

From Belief to Facts in the Theory of Evolution

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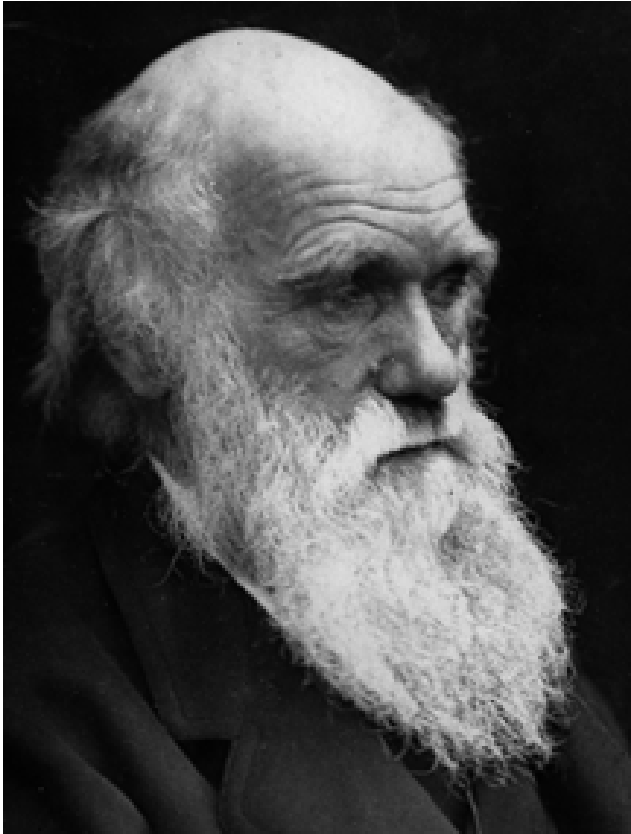
26th International Wittgenstein Symposium

Kirchberg am Wechsel, NÖ, 03.– 09.08.2003

Web-Page for further information:

<http://www.tbi.univie.ac.at/~pks>

- 1. Theory of evolution, science, and religion**
- 2. Genetics and the theory of evolution**
- 3. Evolution experiments in the laboratory**
- 4. Molecular genetics and the tree of life**



Charles Robert Darwin, 1809-1882



Gregor Mendel, 1822-1884,
Abbot of the Augustinian Monastery in Brunn

The two great scholars in nineteenth century biology



The University of Chicago Press:
Chicago, 1979

The publication of the „Origin of Species“ was well prepared by Charles Darwin and his scientific friends. The conclusions of Alfred Russel Wallace drawn from observations made in Brazil (Amazon territory, 1848-1852) and in Indonesia (Malayan Archipelago, 1854-1862) were close to Darwin's thoughts, who did his systematic studies during the voyage around the world on HMS Beagle (1831-1836). Competition with Wallace urged Darwin to publish his comprehensive book.



HMS Beagle



Charles Robert Darwin, 1809-1882

An abstract of an Essay
on the
Origin
&
Species and Varieties
through natural selection
&
Charles Darwin M.A.
Fellow of the Royal Geological Society

London
in 1859

Earlier abstract of the
,Origin of Species'



Alfred Russell Wallace, 1823-1913

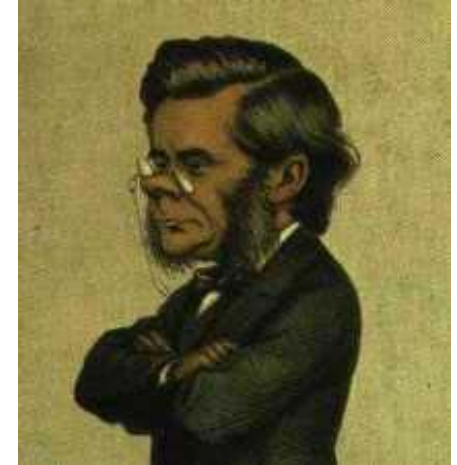
The two competitors in the formulation of evolution by natural selection



Samuel Wilberforce, 1805-1873,
asked Huxley **whether it was through his grandfather or his grandmother that he claimed descent from monkeys.**



British Association for the Advancement of Science: Meeting, Oxford 1860



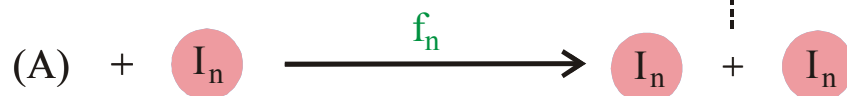
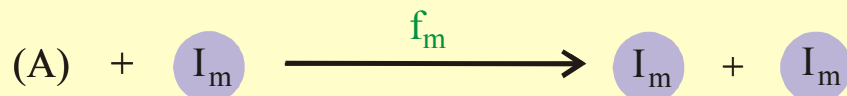
Thomas Henry Huxley, 1825-1895,
replied that if faced with the question, „**would I rather have a miserable ape for a grandfather, or a man highly endowed by nature and possessed of great means and influence, and yet who employs these faculties and that influence to the mere purpose of introducing ridicule into a grave scientific discussion – I unhesitatingly affirm my preference for the ape.**“



Darwin, 1809-1882,
On the Origin of Species by Means of Natural Selection; or the Preservation of Favored Races in the Struggle for Life,
First edition, 24.11.1859,
London: John Murray, Albemarle Street

The Bishop Wilberforce –Huxley debate: Oxford, 30.06.1860

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$$dx_i / dt = f_i x_i - x_i \Phi = x_i (f_i - \Phi)$$

$$\Phi = \sum_j f_j x_j ; \quad \sum_j x_j = 1 ; \quad i, j = 1, 2, \dots, n$$

$$[I_i] = x_i \geq 0 ; \quad i = 1, 2, \dots, n ;$$

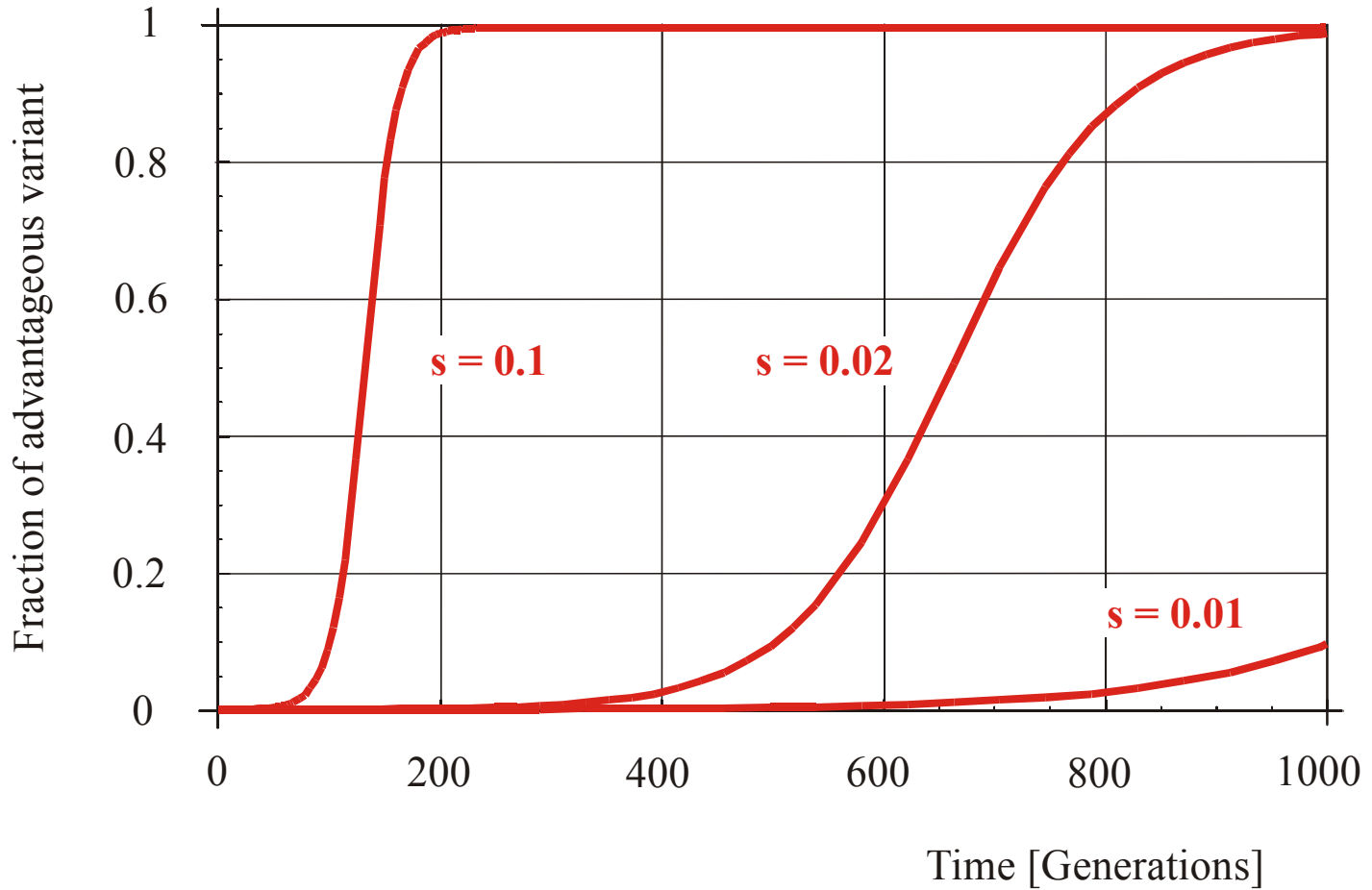
$$[A] = a = \text{constant}$$

$$f_m = \max \{f_j ; j=1, 2, \dots, n\}$$

$$x_m(t) \rightarrow 1 \text{ for } t \rightarrow \infty$$

Reproduction of individuals as basis of selection

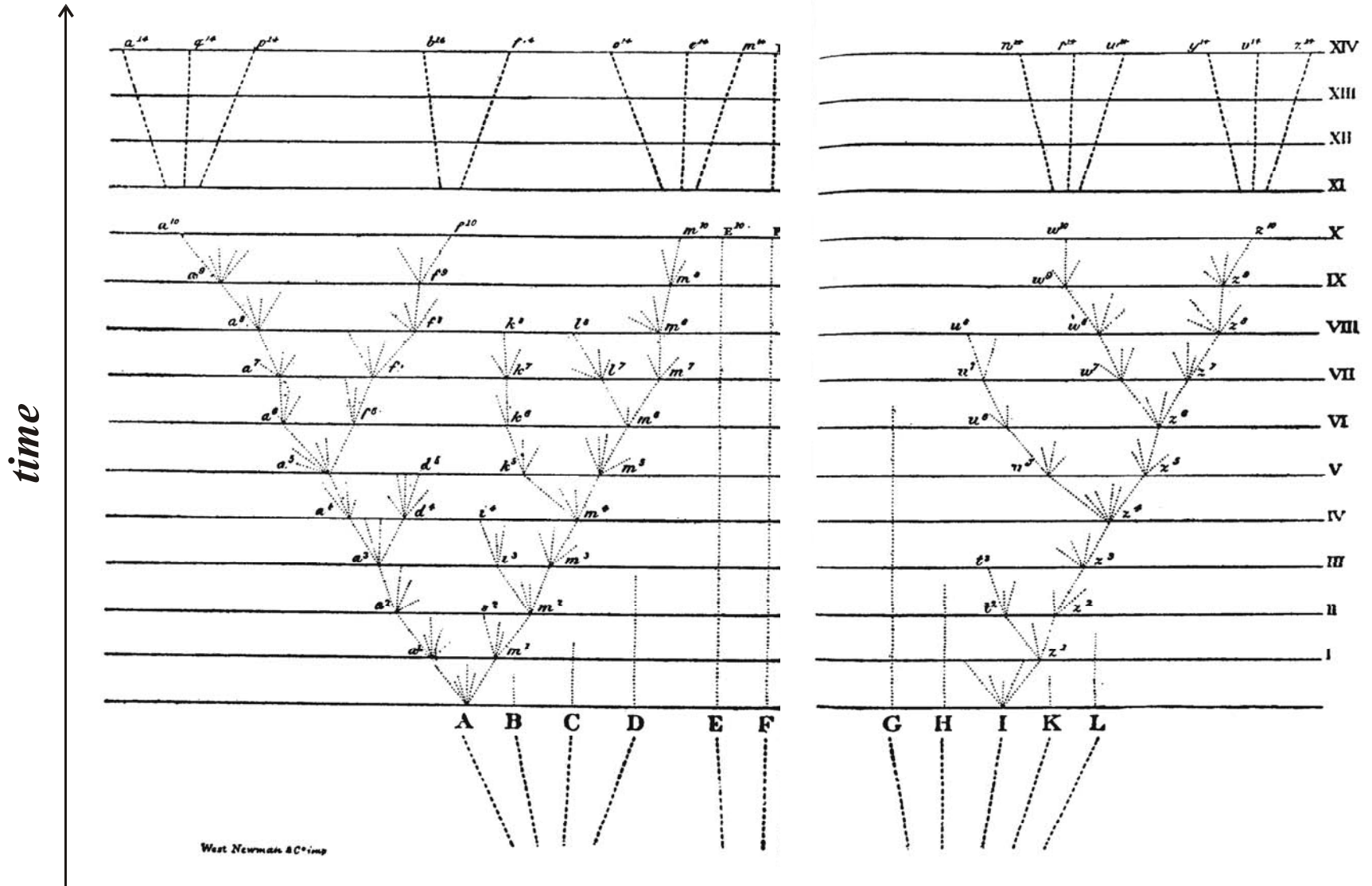
$$s = (f_2 - f_1) / f_1; f_2 > f_1; x_1(0) = 1 - 1/N; x_2(0) = 1/N$$



Selection of advantageous mutants in populations of $N = 10\ 000$ individuals

Key ingredients in Darwin's theory of evolution are:

- (i) Variations** occurring spontaneously and not themselves produced by the environment,
- (ii) Competition** for resources, so that only the best adapted survive to reproduce, and, therefore
- (iii) Selection** by the environment, of which variants will survive and increase in number.



Charles Darwin, *The Origin of Species*, 6th edition.
 Everyman's Library, Vol.811, Dent London, pp.121-122.



