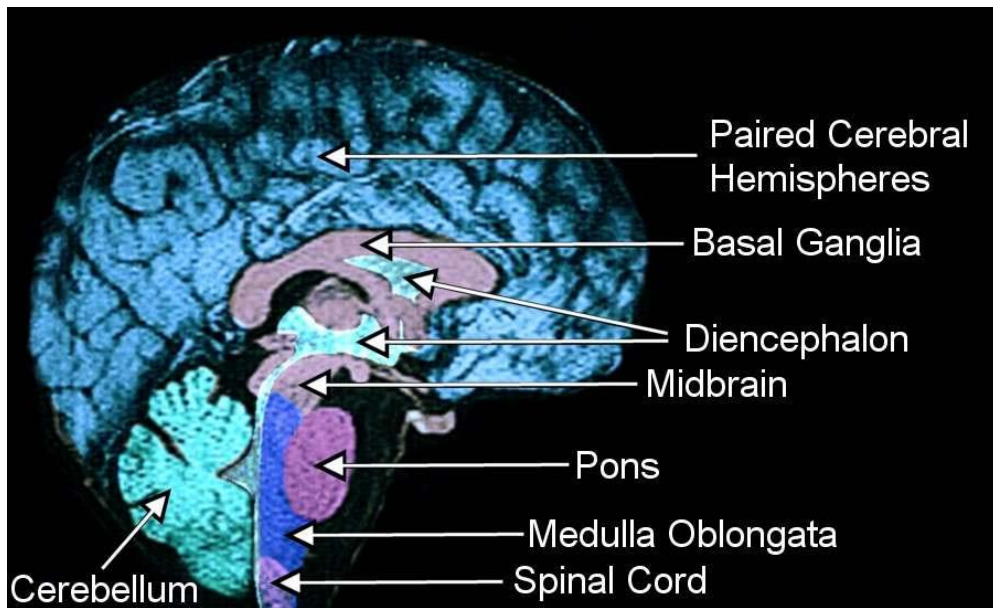
Astrocytes



Photos of neurons and astrocytes



Oligodendrocyte



The human brain

10^{11} neurons connected by $\approx 10^{13}$ to 10^{14} synapses



Computer axial tomography - CAT

Magnetic resonance imaging - MRI

Functional magnetic resonance imaging
- **fMRI**

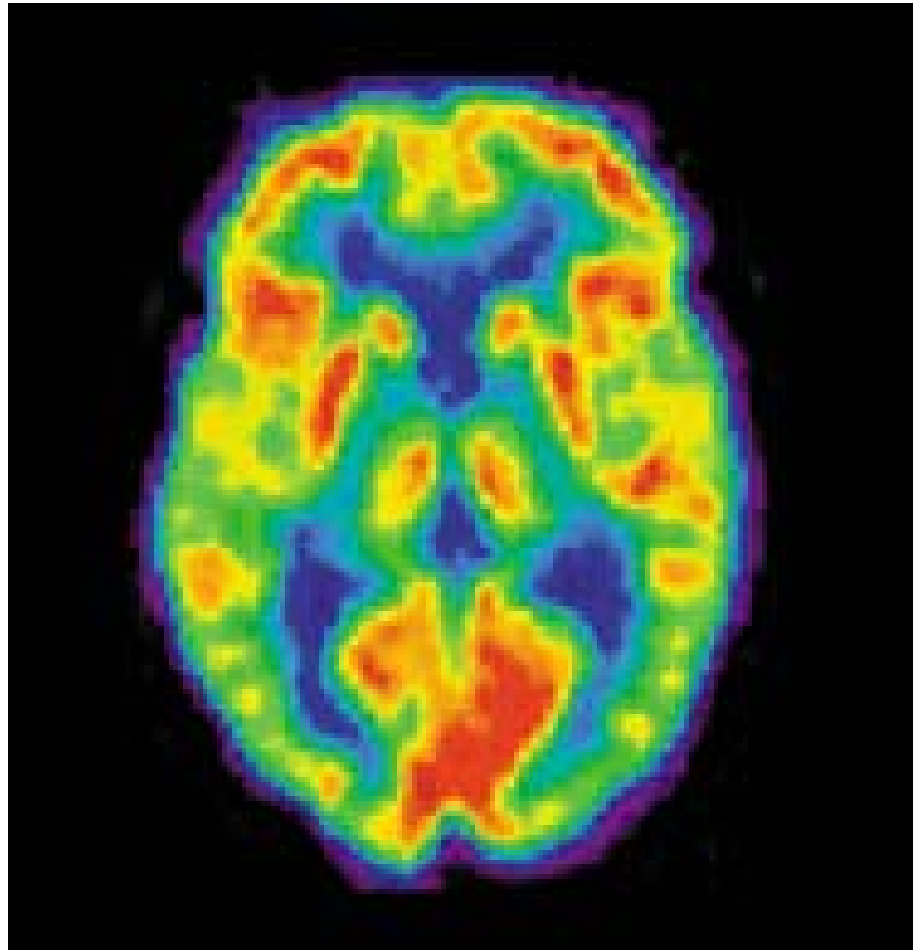
Positron emission tomography - PET

Single photon emission computed
tomography - SPECT

