spontaneous generation of genetic polymers

Ernesto Di Mauro Università di Roma "Sapienza"



or ... ab initio nanotechnology

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atomic transformations

$$3 \ {}^{4}\text{He} \longrightarrow {}^{12}\text{C}$$

$${}^{12}\text{C} + {}^{4}\text{He} \longrightarrow {}^{16}\text{O}$$

$${}^{16}\text{O} + {}^{16}\text{O} \longrightarrow {}^{31}\text{P} + {}^{1}\text{H}$$

$${}^{1}\text{H} \xrightarrow{\text{CN}}{\text{CNO}} {}^{4}\text{He} + {}^{N} \underset{\text{product}}{\text{secondary}}$$

$$8 \ {}^{4}\text{He} \longrightarrow {}^{32}\text{S}$$

















Synthesis of purine and pyrimidine bases, N⁹-formylpurines, and acyclonucleosides from formamide





phosphate minerals

Variscite $AI(PO_4)(H_2O)_2$ Lazulite $Mg[AI(PO_4)(OH)]_2$ • Augelite $Al_2 PO_4(H_2O)_3$ Childrenite $Mn^{2+}[AI(PO_4)(OH)_2(H_2O)]$ • Wavellite $Al_3(OH)_3(PO_4)_2(H_2O)_5$ Triphylite $LiFe^{2+}(PO_{4})$ Hydroxylapatite Ca₅(PO₄)₃OH Eosphorite $Fe^{2+}[AI(PO_4)(OH)_2(H_2O)]$ Hureaulite $Mn^{2+}_{5}(PO_{3}(OH)_{2}(PO_{4})(H_{2}O)_{4})$ • Vauxite $Fe^{2+}AI_2(PO_4)(OH)_2(H_2O)_6$ Reddingite $Mn^{2+}_{3}(PO_{4})_{2}(H_{2}O)_{3}$ Fairfieldite $Ca_2 [Mn^{2+}(PO_4)(H_2O)_2]$ Purpurite Mn³⁺(PO₄) Laueite $Mn^{2+}[Fe^{3+}_{2}(PO_{4})_{2}(OH)_{2}(H_{2}O)_{2}](H_{2}O)_{4}(H_{2}O)_{2}$ • Ludlamite Fe^{2+}_{3} (PO₄)₂(H₂O)₄ Rockbridgeite $Fe^{2+}Fe^{3+}_{4}(PO_{4})_{3}(OH)_{5}$ • Vivianite $Fe^{2+}_3(PO_4)_2(H_2O)_8$ Fluorapatite $Ca_5(PO_4)_3F$ Strengite $Fe^{3+}(PO_4)(H_2O)_2$ Crandallite $CaAl_3(PO_4)_2(OH)_5+H_2O$ Cacoxenite $Fe^{3+}_{25}(PO_4)_{17}O_6(OH)_{12}(H_2O)_{75}$ Anapaite $Ca_2[Fe^{2+}(PO_4)_2(H_2O)_4]$ • Libethenite $Cu^{2+}_{2}(PO_{4})(OH)$ Scholzite $CaZn_2(PO_4)_2(H_2O)_2$ • Cornetite $Cu^{2+}_{3}(PO_{4})(OH)_{3}$ Turquoise $Cu^{2+}Al_{6}(PO_{4})_{4}(OH)_{8}(H_{2}O)_{4}$ Tarbuttite $Zn_2(PO_4)(OH)$ Pyromorphite Pb₅(PO₄)₃Cl Monazite Ce(PO4) Autunite $Ca[(UO_2) (PO_4)]_2(H_2O)_{10-12}$ Montebrasite Li[Al(PO₄)(OH)] Torbenite $Cu^{2+}[(UO_2 (PO_4)]_2 (H_2O)_8]$ Beryllonite $Na[BePO_4]$ Herderite Ca[BePO₄F] Brasilianite $NaAl_3(PO_4)_2(OH)_4$ Pseudomalachite $Cu^{2+}_{5}(PO_{4})_{2}(OH)_{4}$ Wardite $NaAl_3(OH)_4(PO_4)_2(H_2O)_2$ Calcioferrite $Ca_{4}MgFe_{4}(PO_{4})_{6}(OH)_{4}(H_{2}O)_{13}$ Hydroxylapatite $Ca_5(PO_4)_3OH$