Etienne Geoffroy Saint-Hillaire, 1772-1844



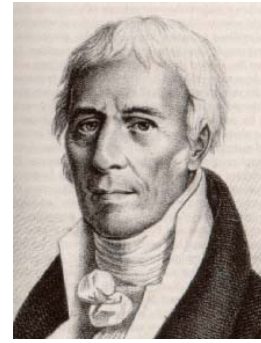
Charles Robert Darwin, 1809-1882



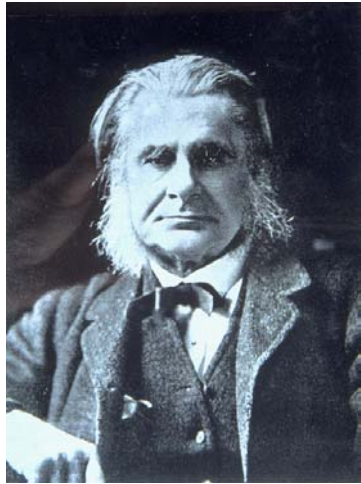
Erasmus Darwin, 1731-1802



Alfred Russel Wallace, 1823-1913



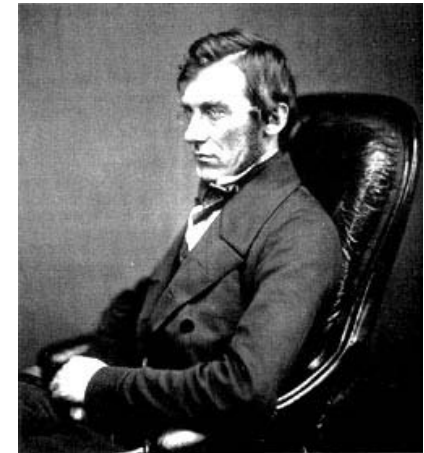
Jean Baptiste Pierre Antoine de Monet, Chevalier de Lamarck, 1744-1829



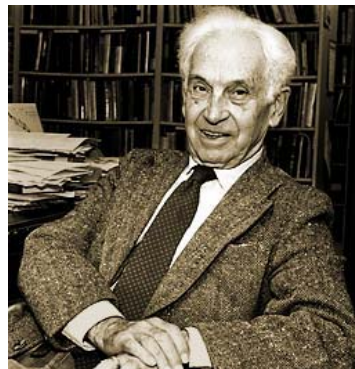
Thomas Henry Huxley, 1825-1895



HMS Beagle, 1831-1836



Joseph Dalton Hooker, 1817-1911



Ernst Mayr, 1904 -

The 'Evolutionists'



John Frederick William Herschel, 1792-1871



Sir Charles Lyell, 1797-1875

„Borderline Evolutionists“ (Michael Ruse, *The Darwinian Revolution*, 1979)



Georges Cuvier, 1769-1832



Bishop Samuel Wilberforce, 1805-1873



Richard Owen, 1804-1892



Jean Louis Rodolphe Agassiz, 1807-1873



Adam Sedgwick, 1785-1873



William Whewell, 1794-1866

The 'Anti-Evolutionists'

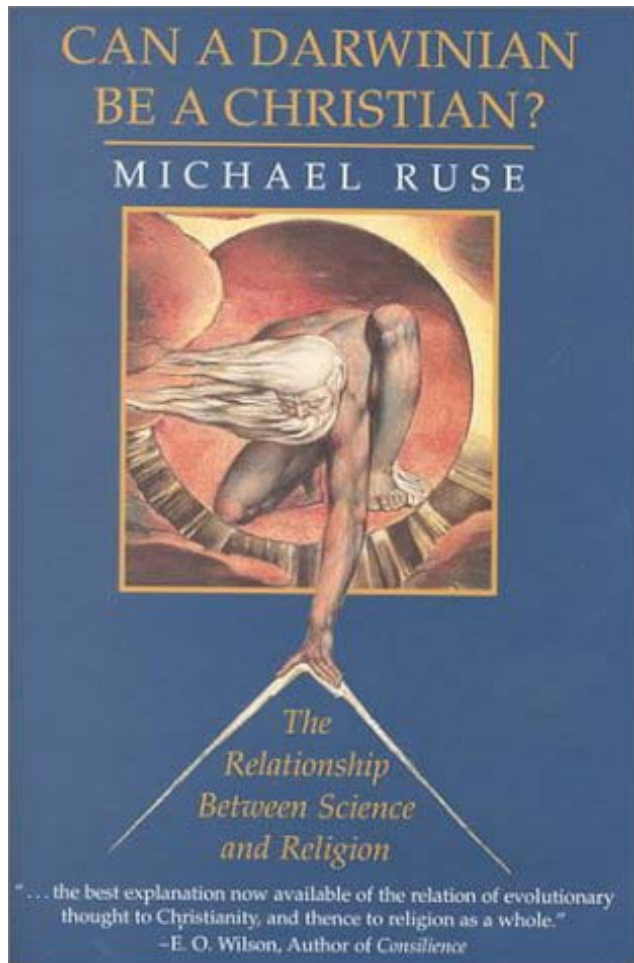
Science – Religion debate in the 21st Century:

Science

Richard Dawkins,
Stephen J. Gould,
E.O. Wilson

Religion

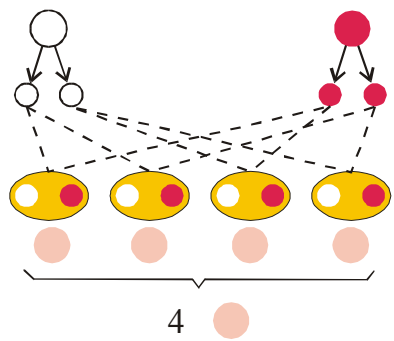
Arthur Peacocke,
Robert J. Russell,
Keith Ward



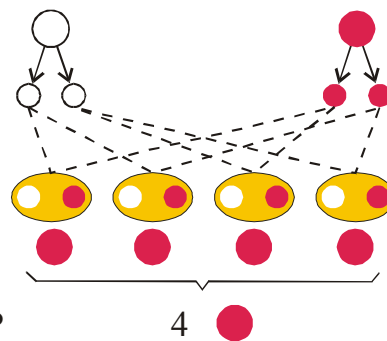
Cambridge University Press,
New York 2002

Michael Ruse argues, that although it is at times difficult for a Darwinian to embrace Christian belief, it is by no means inconceivable. At the same he suggests ways in which a Christian believer should have no difficulty accepting evolution in general, and Darwinism in particular.

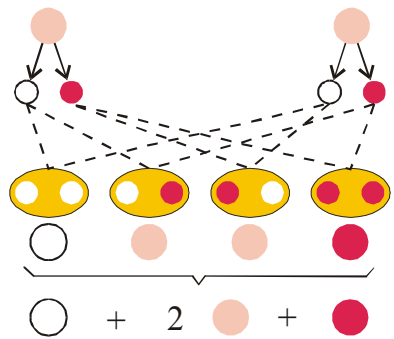
1. Theory of evolution, science, and religion
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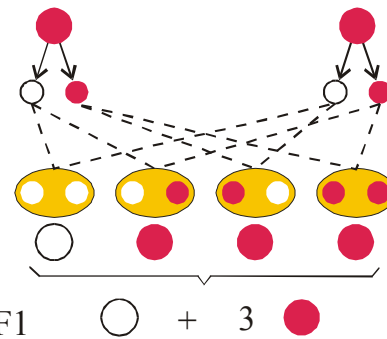
P



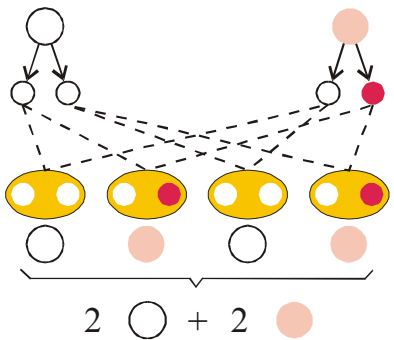
$F1 = P \times P$



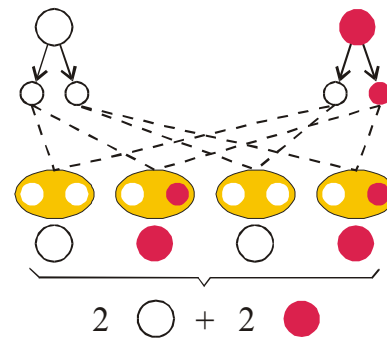
F1



$F2 = F1 \times F1$

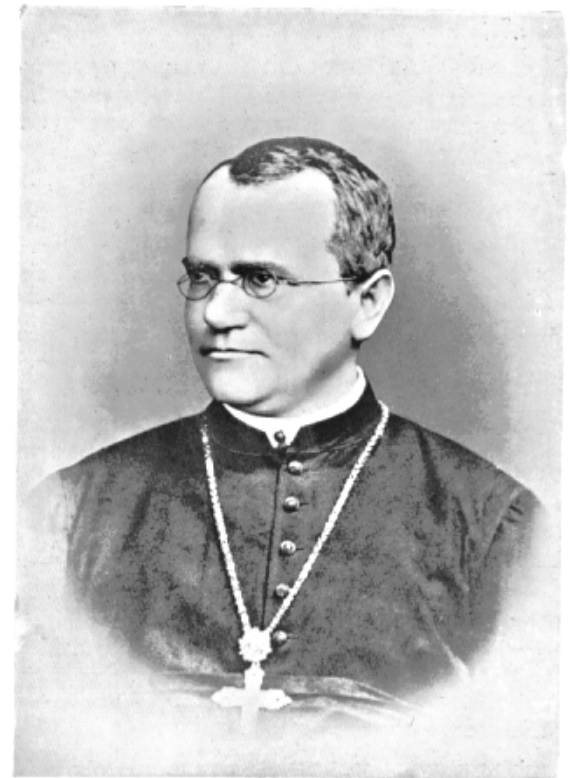


$P \times F1$



Intermediate pair of alleles

Dominant/recessive pair of alleles



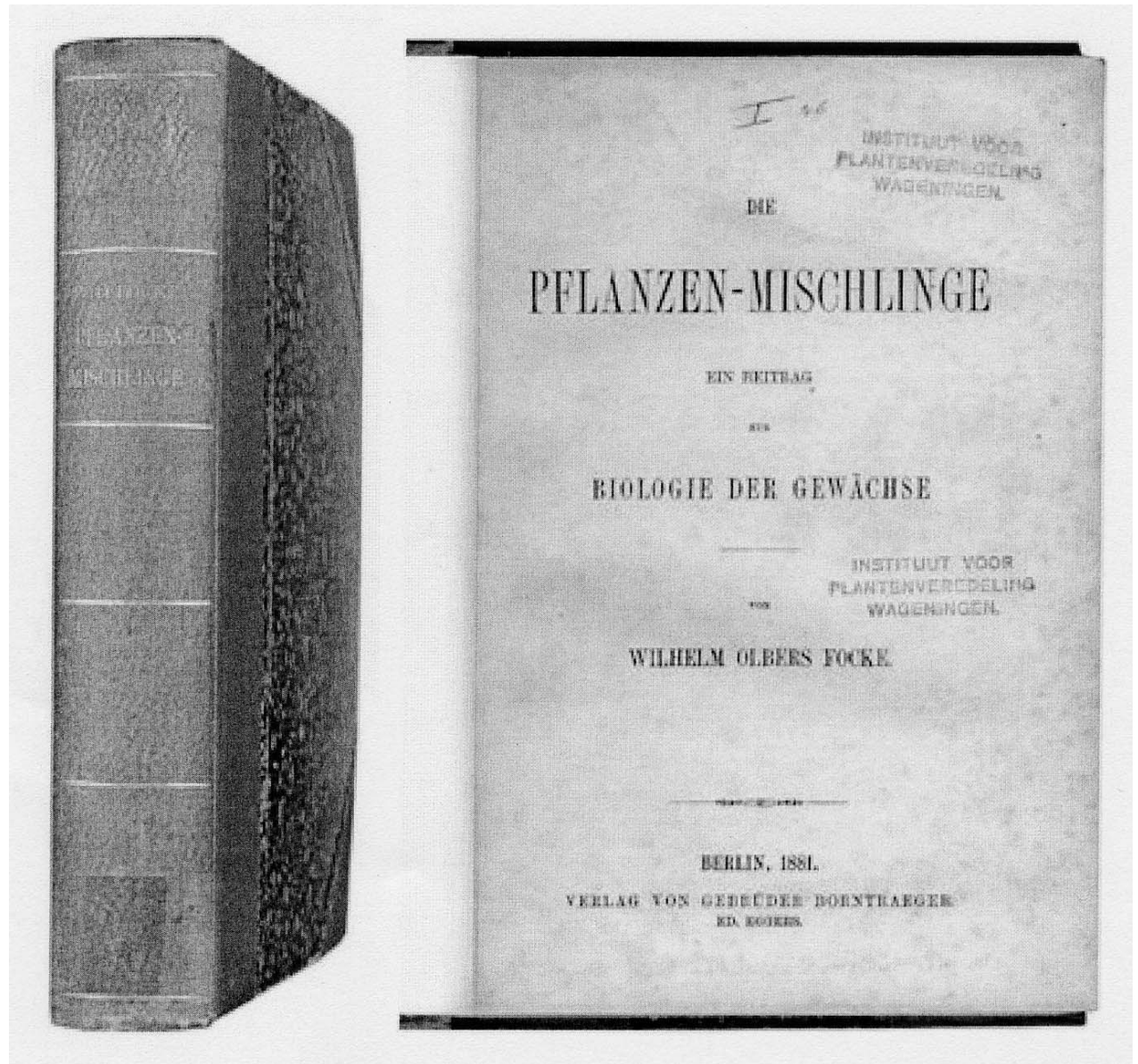
Gregor Mendel's laws of inheritance:

Versuche über Pflanzen-Hybriden.

Verhandlungen des naturforschenden Vereins in Brunn, 4: 3-47 (1865)

Presented at the Meetings of 08.02. and 08.03.1865

Mendel's work cited 1881
in W.O. Focke's
„Die Pflanzen-Mischlinge“





Sir Ronald Aylmer Fisher, 1890-1962



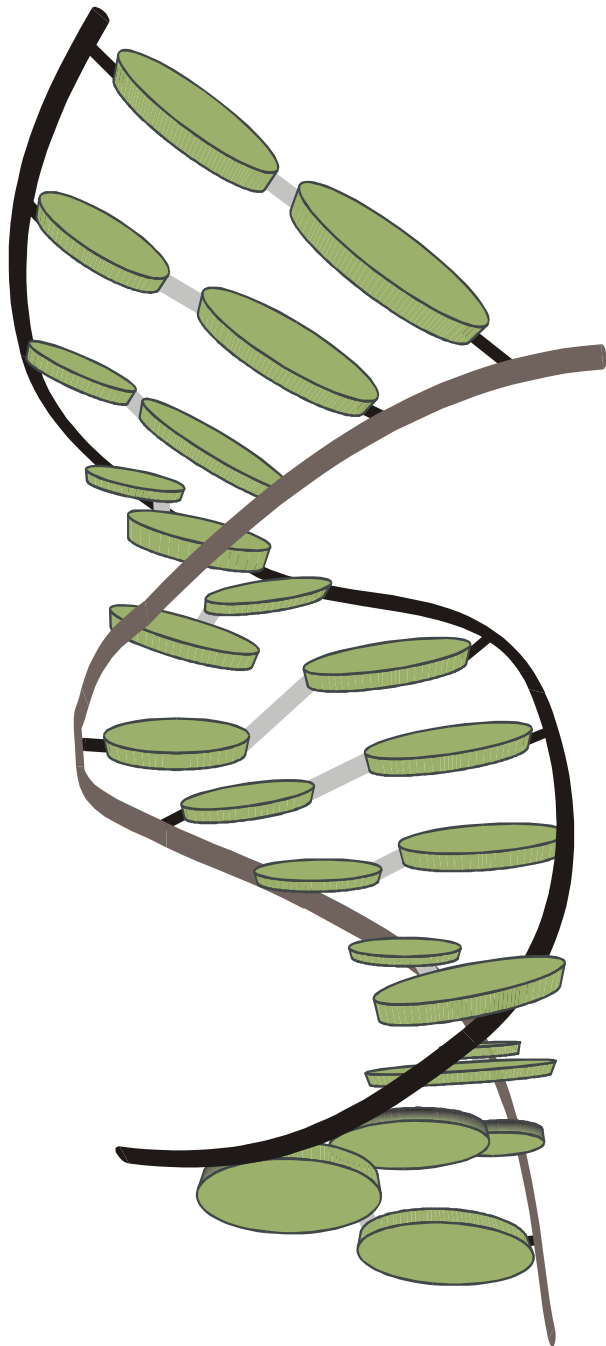
John Burdon Sanderson Haldane, 1892-1964



Sewall Wright, 1889-1988

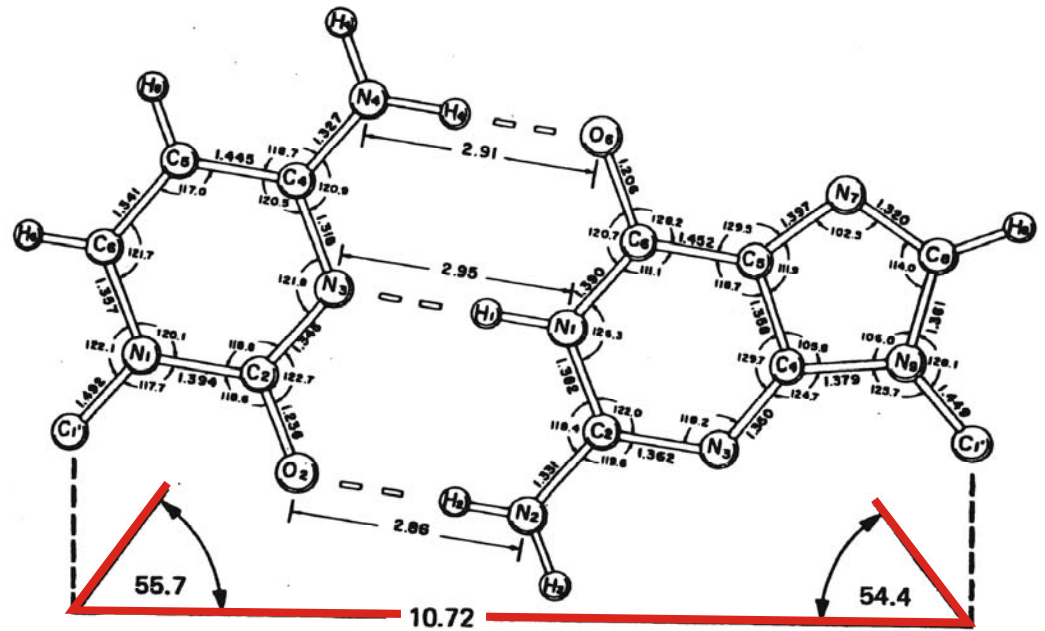
The three scholars of theoretical population biology

