

THE EFFECTS OF TEMPERATURE ON PROTEIN SYNTHESIS

The two aspects of the temperature response of plants that are most interesting are (a) the short-term responses to high and low temperatures and (b) the adaptive responses.

The short-term responses are largely determined by the activation energies of the complex of reactions that make up the process of protein synthesis. In those studies where the response of growth can be compared with that of protein synthesis, an important generality emerges: Protein synthesis occurs at the highest rate at those temperatures that also give the highest rates of growth for the plant. Beevers *et al.* (1965) have shown that the optimum temperature for NR synthesis in leaves of maize was 35°C. This compares with an optimum for leaf growth at 35°C for leaves of maize (Watts, 1972). For radish, the optimum for protein synthesis was 31°C (Beevers *et al.*, 1965). Radish grows well in cool seasons. Leaf growth in bean, which also does well at cool temperatures, has an optimum for leaf growth at 25–30°C (Wilson and Ludlow, 1969).

In the bacteria, a similar correspondence between growth and protein synthesis holds true. Friedman *et al.* (1969) showed that free 30S and 50S ribosomal subunits accumulate when *E. coli*, a mesophile, is grown at 8°C. In this species, cell division virtually ceases at 8°C. These authors conclude that the formation of some portion of the initiation complex was inhibited by cold but that peptide chains could be completed if they had already been initiated.

Micrococcus cryophilus, a psychrophilic bacterium that grows poorly above 25°C, exhibits an inhibition of protein synthesis above this temperature (Malcolm, 1969). In this species, the block can be traced to the amino acyl transferases for glutamate and proline. These two enzymes lose activity for the cognate species of tRNA at these temperatures, in part because of changes in the tRNA's to configurations that are unfavorable for the enzymes and in part due to changes in the enzymes themselves. Somewhat similar results have been found with the psychrophilic yeast *Candida gelida* (Nash *et al.*, 1969). Several amino acyl tRNA synthetases appear to be denatured at temperatures above the growth optimum (25–30°C). Leucyl tRNA synthetase was most affected. In addition to the synthetases, the formation of ribosome-bound polypeptide chains was also inhibited at high temperatures.

It is perhaps not surprising that the temperature sensitivity of growth and protein synthesis are correlated. Cells require new protein for

Structurefunction Relationships Of Proteins Proceedings Of The Third John

Henry Vogel



Structurefunction Relationships Of Proteins Proceedings Of The Third John:

Structure-function Relationships of Proteins Roy Markham,1976 Current Catalog National Library of Medicine (U.S.),1985 First multi year cumulation covers six years 1965 70 **National Library of Medicine Current Catalog** National Library of Medicine (U.S.), *Nucleic Acid-Protein Recognition* Henry Vogel,2012-12-02 Nucleic Acid Protein Recognition covers the proceedings of a symposium on Nucleic Acid Protein Recognition held at Arden House Harriman Campus of Columbia University on May 30 June 1 1976 The symposium inaugurated the P S Biomedical Sciences Symposia under the sponsorship of the College of Physicians and Surgeons of Columbia University This book is organized into nine part encompassing 31 chapters The opening parts describe the principles of DNA replication and the unique chromatin structure These parts also examine the physical chemistry of the interactions of melting proteins with nucleic acids The third part presents the different types of approaches that can be used to study the function of RNA polymerases and the development of a cell free system that favors Pol II catalyzed transcription from type 2 adenovirus DNA Parts IV and V deal with the sequence determination of wild type and mutant repressor and the restriction and modification of DNA endonucleases while parts VI and VII focus of the recognition of tRNA Part VIII discusses some significant studies on the assembly of ribosomes and the principles of ribosomal interactions Lastly Part IX considers the role of small RNA template in the reaction mechanism of RNA replicases and ribonucleases This part also surveys the so called RNase III cleavage of different types of RNA and the structure of nucleic acid protein complexes *Gene Function* S. Rosenthal,H. Bielka,Ch. Coutelle,2014-05-09 Gene Function contains the proceedings of the 12th Meeting of the Federation of European Biochemical Societies held in Dresden Germany in 1978 The meeting provided a forum for discussing progress in the understanding of gene function and covered topics ranging from the functional organization of chromatin to principles of interactions and recognition models The role of DNA sequence in the recognition of restriction endonucleases and modification enzymes is also examined along with gene expression RNA processing and modification and isolation and synthesis of genes Comprised of 49 chapters this volume begins with an overview of what can be learned from the genetic analysis of the lac repressor followed by a discussion on the topography of the interaction the lac repressor RNA polymerase and histones with DNA The reader is then introduced to complementarity and recognition code between regulatory proteins and DNA chromatin replication in vitro and the cytoplasmic petite mutation in *Saccharomyces cerevisiae* Subsequent chapters explore arc like and helical arrangements of nucleosome cores changes in gene expression during cellular differentiation polyadenylation and processing of pre messenger RNA and the molecular biology of bacteriophages T3 and T7 This book will be of interest to geneticists biochemists and molecular biologists Protein, Structure, Function, and Industrial Applications Federation of European Biochemical Societies,1979 Current Catalog ,1986 First multi year cumulation covers six years 1965 70 *Protein Structure-Function Relationships in Foods* Rickey Y. Yada,R.L. Jackman,2012-12-06 Food proteins constitute a diverse and

complex collection of biological macro molecules Although contributing to the nutritional quality of the foods we consume proteins also act as integral components by virtue of their diverse functional properties The expression of these functional properties during the preparation processing and storage of foods is largely dictated by changes to the structure or structure related properties of the proteins involved Therefore germane to the optimal use of existing and future food protein sources is a thorough understanding of the nature of the relationships between structure and function It is the goal of this book to aid in better defining these relationships Two distinct sections are apparent firstly those chapters which address structure function relationships using a variety of food systems as examples to demonstrate the intricacies of this relationship and secondly those chapters which discuss techniques used to either examine structural parameters or aid in establishing quantitative relationships between protein structure and function The editors would like to thank all contributors for their assistance co operation and above all their patience in putting this volume together and the following companies organizations for their financial support without which it would not have been the success it was Ault Foods Limited Best Foods Canada Limited Natural Sciences and Engineering Research Council of Canada Ontario Ministry of Agriculture and Food Quest International Canada Inc and University of Guelph R Y Y R LJ

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