Synthesis of purine and pyrimidine bases from phosphate minerals







RNA polymerization in water



Costanzo et al. J. Biol. Chem. 2009

Polymerizing 3',5'-cyclic GMP



Non-enzymatic polymerization of 3',5'-cGMP



Non-enzymatic polymerization of 3',5'-cAMP



85°C; TrisHCl pH 8,2

the warm little pond



NI

Non-enzymatic mechanisms for the generation of long RNA sequences in water

- Synthesis of RNA chains from cyclic nucleotides
- RNA chain extension
- RNA ligation

• RNA stability in different conditions of temperature and pH

RNA chain extension in water



Costanzo et al. J.Biol. Chem. 2009





Polymerizing 3',5'-cyclic GMP on a 5'A₁₂C₁₂3' oligo



The plausible mechanism for sequence extension



Ar explosive, concentration-dependent chain-extension reaction



Sequence-directed terminal ligation



PolyG[®] + PolyC sequence-directed terminal ligation



5'

5'

G₂₄

C₂₄

P

PolyC₂₄ pH 5 60°C





PolyC[®]+ PolyG sequence-directed terminal ligation



Stacking-directed terminal ligation



Pino et al. J. Biol. Chem. 2008

RNA ligation in H₂O 23-, 24-mer







Combinations thereof — heterogeneous sequences

RNA multiplication in water



Differential stability of phosphoester bonds in ribo-monomers and oligomers



Saladino et al. J. Biol. Chem. 2006

formamide %



Environment induces selection for sequence complexity



"Something came from nothing because it is more stable than nothing"

The Comprehensible Cosmos, Victor Stenger

Directed Assembly of One-Dimensional Nanostructures into Functional Networks

Yu Huang,^{1*} Xiangfeng Duan,^{1*} Qingqiao Wei,¹ Charles M. Lieber^{1,2}†

Fig. 1. Schematic of fluidic channel structures for flow assembly. (A) A channel formed when the PDMS mold was brought in contact with a flat substrate. NW assembly was carried out by flowing an NW suspension inside the channel with a controlled flow rate for a set duration. Parallel arrays of NWs are observed in the flow direction on the substrate when the PDMS mold is removed. (B) Multiple crossed NW arrays can be obtained by changing the flow direction sequentially in a layerby-layer assembly process.







Directed Assembly of One-Dimensional Nanostructures into Functional Networks

Yu Huang,¹* Xiangfeng Duan,¹* Qingqiao Wei,¹ Charles M. Lieber^{1,2}†

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United States Patent [19]

Hollenberg et al.

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[45] Date of Patent:

Oct. 1, 1996

[54] DNA AND DNA TECHNOLOGY FOR THE CONSTRUCTION OF NETWORKS TO BE USED IN CHIP CONSTRUCTION AND CHIP PRODUCTION (DNA-CHIPS)

- [76] Inventors: Cornelis P. Hollenberg, Chopinstrasse 7, 4000, Düsseldorf, Germany; Ernesto di Mauro, Via Andrea Fulvio 10, 00162, Rome, Italy
- [21] Appl. No.: 532,542
- [22] Filed: Sep. 25, 1995

Related U.S. Application Data

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 [58] Field of Search 435/6, 810, 287,
- 435/299; 536/22.1, 25.3; 437/1, 15, 16,



D. Eigler et al., "An Atomic Switch Realized With The Scanning Tunnelling Microscope," Nature, vol. 352, pp. 600–603 (1991).

A. Oliphant et al., "Defining the Sequence Specificity of DNA-Binding Proteins by Selecting Binding Sites from Random-Sequence Oligonucleotides: Analysis of Yeast GCN4 Protein," Molecular and Cellular Biology, 9:2944 (1989).