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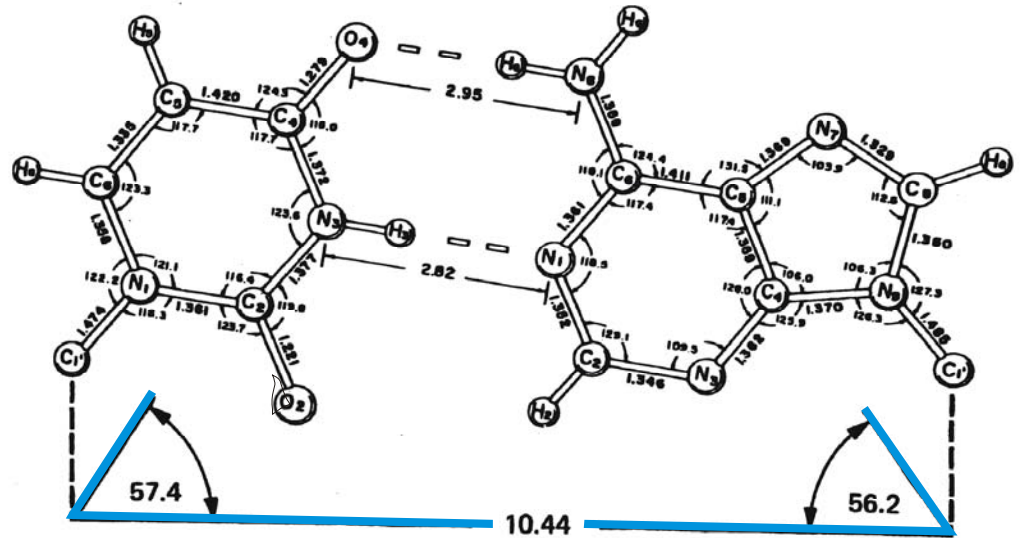
James D. Watson, 1928- , and Francis Crick, 1916- ,
Nobel Prize 1962

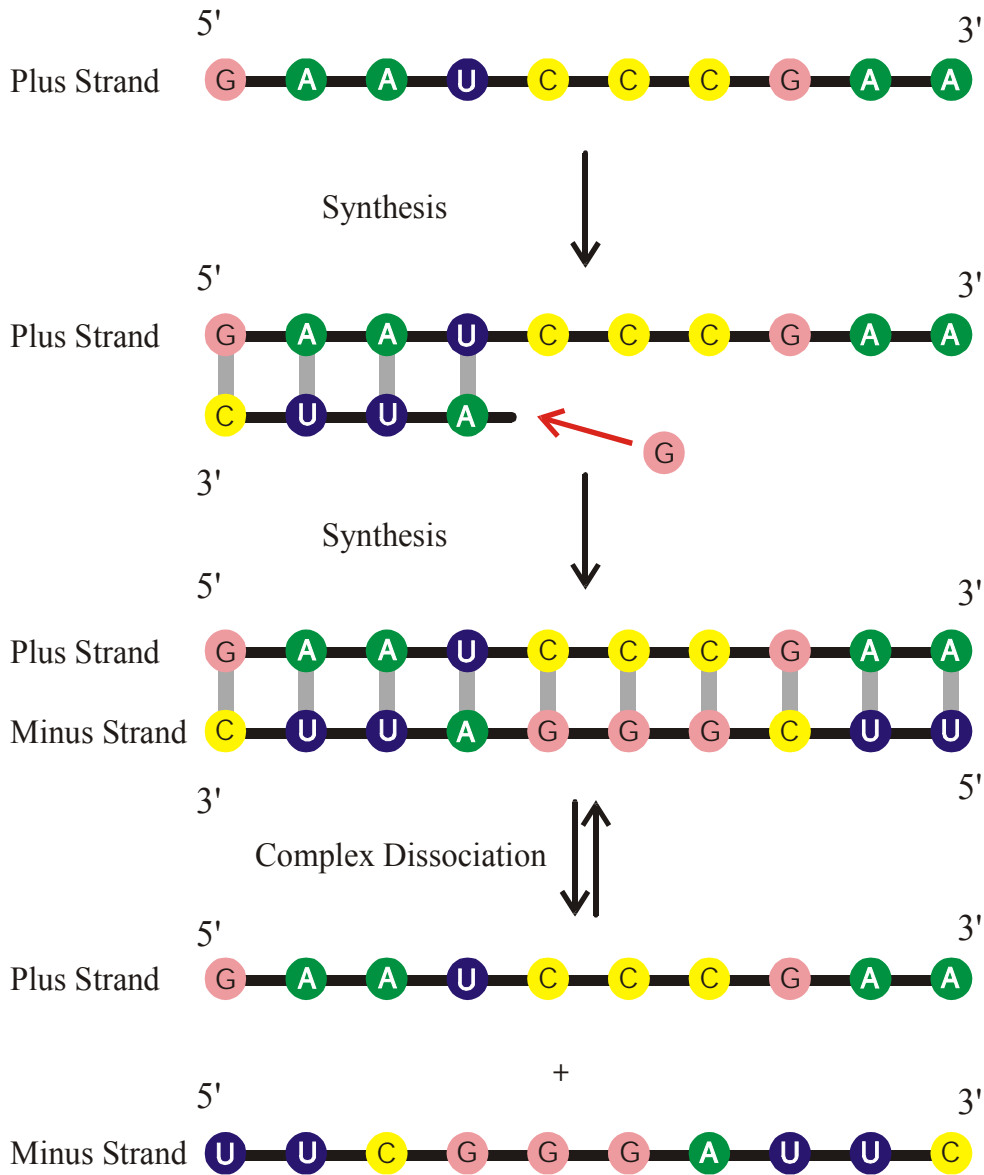
The three-dimensional structure of a
short double helical stack of B-DNA



Canonical Watson-Crick
base pairs:

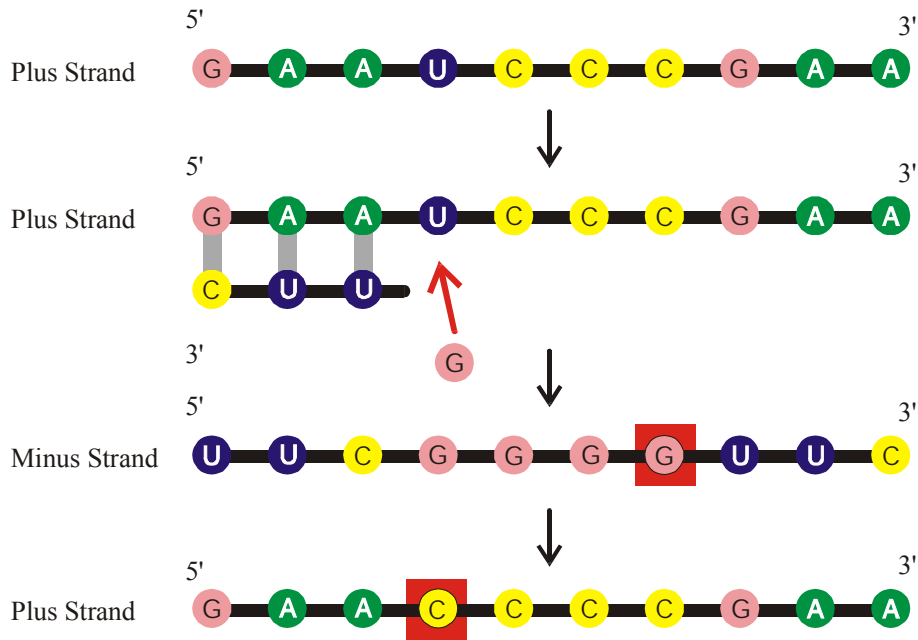
cytosine – guanine
uracil – adenine





Complementary replication as the simplest copying mechanism of RNA
 Complementary is determined by Watson-Crick base pairs:

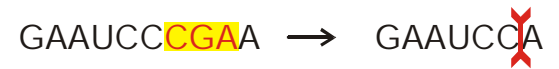




Point Mutation



Insertion

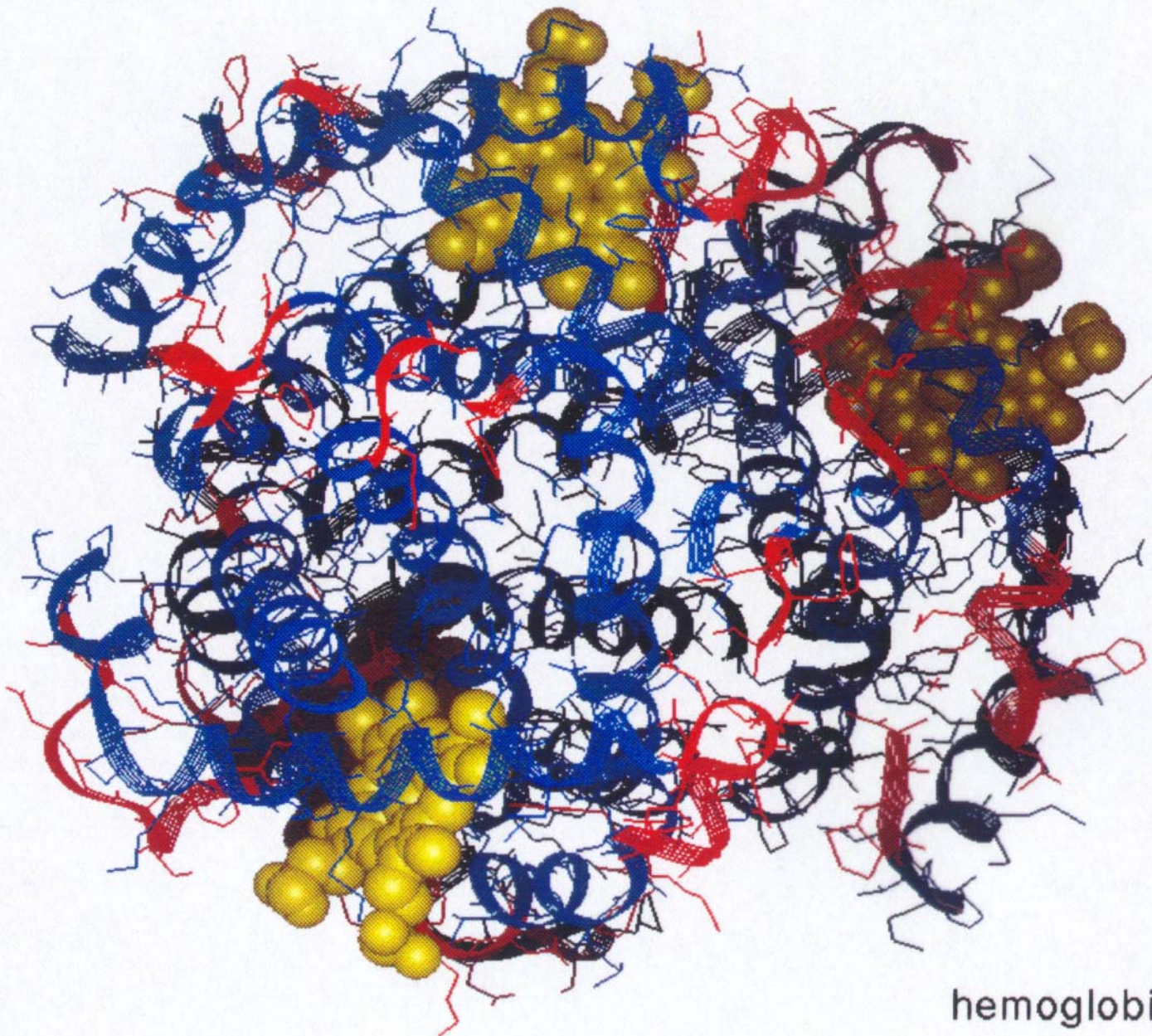


Deletion

Mutations in nucleic acids represent the mechanism of **variation** of **genotypes**.



Max Perutz, 1914-2002, at the opening
of the Max Perutz-Library, Vienna
BioCenter, in 1994
Nobel Prize 1962



hemoglobin

Evolution of RNA molecules based on Q β phage

D.R.Mills, R.L.Peterson, S.Spiegelman, *An extracellular Darwinian experiment with a self-duplicating nucleic acid molecule*. Proc.Natl.Acad.Sci.USA **58** (1967), 217-224

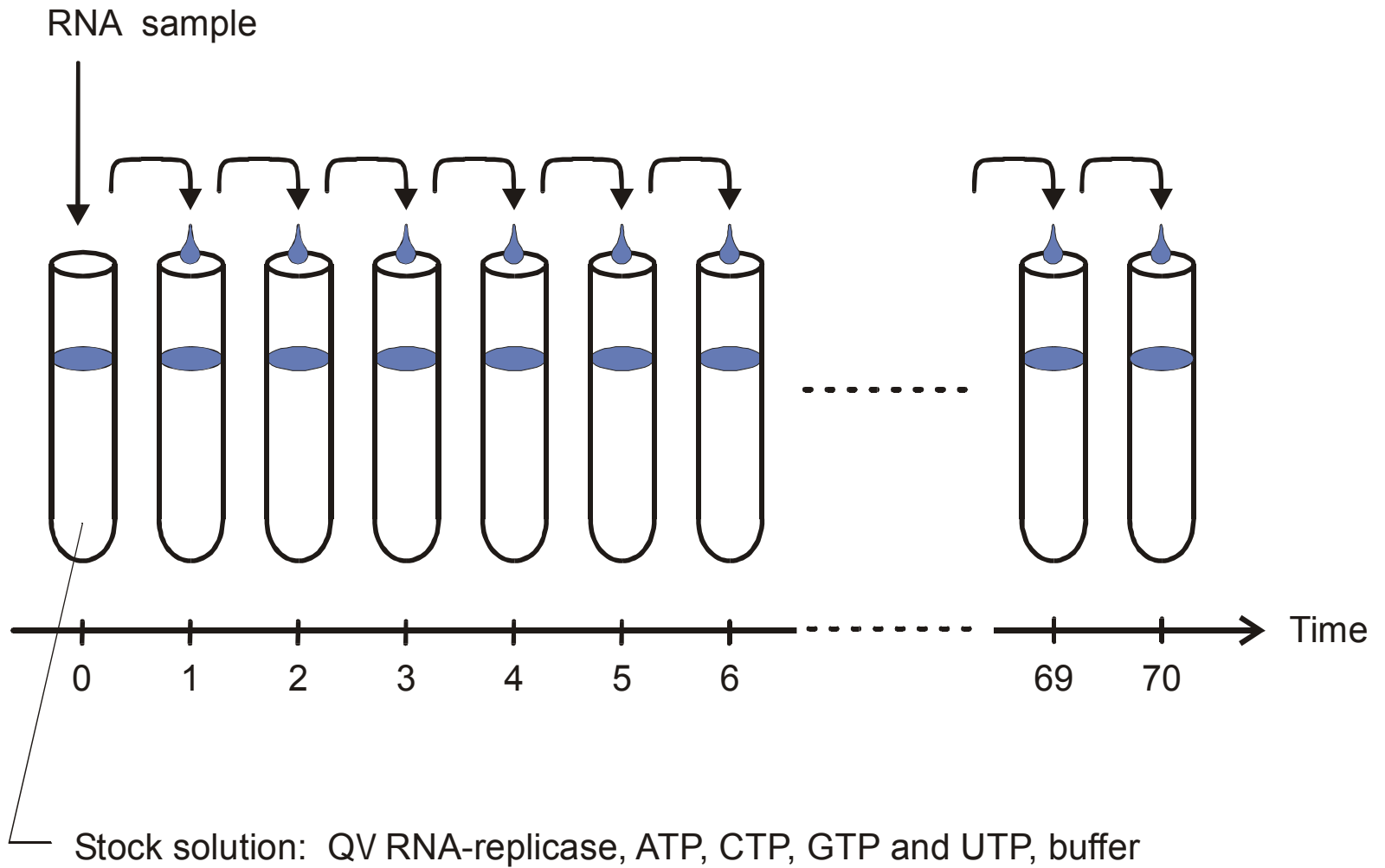
S.Spiegelman, *An approach to the experimental analysis of precellular evolution*. Quart.Rev.Biophys. **4** (1971), 213-253

C.K.Biebricher, *Darwinian selection of self-replicating RNA molecules*. Evolutionary Biology **16** (1983), 1-52