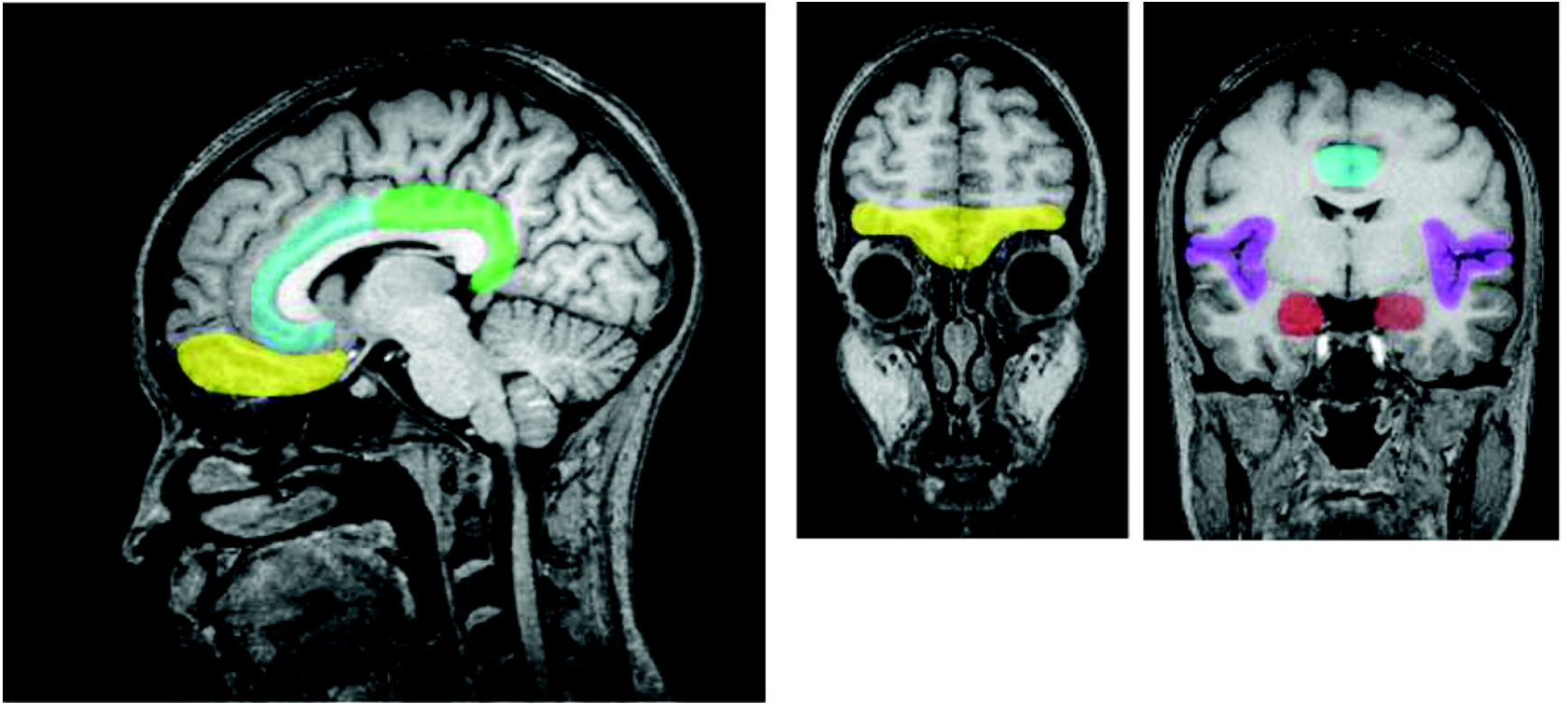
Diffuse Optical Tomography - DOT



Neuroimaging techniques



Positron emission tomography - PET



Brain regions involved in emotional experience: **Amygdala** (linking perception, automatic emotional response and memory), **orbitofrontal cortex**, **insular cortex**, **anterior** and **posterior cingulate cortices**.

Picture taken from *Science* **298**, 1191-1194 (2002)

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