C. Cantor et al., "Orientation in Electric Fields," Biophysical Chemistry part II, 665-668 (1980).



子は、以下の方法によりマトリックス(この上に 核酸回路網が形成される)に結合させることがで きる:

マイクロマニピュレーターを用いて特異的 (1) 蛋白(即ちラムダー蛋白リプレッサー:下記を参 照)溶液の微細な1滴を、ポリエチレンのような 疎水性表面の上にのせ、乾燥させる。

配列6は自己アニーリングした二本銷構造 (2)の左の末端に位置するような配列で、配列(4)

3' and 5' indicate the 2 extremities of the nucleic acid strand. The enzymatic polymerization of DNA by the enzyme DNA polymerase (ref.7) proceeds by addition of monomers to the 3'-extremity (see ref.1). The letters G,A,C,T are acronyms that indicate the monomeric constituents of the DNA strand; they are nucleotide monophosphates containing respectively a purine (guanine for G, adenine for A) or a pyrimidine (cytosine for C., thymine for T) residue. In DNA polymers these compounds can base pair specifically: G couples 10 always with C, A with T. Therefore a self-annealing reaction in a solution containing the appropriate buffer (2x SSC solution, ref.8 p.447) at 20° C. will produce the molecule. (2) 15

AGATCAAGTCAATT3 TCTAGTTCAGTTAACTATTTTGGAAGCGTAGCTTCC-

- ACTAATCCT5

The left extremity of molecule (2) is the DWIP, the right extremity is the growing point (that is the point onto which additional hybridization or synthetic reactions can be perpoints or switches. Elongation may be obtained by hybridization of a preformed DNA molecule or a reaction of DNA synthesis. Hybridization of nucleic acids is a procedure that exploits the tendency of nucleic acids to anneal to double strand structures (according to the rules mentioned above: A with T, G with C), if the complementary order of the nucleotides that compose the DNA sequence permits it. One synthesizes according to the procedure mentioned

above the following molecule:

GATAAAACCTTCCATAACAAAGTGGTTGAA

The hybridization reaction between molecules (2)+(3) will 40 produce molecule (4):

> AGATCAAGTCAATTGATAAAACCTTC-TCTAGTTCAGTTAACTATTTTGGAAG--CAT AACAAAGTGGTTGAA

-GTAGCTTCCACTAATCCT

This molecule produced by synthesis and hybridization has one DWIP (left) (defined above as "blunt end") and a branched extremity (right). This branched extremity now provides two different growing points that can be used for 55 -TTTTGGAAGGTAGCTTCCACTAATCCT further elongation and branching of the molecule, to produce a network (Scheme 1)). Many DNA sequences can lead to the shown below structure. The length is variable.

Scheme 1:



4 -continued



Single strand interruptions in the DNA strands (indicated in Scheme 1 by the arrows), can be easily filled up by the reaction of the enzyme DNA ligase (commercially available, i.e., from Bethesda Research Laboratories, Bochringer Manformed in order to elongate the chain and/or create branch 25 nheim, etc, see refs. 8,9). The synthesis of oligonucleotides (molecules 1 and 3) can be performed with commercially available apparatus (i.e., from Applied Biosystems or New Brunswick Scientific Company).

The DWIP can be fixed to a solid matrix by several techniques e.g., locally fixed charged molecules or sequence specific DNA binding proteins (as bacteriophage DNA binding proteins, Adenovirus binding protein, lac repressor or synthetic DNA binding proteins) or covalent chemical binding.

Outline 2

35

(3)

(4)

A DNA molecule such as molecule (4) described in outline 1 can be fixed by the following procedure to a matrix onto which the nucleic acid network will be formed:

(i) Place, by the use of a micromanipulator, a microdrop of a solution of a specific protein (i.e. lambda-protein repressor; see below) on a hydrophobic surface like polyethylene and let it dry.

(ii) Synthesize a molecule (5) which contains the sequence (4) and (6) in such an arrangement that sequence 6 is located at the left end of the self-annealed double strand structure:

TACCTCTGGCGGTGATAAGATCAAGTCAATTGAT-

ATGGAGACCGCCACTATTCTAGTTCAGTTAACTA-

Sequence 6 (plus GAACG) Sequence 4 (without GAACG)

-АЛЛАССТТССАТ АЛСАЛАСТОСТТСАА

(iii) Treat the hydrophobic surface with a solution of DNA molecule (5). The specific binding of the DNA molecule to the protein molecule is ensured by the use of the specific DNA-protein interaction. Specificity of such interaction is a well-known phenomenon in biological processes and several DNA-protein interaction systems can be chosen, as detailed in the following paragraph.

Repressors are proteins which regulate gene expression, well described for bacteria and bacteriophages systems

5,561,071

Ar explosive, concentration-dependent chain-extension reaction



Louis Pasteur





• "Omne vivum ex vivo" (1864)



Alexander I. Oparin



"Proiskhozhdenie zhizny" (The Origin of Life") 1924