G.Bauer, H.Otten, J.S.McCaskill, *Travelling waves of in vitro evolving RNA*. Proc.Natl.Acad.Sci.USA **86** (1989), 7937-7941

C.K.Biebricher, W.C.Gardiner, *Molecular evolution of RNA in vitro*. Biophysical Chemistry **66** (1997), 179-192

G.Strunk, T.Ederhof, *Machines for automated evolution experiments in vitro based on the serial transfer concept*. Biophysical Chemistry **66** (1997), 193-202



The serial transfer technique applied to RNA evolution *in vitro*

Reproduction of the original figure of the serial transfer experiment with Q β RNA

D.R.Mills, R.L.Peterson, S.Spiegelman,
*An extracellular Darwinian experiment
 with a self-duplicating nucleic acid
 molecule.* Proc.Natl.Acad.Sci.USA
58 (1967), 217-224

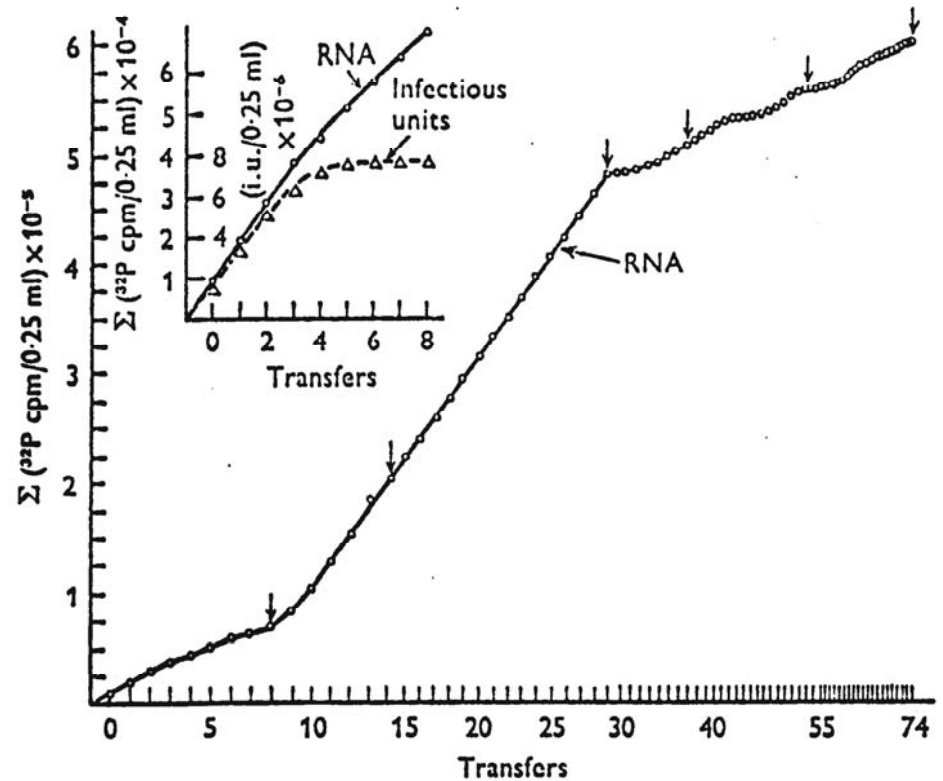
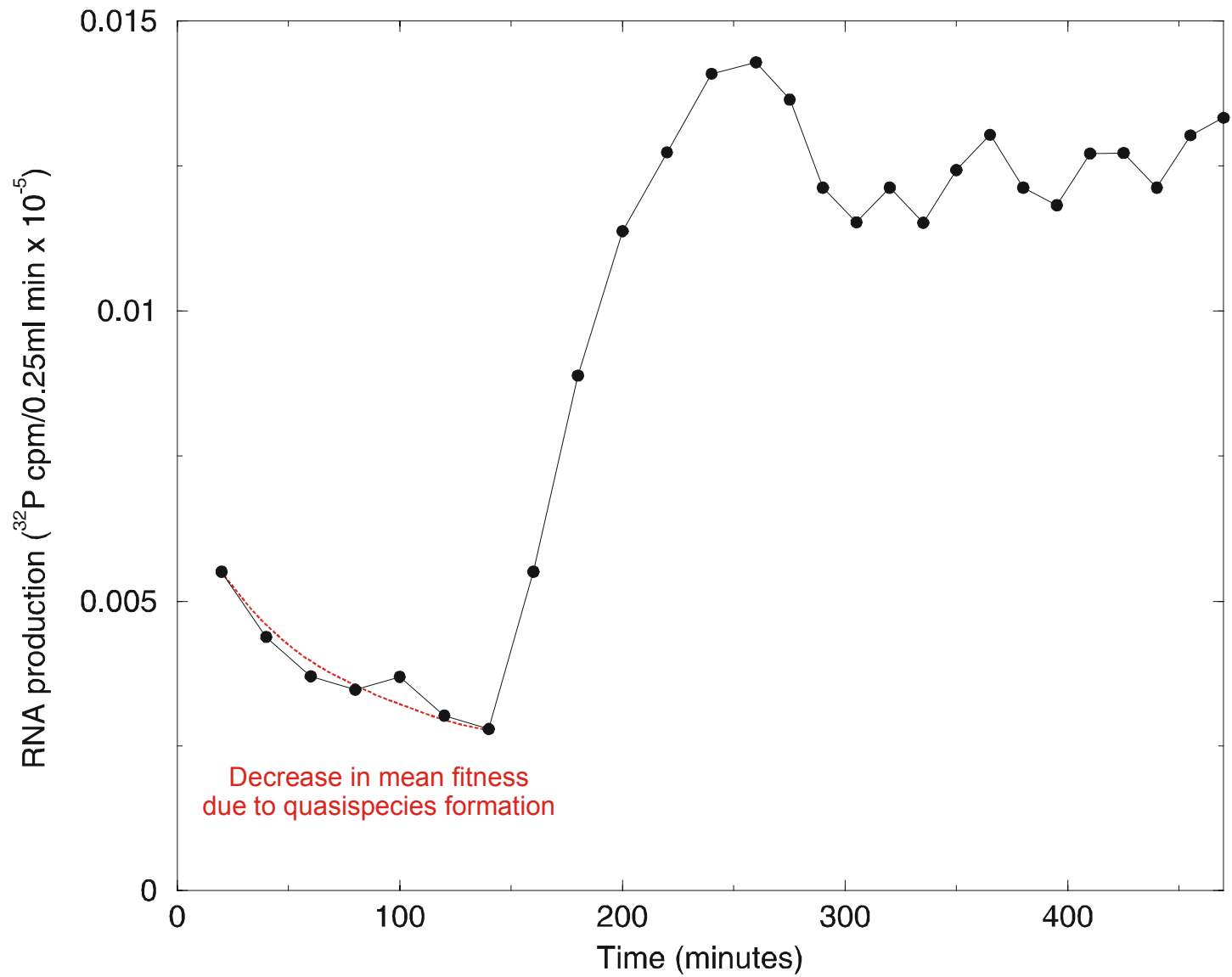


Fig. 9. Serial transfer experiment. Each 0.25 ml standard reaction mixture contained 40 μg of Q β replicase and ^{32}P -UTP. The first reaction (0 transfer) was initiated by the addition of 0.2 μg ts-1 (temperature-sensitive RNA) and incubated at 35 $^{\circ}\text{C}$ for 20 min, whereupon 0.02 ml was drawn for counting and 0.02 ml was used to prime the second reaction (first transfer), and so on. After the first 13 reactions, the incubation periods were reduced to 15 min (transfers 14-29). Transfers 30-38 were incubated for 10 min. Transfers 39-52 were incubated for 7 min, and transfers 53-74 were incubated for 5 min. The arrows above certain transfers (0, 8, 14, 29, 37, 53, and 73) indicate where 0.001-0.1 ml of product was removed and used to prime reactions for sedimentation analysis on sucrose. The inset examines both infectious and total RNA. The results show that biologically competent RNA ceases to appear after the 4th transfer (Mills *et al.* 1967).



The increase in RNA production rate during a serial transfer experiment

Bacterial Evolution

S. F. Elena, V. S. Cooper, R. E. Lenski. *Punctuated evolution caused by selection of rare beneficial mutants*. Science **272** (1996), 1802-1804

D. Papadopoulos, D. Schneider, J. Meier-Eiss, W. Arber, R. E. Lenski, M. Blot. *Genomic evolution during a 10,000-generation experiment with bacteria*. Proc.Natl.Acad.Sci.USA **96** (1999), 3807-3812

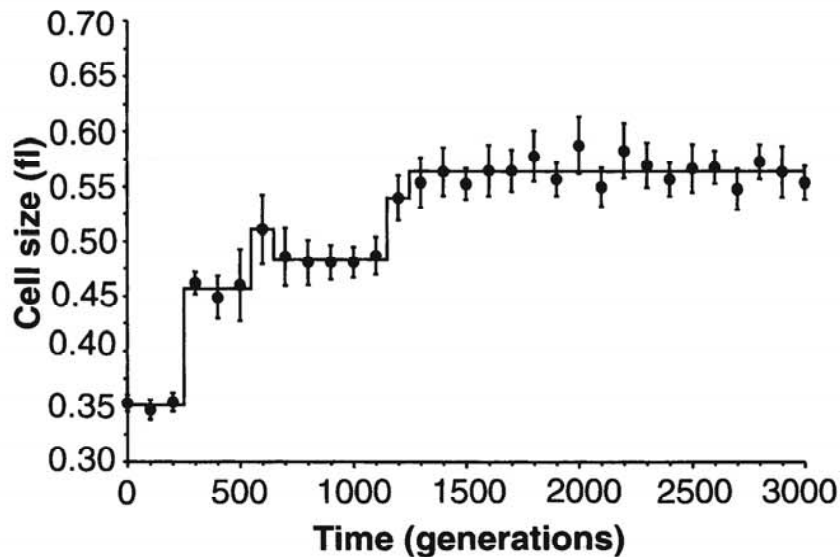


Fig. 1. Change in average cell size (1 fl = 10^{-15} L) in a population of *E. coli* during 3000 generations of experimental evolution. Each point is the mean of 10 replicate assays (22). Error bars indicate 95% confidence intervals. The solid line shows the best fit of a step-function model to these data (Table 1).

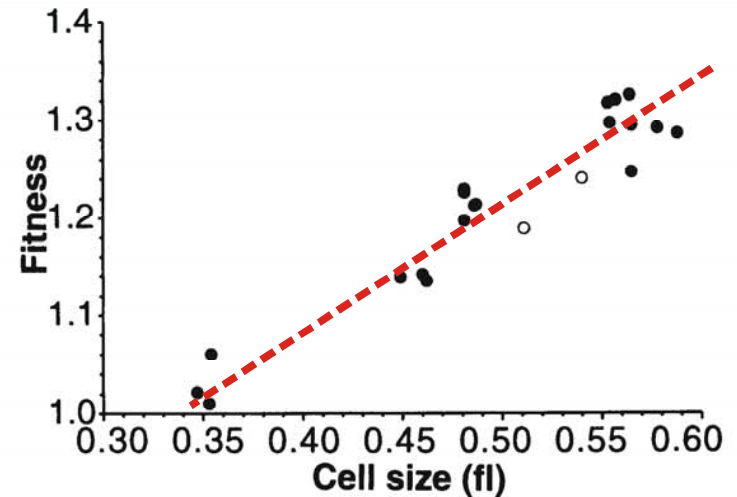


Fig. 2. Correlation between average cell size and mean fitness, each measured at 100-generation intervals for 2000 generations. Fitness is expressed relative to the ancestral genotype and was obtained from competition experiments between derived and ancestral cells (6, 7). The open symbols indicate the only two samples assigned to different steps by the cell size and fitness data.

Epochal evolution of bacteria in serial transfer experiments under constant conditions

S. F. Elena, V. S. Cooper, R. E. Lenski. *Punctuated evolution caused by selection of rare beneficial mutants.* Science **272** (1996), 1802-1804

Evolutionary design of RNA molecules

D.B.Bartel, J.W.Szostak, *In vitro selection of RNA molecules that bind specific ligands.* Nature **346** (1990), 818-822

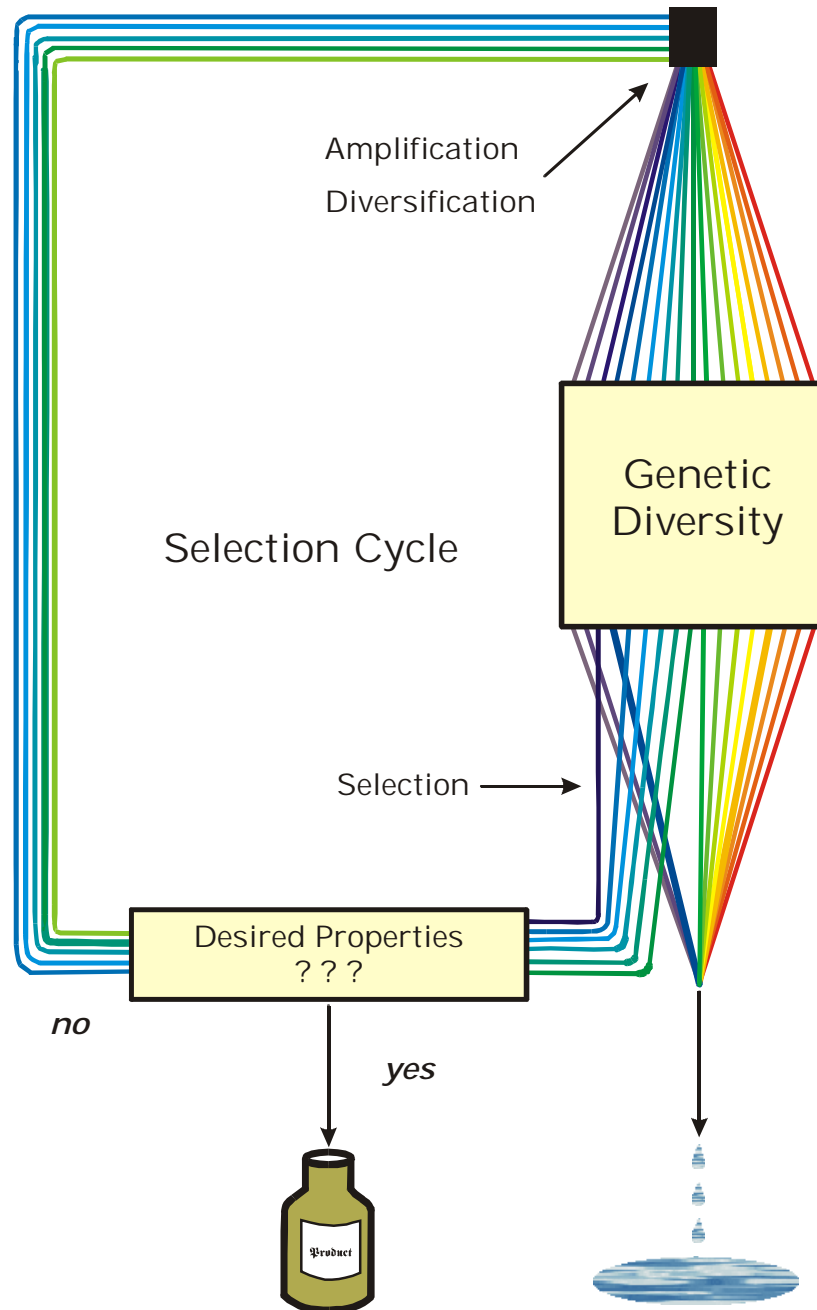
C.Tuerk, L.Gold, *SELEX - Systematic evolution of ligands by exponential enrichment: RNA ligands to bacteriophage T4 DNA polymerase.* Science **249** (1990), 505-510

D.P.Bartel, J.W.Szostak, *Isolation of new ribozymes from a large pool of random sequences.* Science **261** (1993), 1411-1418

R.D.Jenison, S.C.Gill, A.Pardi, B.Poliski, *High-resolution molecular discrimination by RNA.* Science **263** (1994), 1425-1429

Y.Wang, R.R.Rando, *Specific binding of aminoglycoside antibiotics to RNA.* Chemistry & Biology **2** (1995), 281-290

L.Jiang, A.K.Suri, R.Fiala, D.J.Patel, *Saccharide-RNA recognition in an aminoglycoside antibiotic-RNA aptamer complex.* Chemistry & Biology **4** (1997), 35-50

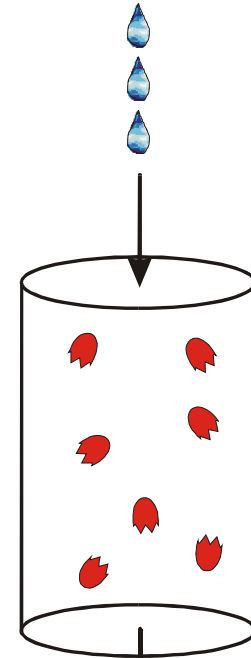
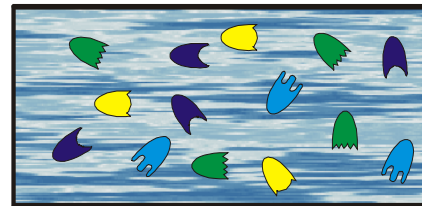
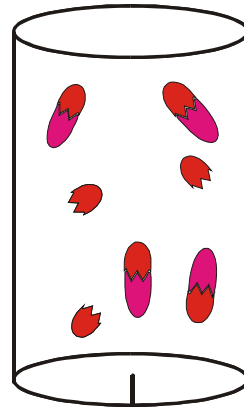
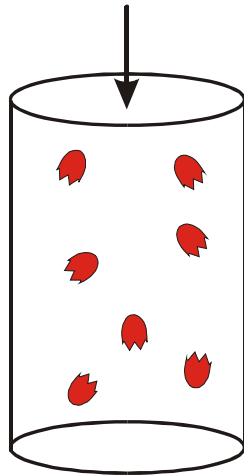
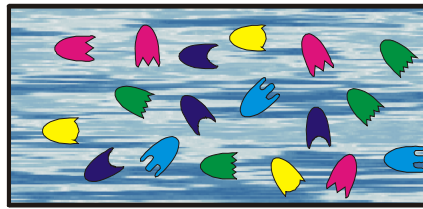


Selection cycle used in applied molecular evolution to design molecules with predefined properties

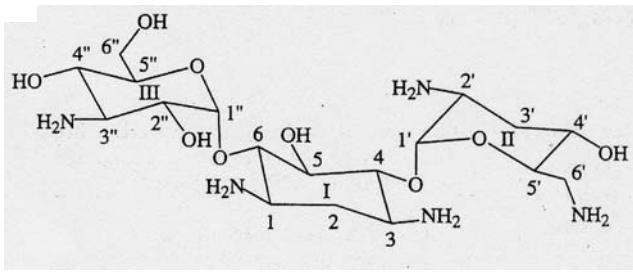
Retention of binders

Elution of binders

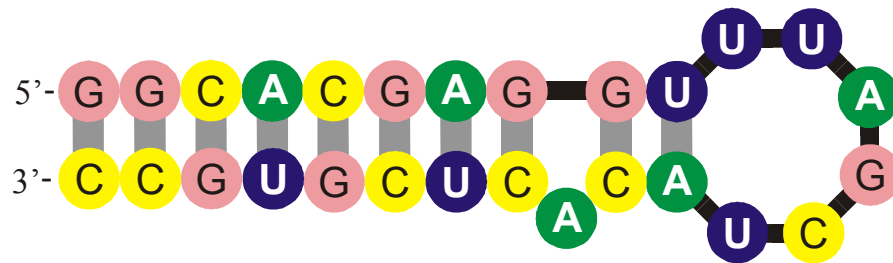
Chromatographic column



The SELEX technique for the evolutionary design of *aptamers*



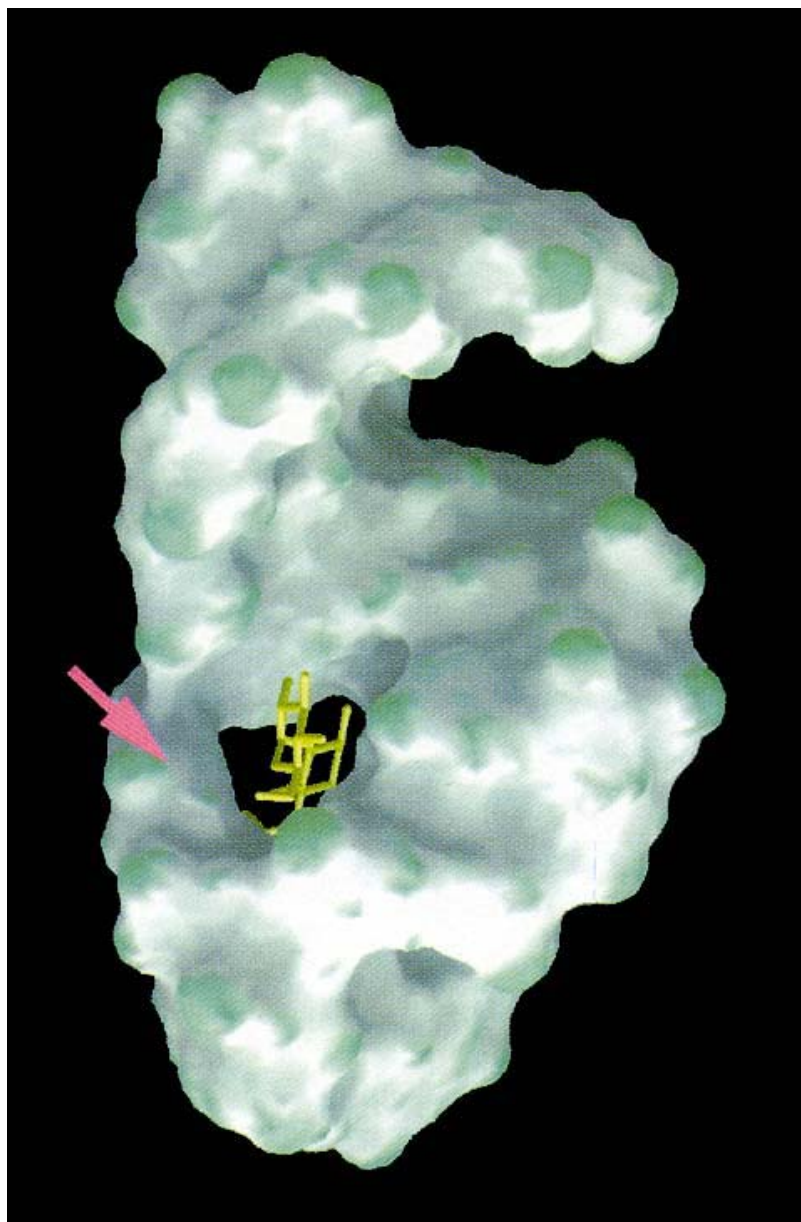
tobramycin



RNA aptamer

Formation of secondary structure of the tobramycin binding RNA aptamer

L. Jiang, A. K. Suri, R. Fiala, D. J. Patel, *Saccharide-RNA recognition in an aminoglycoside antibiotic-RNA aptamer complex*. *Chemistry & Biology* 4:35-50 (1997)



The three-dimensional structure of the tobramycin aptamer complex

L. Jiang, A. K. Suri, R. Fiala, D. J. Patel,
Chemistry & Biology 4:35-50 (1997)

*No new principle will declare itself
from below a heap of facts.*

Sir Peter Medawar, 1985

Theory of molecular evolution

M.Eigen, *Self-organization of matter and the evolution of biological macromolecules*.

Naturwissenschaften **58** (1971), 465-526

C.J. Thompson, J.L. McBride, *On Eigen's theory of the self-organization of matter and the evolution of biological macromolecules*. Math. Biosci. **21** (1974), 127-142

B.L. Jones, R.H. Enns, S.S. Rangnekar, *On the theory of selection of coupled macromolecular systems*. Bull.Math.Biol. **38** (1976), 15-28

M.Eigen, P.Schuster, *The hypercycle. A principle of natural self-organization. Part A: Emergence of the hypercycle*. Naturwissenschaften **58** (1977), 465-526

M.Eigen, P.Schuster, *The hypercycle. A principle of natural self-organization. Part B: The abstract hypercycle*. Naturwissenschaften **65** (1978), 7-41

M.Eigen, P.Schuster, *The hypercycle. A principle of natural self-organization. Part C: The realistic hypercycle*. Naturwissenschaften **65** (1978), 341-369

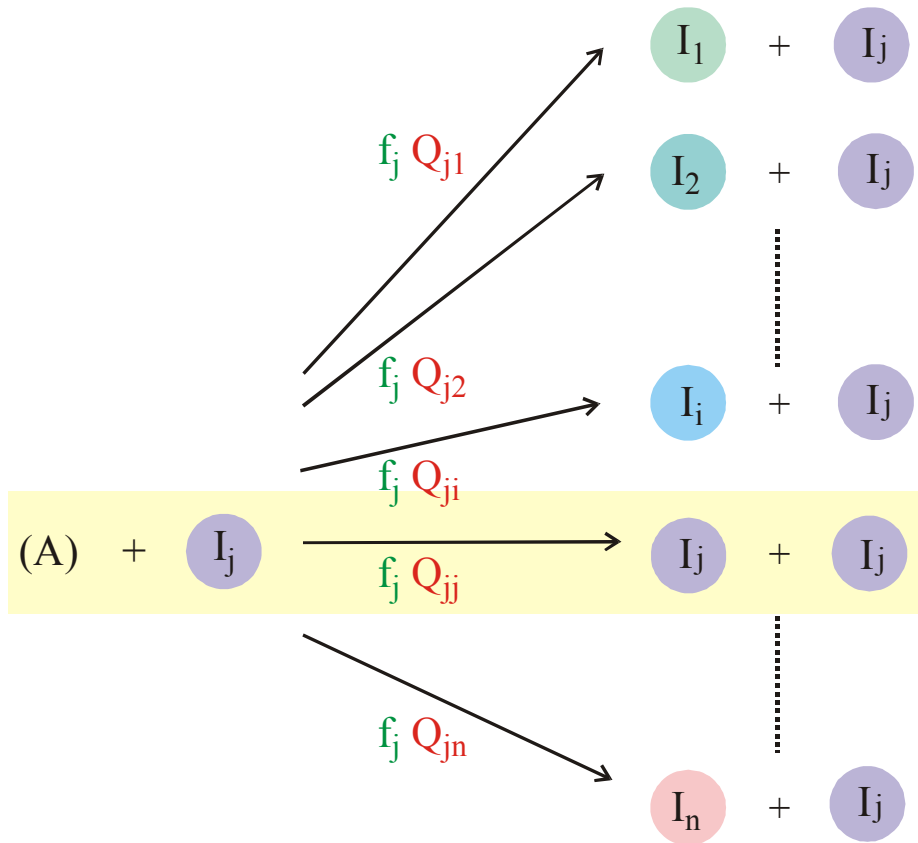
J. Swetina, P. Schuster, *Self-replication with errors - A model for polynucleotide replication*.

Biophys.Chem. **16** (1982), 329-345

J.S. McCaskill, *A localization threshold for macromolecular quasispecies from continuously distributed replication rates*. J.Chem.Phys. **80** (1984), 5194-5202

M.Eigen, J.McCaskill, P.Schuster, *The molecular quasispecies*. Adv.Chem.Phys. **75** (1989), 149-263

C. Reidys, C.Forst, P.Schuster, *Replication and mutation on neutral networks*. Bull.Math.Biol. **63** (2001), 57-94



$$\frac{dx_i}{dt} = \sum_j f_j Q_{ji} x_j - x_i \Phi$$

$$\Phi = \sum_j f_j x_j ; \quad \sum_j x_j = 1 ; \quad \sum_i Q_{ij} = 1$$

$$[I_i] = x_i \ll 0 ; \quad i = 1, 2, \dots, n ;$$

$$[A] = a = \text{constant}$$

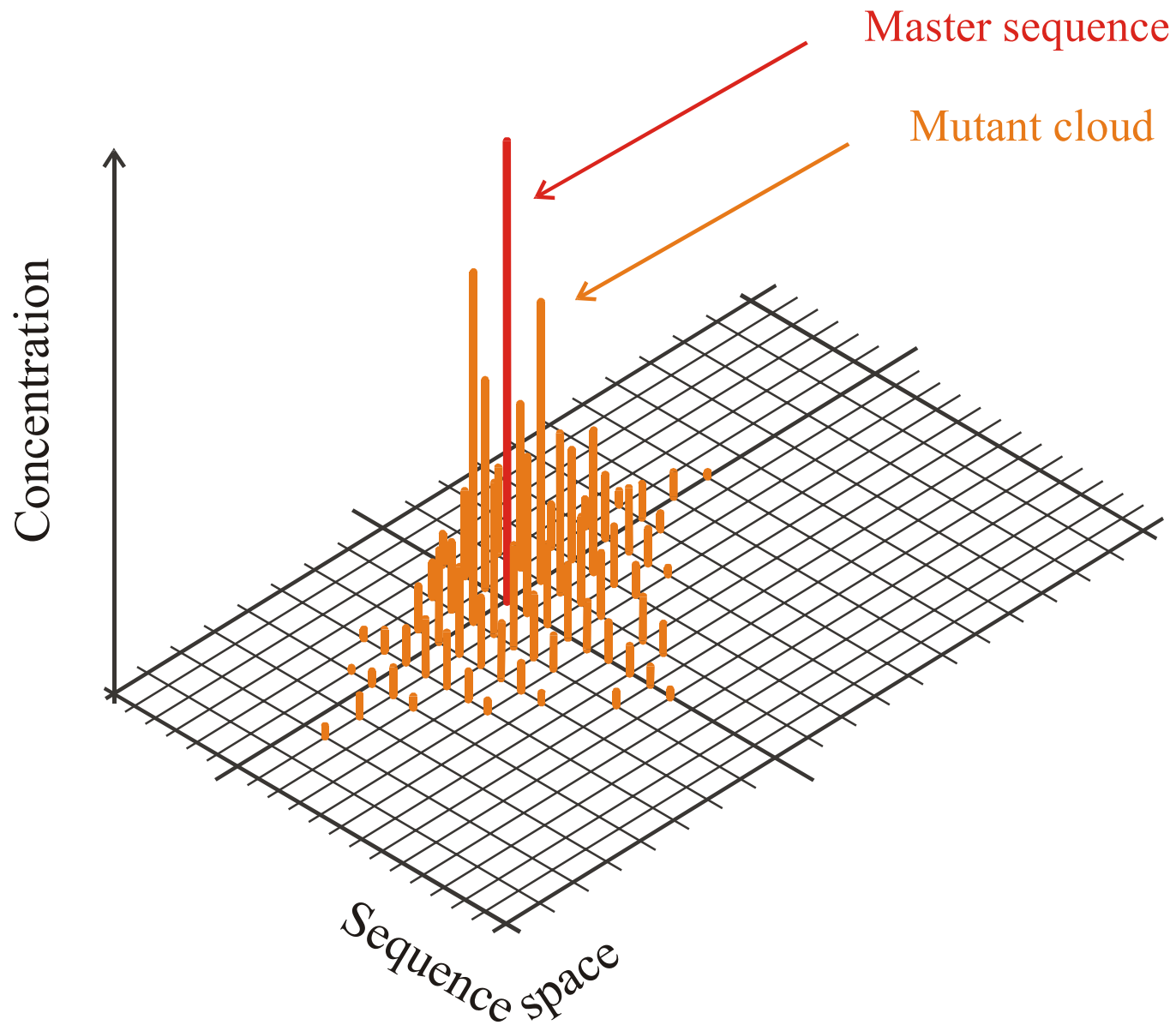
$$Q_{ij} = (1-p)^{\ell-d(i,j)} p^{d(i,j)}$$

p Error rate per digit

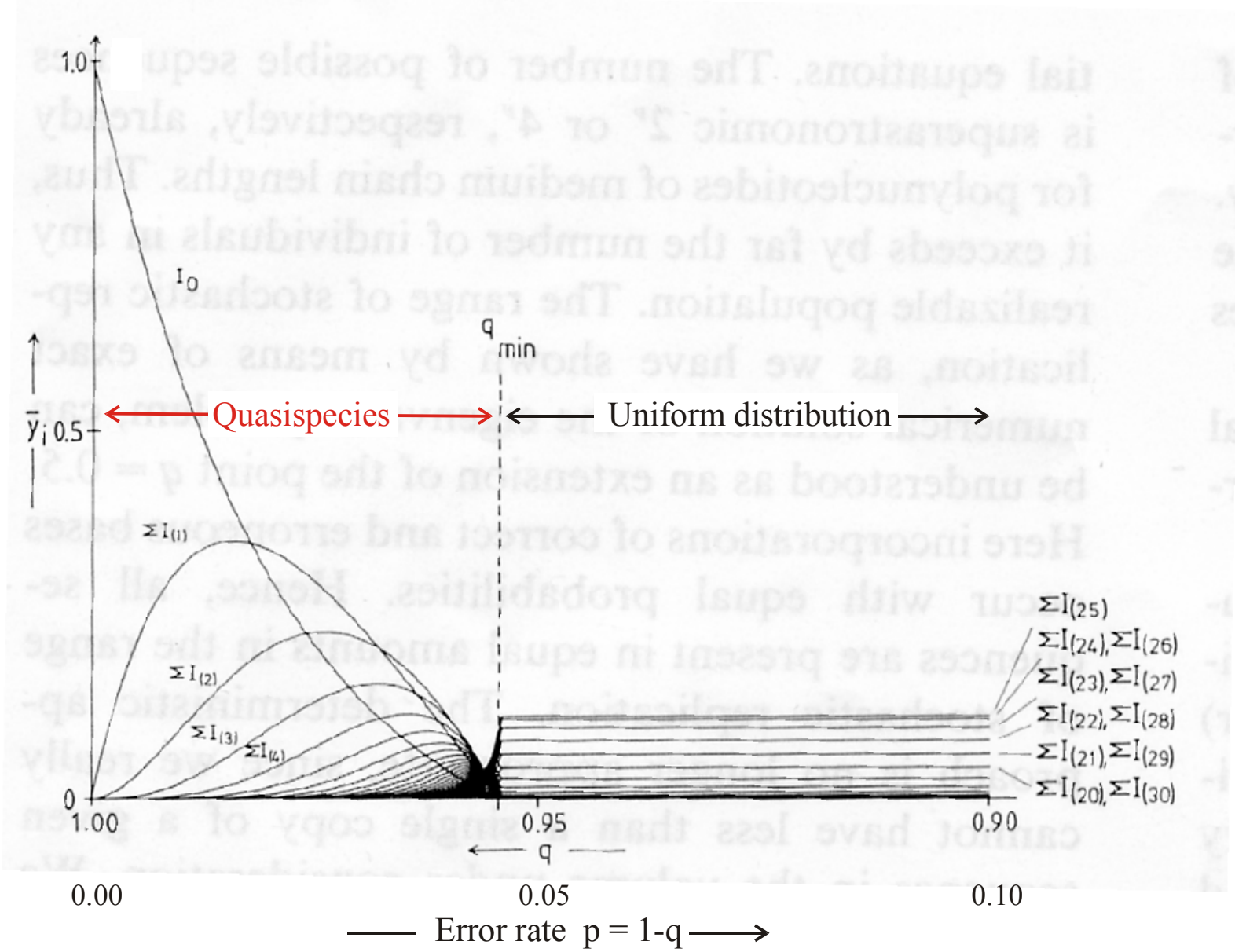
ℓ Chain length of the polynucleotide

$d(i,j)$ Hamming distance between I_i and I_j

Chemical kinetics of replication and mutation as parallel reactions



The molecular quasispecies in sequence space



Quasispecies as a function of the replication accuracy q

Optimization of RNA molecules *in silico*

W.Fontana, P.Schuster, *A computer model of evolutionary optimization*. Biophysical Chemistry **26** (1987), 123-147

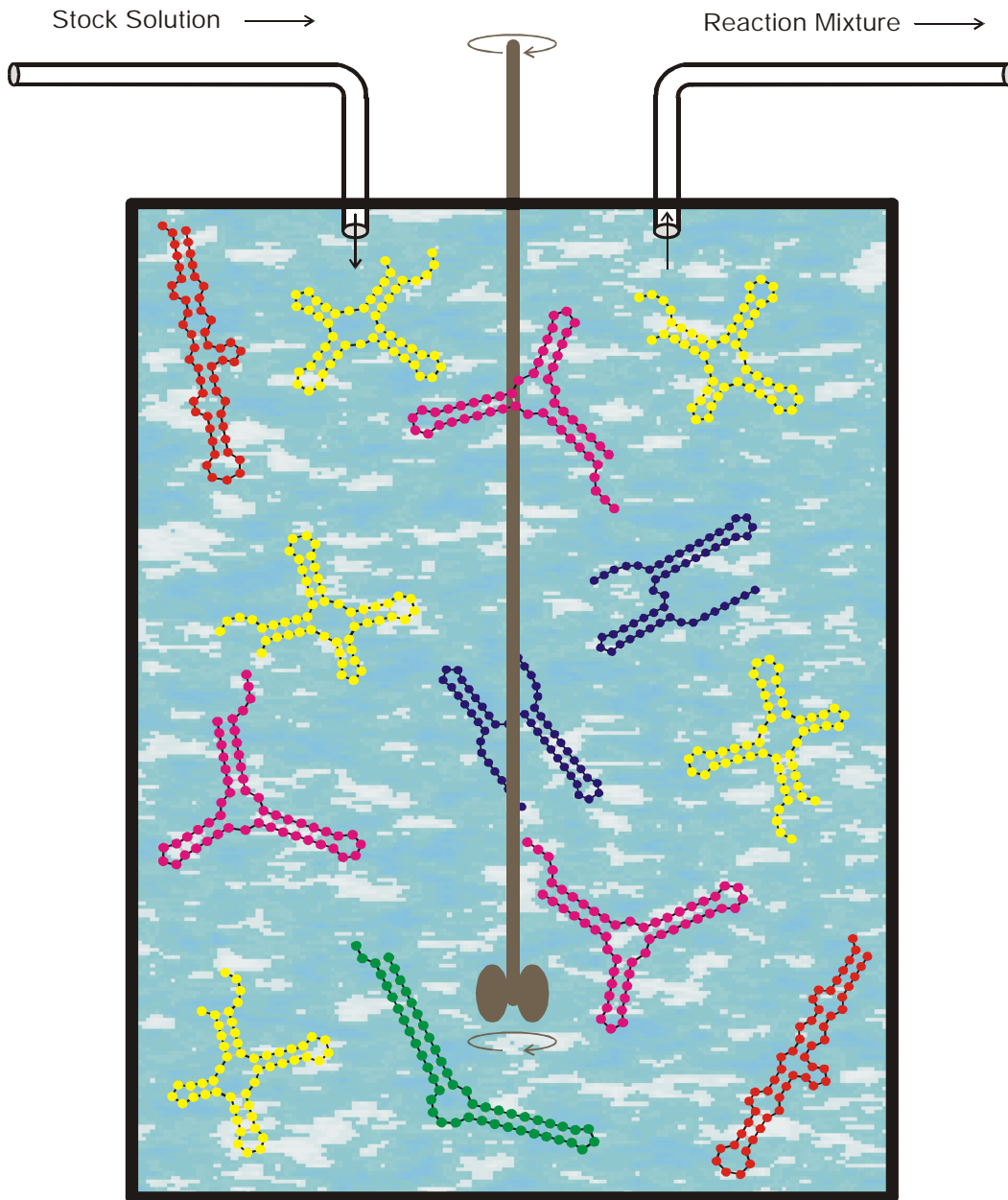
W.Fontana, W.Schnabl, P.Schuster, *Physical aspects of evolutionary optimization and adaptation*. Phys.Rev.A **40** (1989), 3301-3321

M.A.Huynen, W.Fontana, P.F.Stadler, *Smoothness within ruggedness. The role of neutrality in adaptation*. Proc.Natl.Acad.Sci.USA **93** (1996), 397-401

W.Fontana, P.Schuster, *Continuity in evolution. On the nature of transitions*. Science **280** (1998), 1451-1455

W.Fontana, P.Schuster, *Shaping space. The possible and the attainable in RNA genotype-phenotype mapping*. J.Theor.Biol. **194** (1998), 491-515

B.M.R.Stadler, P.F.Stadler, G.P.Wagner, W.Fontana, *The topology of the possible: Formal spaces underlying patterns of evolutionary change*. J.Theor.Biol. **213** (2001), 241-274

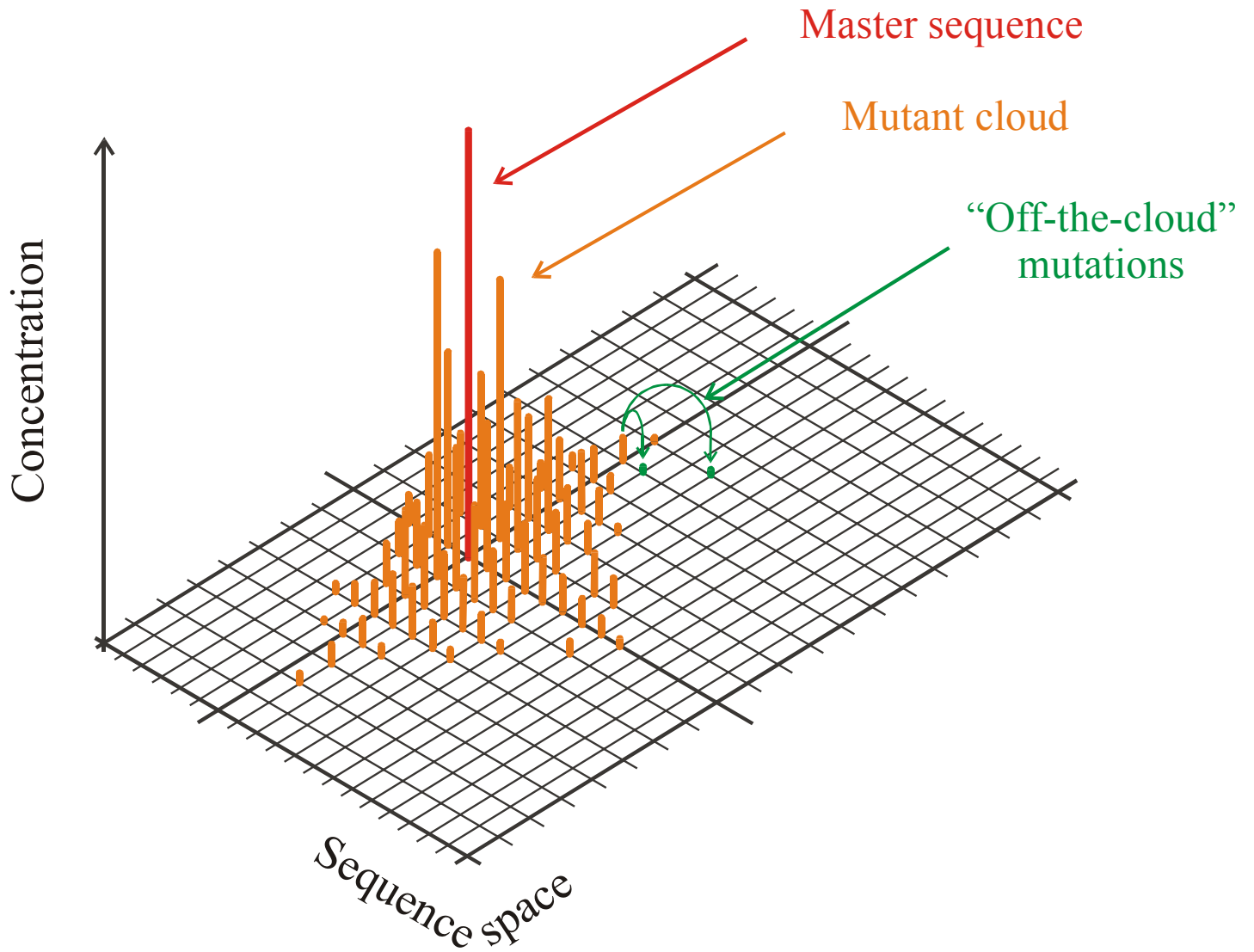


Fitness function:

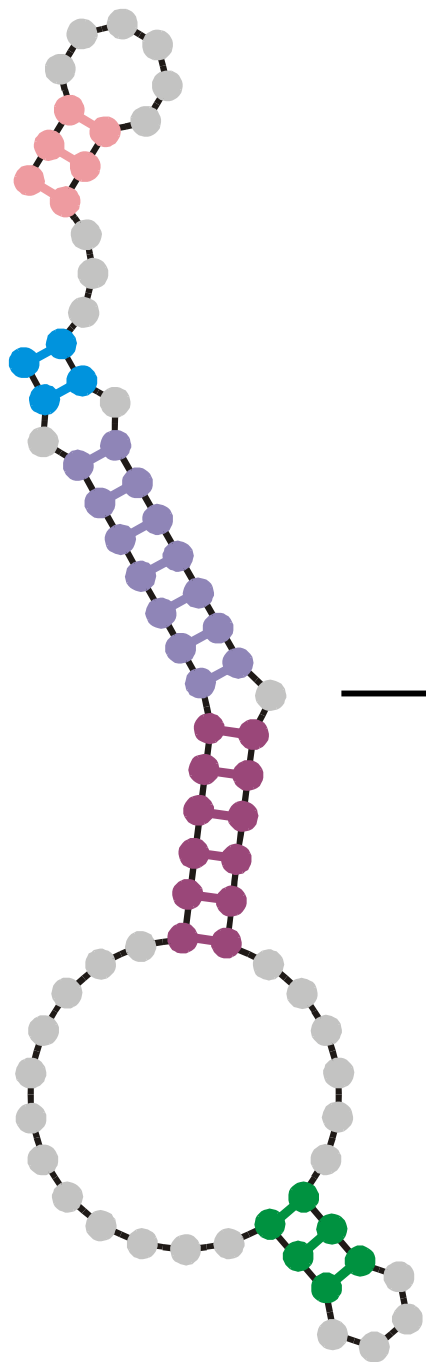
$$f_k = [/ [U + \delta d_S^{(k)}]$$

$$\delta d_S^{(k)} = d^s(I_k, I_h)$$

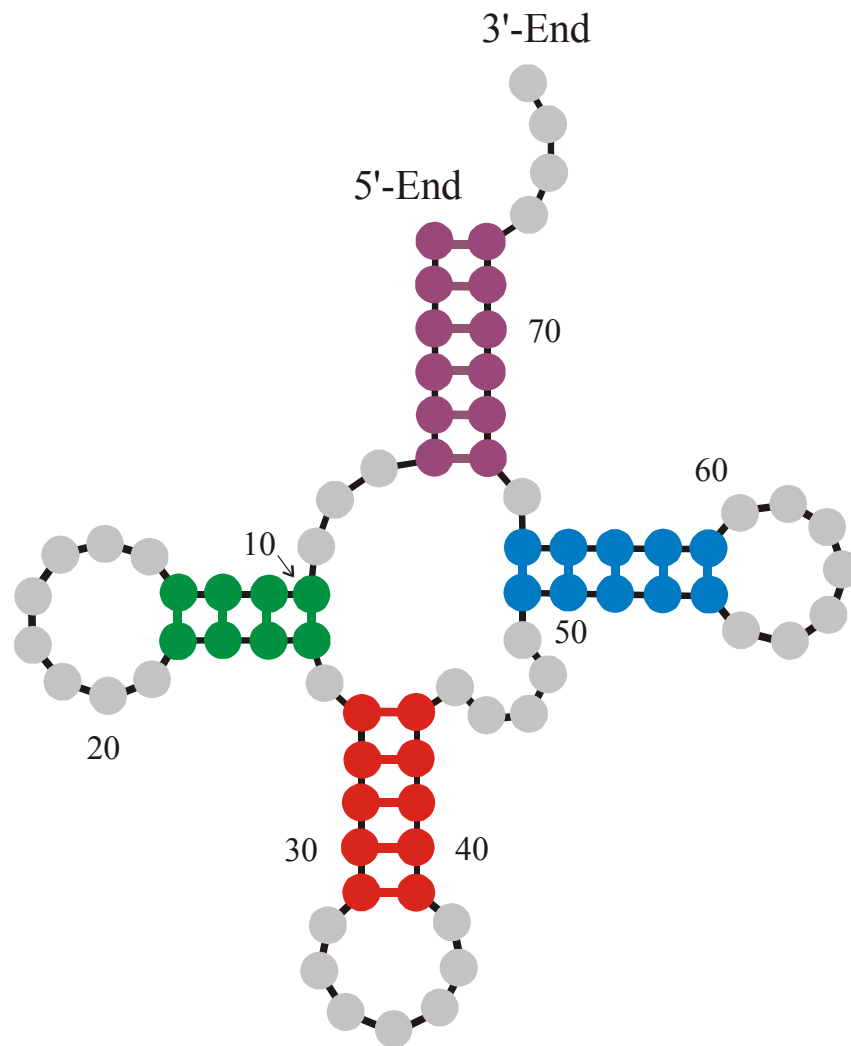
The flowreactor as a device for studies of evolution *in vitro* and *in silico*



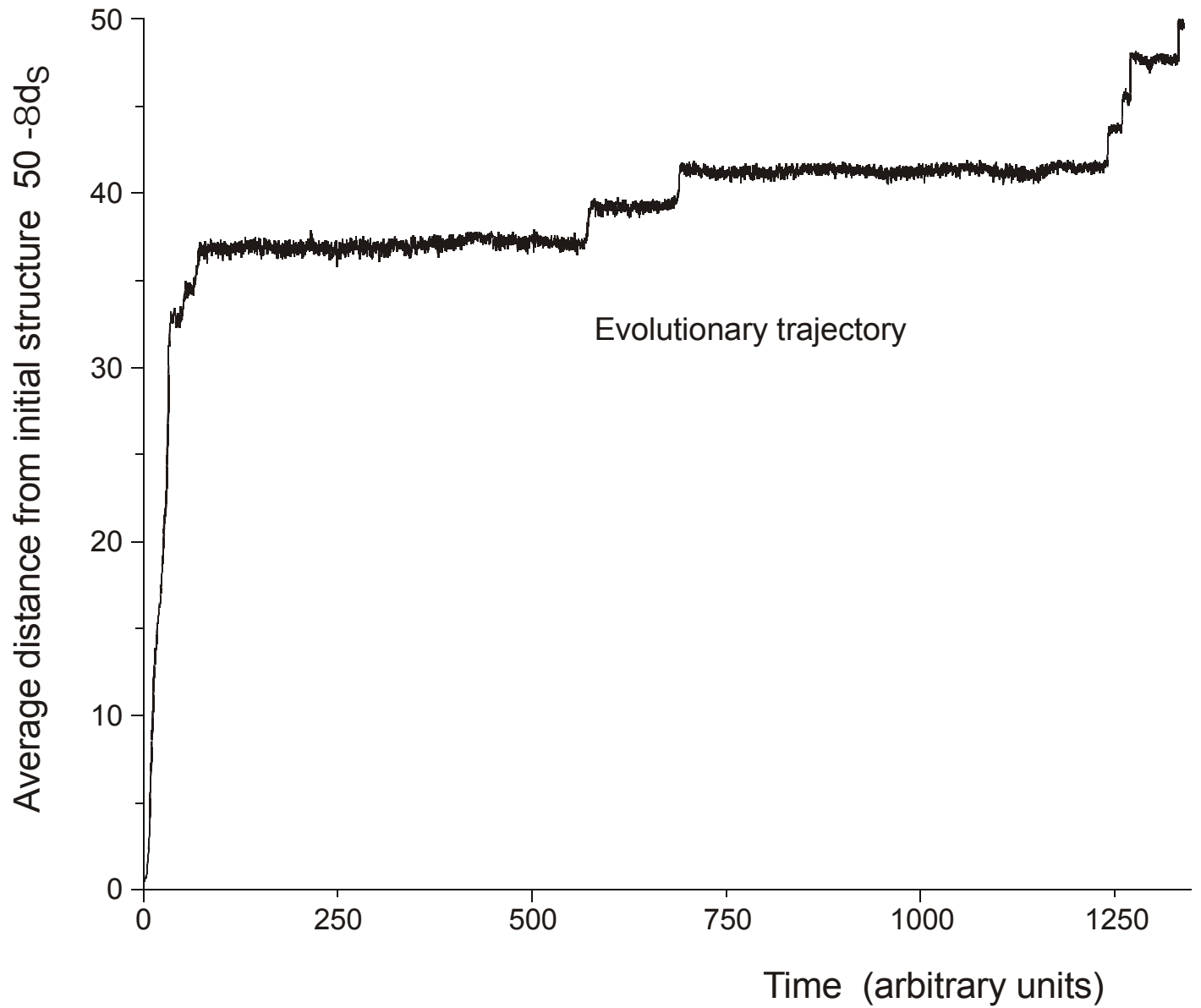
The molecular quasispecies
in sequence space



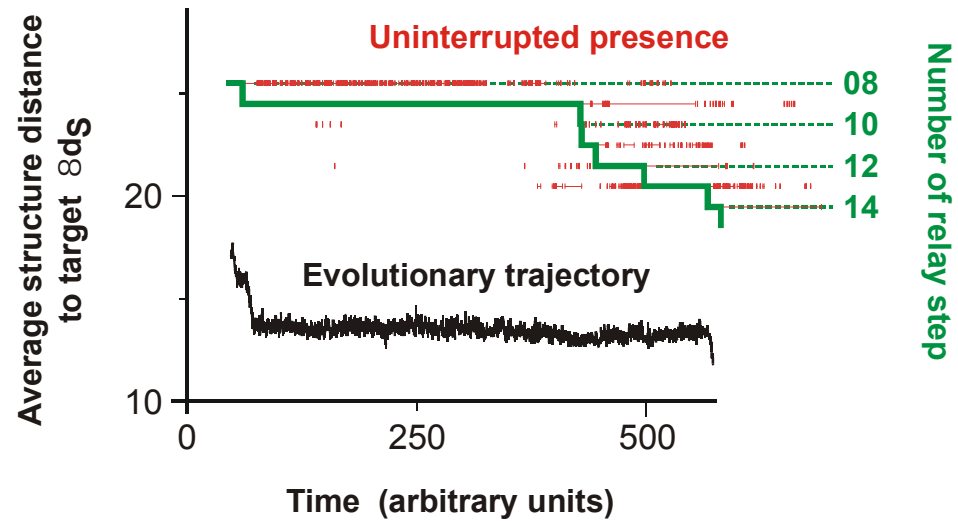
Randomly chosen
initial structure



Phenylalanyl-tRNA as
target structure



In silico optimization in the flow reactor: Trajectory (**biologists' view**)



entry 8 GGUAUGGGCGUUGAAUAGUAGGGUUUAAACCAAUCGGCAACGAUCUCGUGUGCGCAUUUCAUAUCCCGUACAGAA
 .(((((((((((((. (((.))))))))(((((.)))))))))
 exit 8 GGUAUGGGCGUUGAAUAUAJAGGGUUUAAACCAAUCGGCCAACGAUCUCGUGUGCGCAUUUCAUAUCCAUAACAGAA
 entry 9 GGUAUGGGCGUUGAAUAAUAGGGUUUAAACCAAUCGGCCAACGAUCUCGUGUGCGCAUUUCAUAUAACCAUACAGAA
 .(((((((((. (((.))))))))(((((.)))))))))
 exit 9 UGGAUGGACGUUGAAUAACAAGGUAUCGACCAAACAACCAACGAGUAAGUGUGUACGCCCCACACACCGUCCCAAG
 entry 10 UGGAUGGACGUUGAAUAACAAGGUAUCGACCAAACAACCAACGAGUAAGUGUGUACGCCCCACACACCGUCCCAAG
 .((((((. (((.))))))))(((((.))))))
 exit 10 UGGAUGGACGUUGAAUAACAAGGUAUCGACCAAACAACCAACGAGUAAGUGUGUACGCCCCACACAGCGUCCCAAG

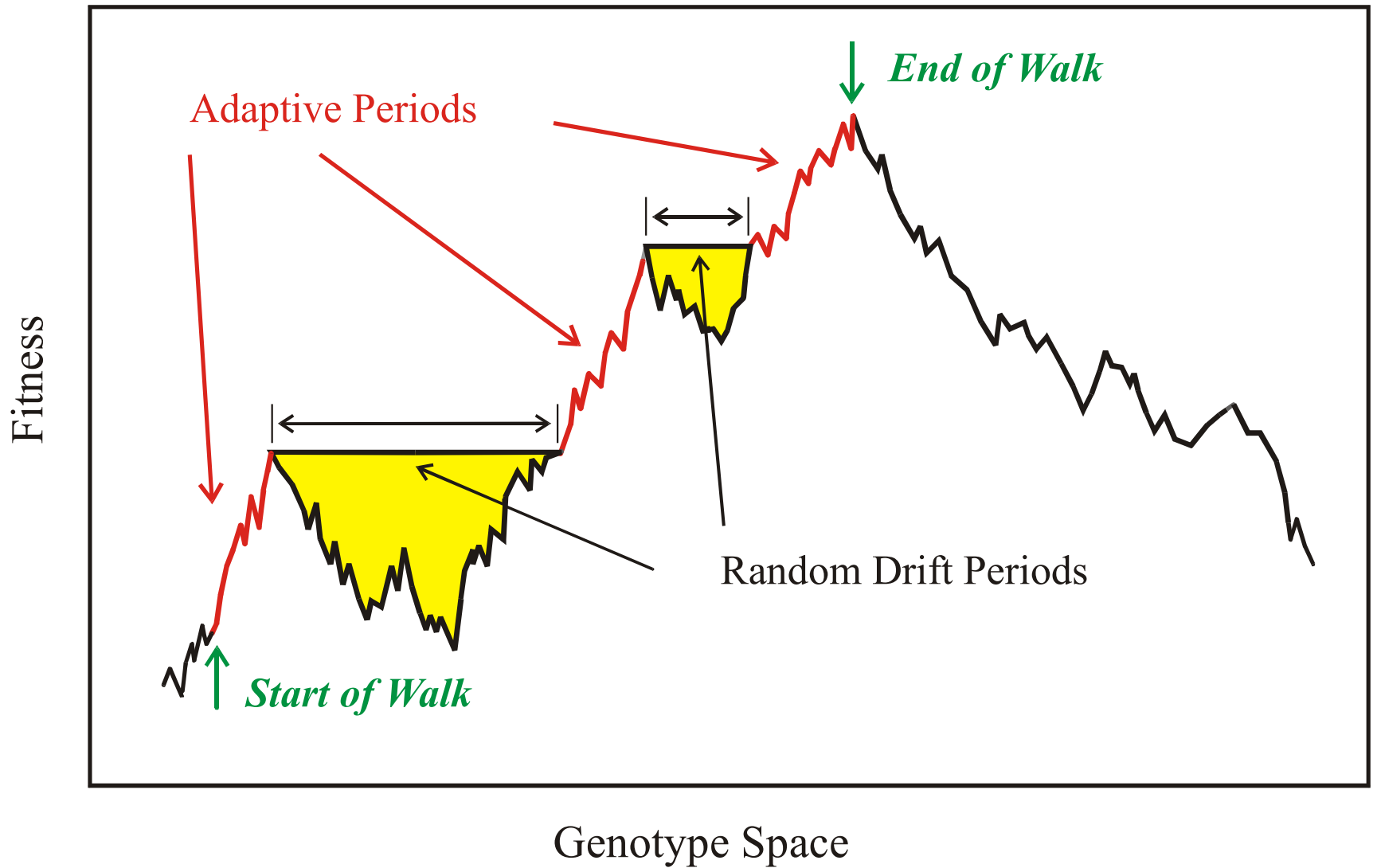
Transition inducing point mutations

Neutral point mutations

Neutral genotype evolution during phenotypic stasis

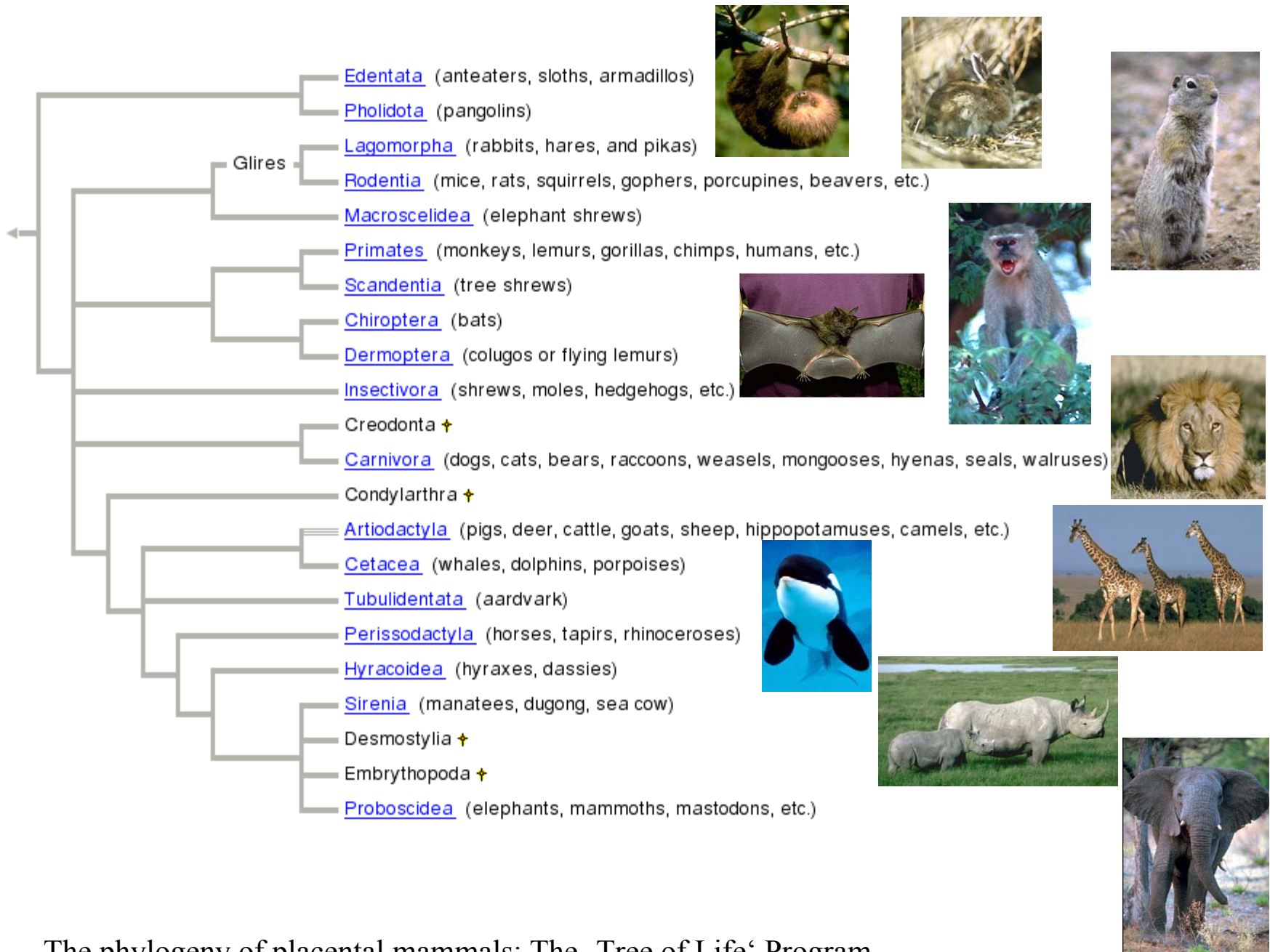
„... Variations neither useful nor injurious would not be affected by natural selection, and would be left either a fluctuating element, as perhaps we see in certain polymorphic species, or would ultimately become fixed, owing to the nature of the organism and the nature of the conditions. ...“

Charles Darwin, Origin of species (1859)



Evolution in genotype space sketched as a non-descending walk in a fitness landscape

1. Theory of evolution, science, and religion
2. Genetics and the theory of evolution
3. Evolution experiments in the laboratory
4. **Molecular genetics and the tree of life**



The phylogeny of placental mammals: The 'Tree of Life' Program

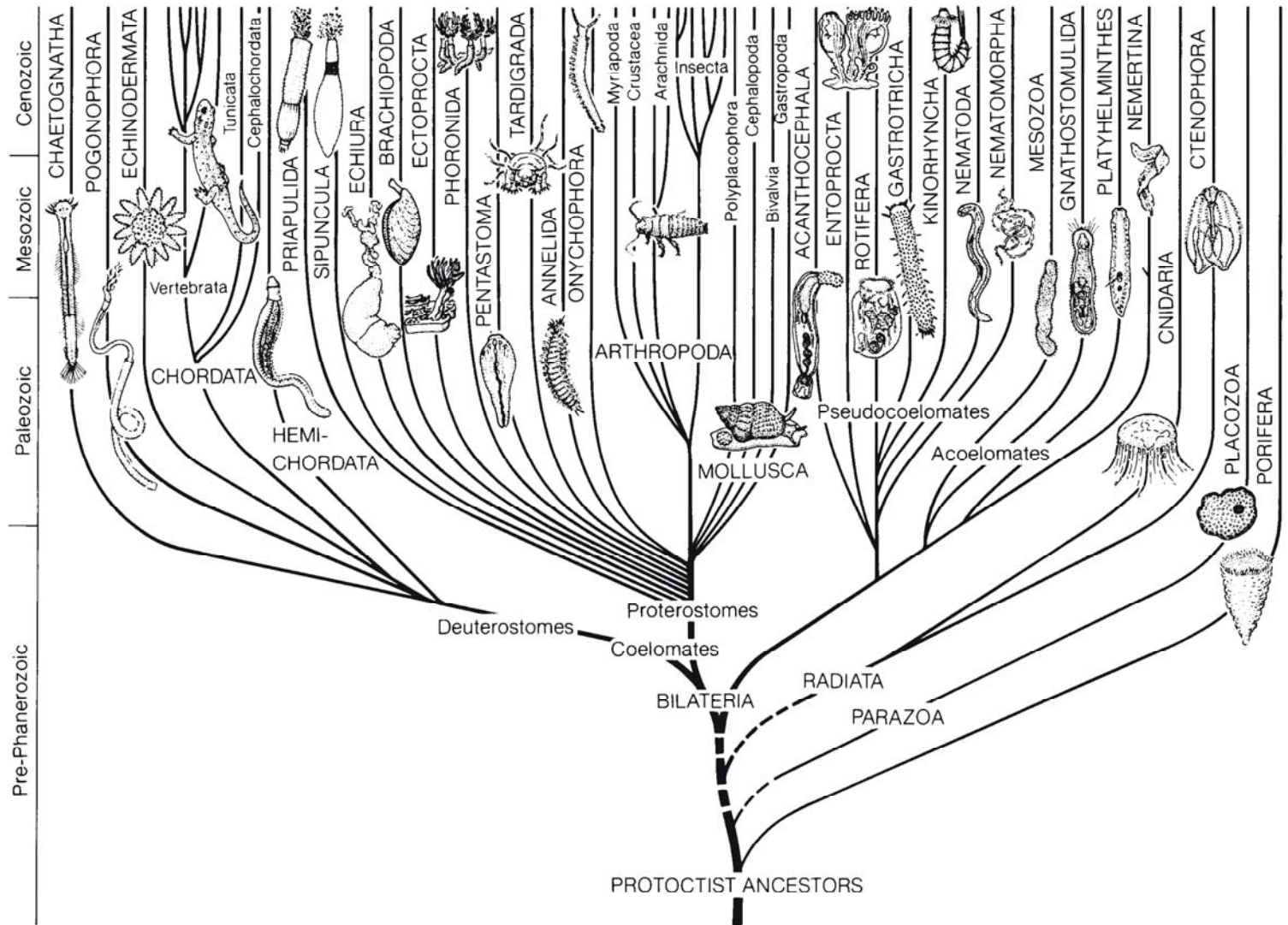
Gerhard Braunitzer, 1929 - 1989



		A														B																
α	Val	Leu	Ser	Pro	Ala	Asp	Lys	Thr	Asp	Val	Lys	Ala	Ala	Try	Gly	Lys	Val	Gly	Ala	His	Ala	Gly	Glu	Tyr	Gly	Ala	Glu	Ala	Leu			
β	Val	His	Leu	Thr	Pro	Glu	Glu	Lys	Ser	Ala	Val	Thr	Ala	Leu	Try	Gly	Lys	Val	Asp		Val	Asp	Glu	Val	Gly	Gly	Glu	Ala	Leu			
I																								Gly								
		C										CD							D					E								
α	Glu	Arg	Met	Phe	Leu	Ser	Phe	Pro	Thr	Thr	Lys	Thr	Tyr	Phe	Pro	His	Phe		Asp	Leu	Ser	His						Gly	Ser	Ala		
β	Gly	Arg	Leu	Leu	Val	Val	Tyr	Pro	Try	Thr	Glu	Arg	Phe	Phe	Glu	Ser	Phe	Gly	Asp	Leu	Ser	Thr	Pro	Asp	Ala	Val	Met	Gly	Asp	Pro		
I								Pro								Phe																
		EF														F																
α	Glu	Val	Lys	Gly	His	Gly	Lys	Lys	Val	Ala	Asp	Ala	Leu	Thr	Asp	Ala	Val	Ala	His	Val	Asp	Asp	Met	Pro	Asp	Ala	Leu	Ser	Ala	Leu	Ser	Asp
β	Lys	Val	Lys	Ala	His	Gly	Lys	Lys	Val	Leu	Gly	Ala	Phe	Ser	Asp	Gly	Leu	Ala	His	Leu	Asp	Asp	Leu	Lys	Gly	Thr	Phe	Ala	Thr	Leu	Ser	Glu
I															His											Leu						
		FG				G										GH																
α	Leu	His	Ala	His	Lys	Leu	Arg	Val	Asp	Pro	Val	Asp	Phe	Lys	Leu	Leu	Ser	His	Cys	Leu	Leu	Val	Thr	Leu	Ala	Ala	His	Leu	Pro		Ala	Glu
β	Leu	His	Cys	Asp	Lys	Leu	His	Val	Asp	Pro	Glu	Asp	Phe	Arg	Leu	Leu	Gly	Asp	Val	Leu	Val	Cys	Val	Leu	Ala	His	His	Phe	Gly		Lys	Glu
I					His																											
		H																														
α	Phe	Thr	Pro	Ala	Val	His	Ala	Ser	Leu	Asp	Lys	Phe	Leu	Ala	Ser	Val	Ser	Thr	Val	Leu	Thr	Ser	Lys	Tyr	Arg							
β	Phe	Thr	Pro	Pro	Val	Glu	Ala	Ala	Tyr	Glu	Lys	Val	Val	Ala	Gly	Val	Ala	Asp	Ala	Leu	Ala	His	Lys	Tyr	His							
I											Lys											Tyr										

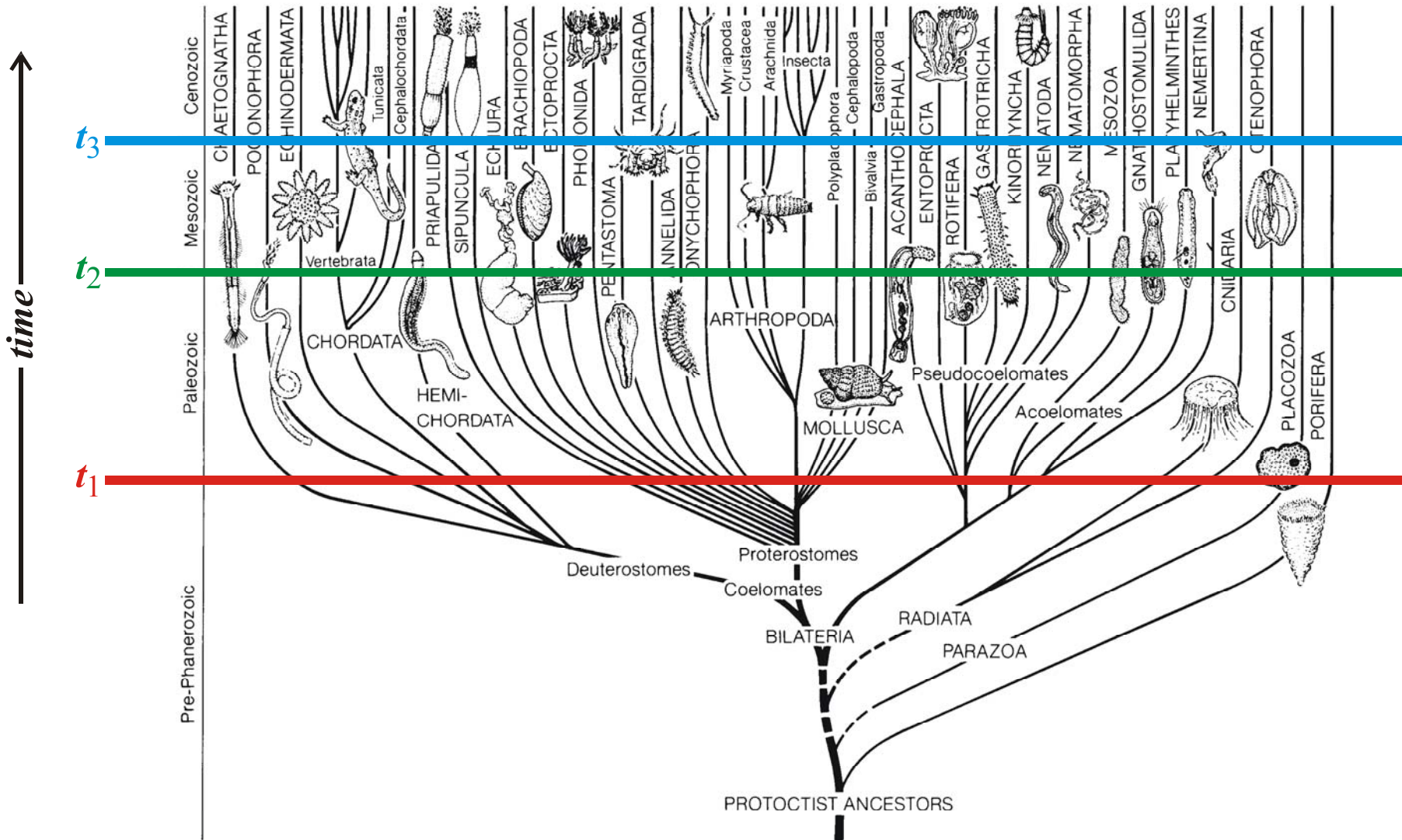
Tab. 4: Die invarianten Reste (I) der Hämoglobine der Vertebraten.

Hemoglobin sequences in different vertebrates



Phylogenetic tree of animal kingdom

Lynn Margulis & Karlene V. Schwarz, *Five Kingdoms. An illustrated guide to the Phyla of Life on Earth*. W.H. Freeman & Co., San Francisco, 1982, p. 160.



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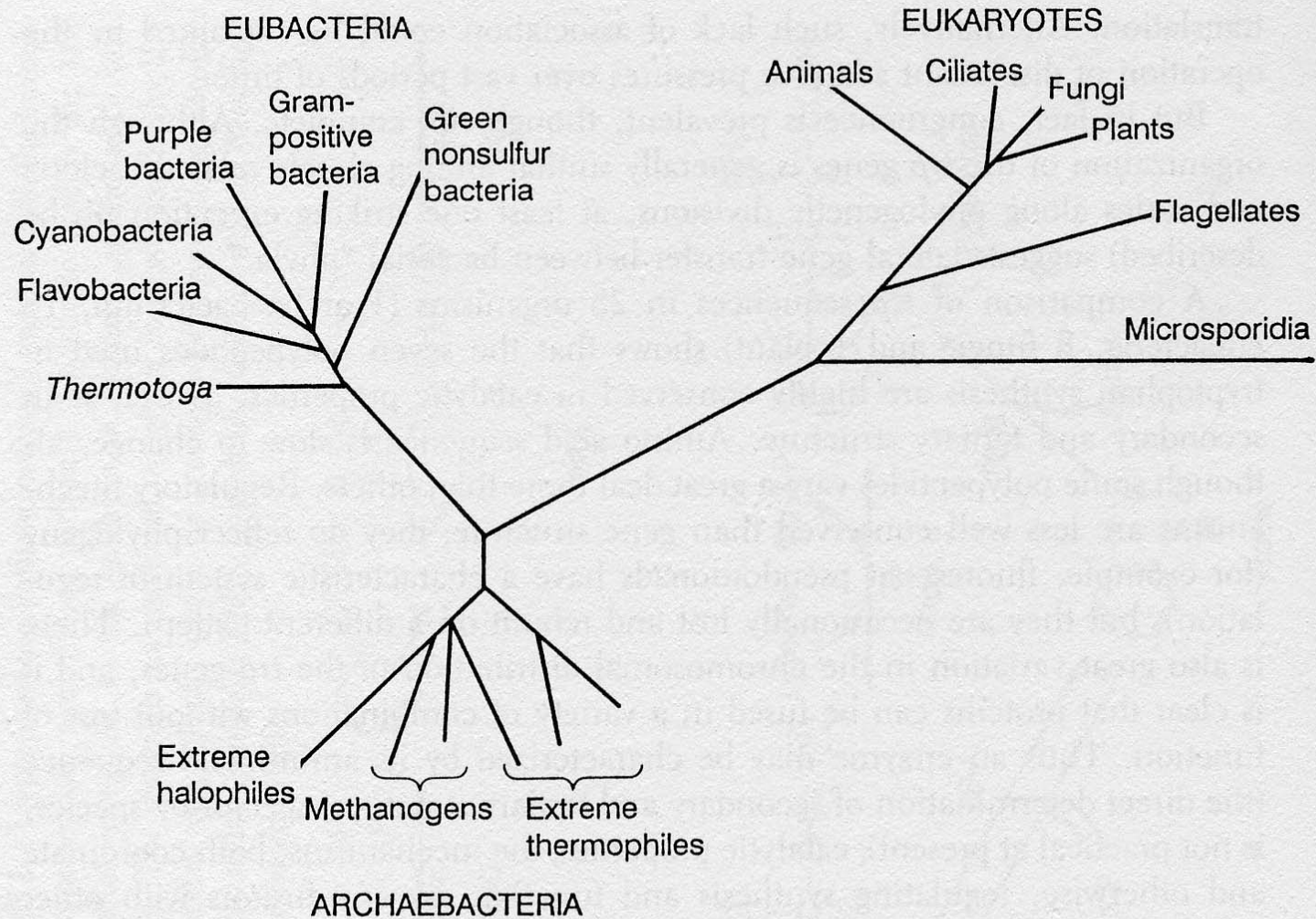


FIGURE 2. The two bacterial phylogenies, taken from the universal phylogenetic tree determined from rRNA sequence comparisons (Woese, 1987).

Evolution at the molecular level.

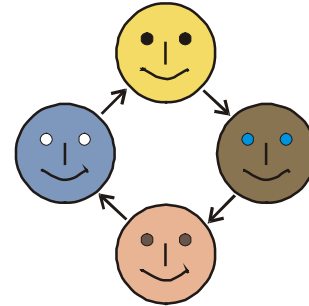
R.K. Selander, A.G. Clark, T.S. Whittam, eds. Sinauer Associates, 1991.

Stage I:
**Independent Competing
Replicators**



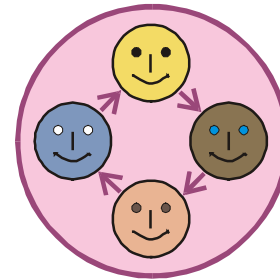
Darwinian evolution

Stage III:
**Functionally Coupled
Replicators**



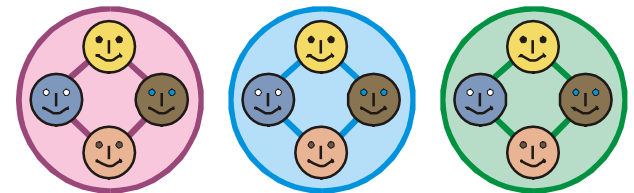
Parasite

Stage IV:
New Unit of Selection



Parasite

Stage V:
**Independent Competing
Units**



Darwinian evolution

A mechanism for major transitions
in evolution

Zur gleichen Zeit schreien viele nach einer neuen Biologie. Man liest, sie wollen „Integrative Biologie“ machen, oder „Systembiologie“. Kaum einer nennt es beim richtigen Namen: Theoretische Biologie. Weil diese einen schlechten Klang hat. Ich jedoch denke, ich kann die Sünden der Vergangenheit vergeben und nehme das Wort: Wir brauchen eine Theorie, die das alles einschließt. Stellen Sie sich doch nur mal vor, wir müssen am Ende all dieses Zeug nicht nur unter Fachleuten besprechen, sondern müssen es an Universitäten lehren, in der Schule, und es der Öffentlichkeit erklären. Wie sollen wir das machen ohne umfassende Theorie? Das, denke ich, ist die Herausforderung, der wir uns stellen müssen.

At the same time people are crying for a new biology. They say, they want to make “Integrative Biology” or “Systems Biology”. Hardly anyone calls it by its proper name: Theoretical Biology. Because it has a bad reputation. I think, however, I can remit the sins of the past and declare: We need a theory, which comprises all that (*Molecular, Structural, Cellular, Developmental, , and Evolutionary Biology*). Imagine, eventually, we not only need to discuss all this stuff with our expert colleagues, but we have to teach it at universities, at schools, and to the public. How could we manage without a comprehensive theory? This is the challenge we have to meet.

Sydney Brenner, Nobelpreisträger 2002, im Gespräch: *„Eine einsame Stimme aus der Prägenomik Ära“*. Laborjournal 2002, Heft 4:28 – 33.

