

Methods of Protein Microcharacterization

by Thomas E. Creighton

Edited by

John E. Salierno

Methods Of Protein Microcharacterization A Practical Handbook

John E. Shively



Methods Of Protein Microcharacterization A Practical Handbook:

Methods of Protein Microcharacterization John E. Shively, 1986-07-08 The book discusses a series of higher dimensional moduli spaces of abelian varieties cubic and K3 surfaces which have embeddings in projective spaces as very special algebraic varieties Many of these were known classically but in the last chapter a new such variety a quintic fourfold is introduced and studied The text will be of interest to all involved in the study of moduli spaces with symmetries and contains in addition a wealth of material which has been only accessible in very old sources including a detailed presentation of the solution of the equation of 27th degree for the li *Methods of Protein Microcharacterization* John E. Shively, 2008-02-22 Milestones in the techniques and methodology of polypeptide structure determination include the determination of the sequence of insulin by Sanger in 1951 1 and the introduction of the repetitive degradation of proteins with phenylisothiocyanate by Edman in 1959 2 The automation of Edman chemistry 3 played a major role in the determination of polypeptide structures Important modifications of Edman chemistry include the solid phase approach by Laursen in 1971 4 and the use of modified Edman reagents such as 4 N N dimethylaminoazobenzene 4 isothiocyanate DABITC for manual sequencing by Chang et al 5 in 1976 A second major breakthrough in the analysis of polypeptides was automated amino acid analysis described by Spackman et al in 1958 6 However during the period from 1975 to 1980 it became increasingly clear that the amount of material required for structural analysis was more than could be easily isolated for the vast majority of proteins The field was criticized for its lack of sensitive techniques for the analysis of growth factors immune modulators membrane receptors and peptide hormones In addition very little had been done to modernize and improve the original instruments introduced in the mid 1960s The first indications of improved instrumentation for Edman chemistry came from Wittmann Liebold's laboratory 7 followed by the introduction of a micro sequencer by Hunkapiller and Hood in 1978 8 The movement toward improved instrumentation culminated in the gas Phase sequencer of Hewick et al 9 in 1981

Biomedical Applications of Mass Spectrometry Clarence H. Suelter, J. Throck Watson, 2009-09-25 Biomedical Applications of Mass Spectrometry Edited by Clarence H Suelter and J Throck Watson This unusual text is not simply a compilation of mass spectrometric methods but provides instead insight into specific approaches mass spectroscopists use when applying the technique to a variety of biological problems Each chapter provides guidance in using the appropriate methods for isolating and purifying the compound class prior to mass spectrometric analysis Covered in depth are the mass spectrometry of carbohydrates peptide sequencing by mass spectrometry mass spectrometry of nucleic acid components and mass spectrometry in pharmacology This definitive look at a growing facet of the science is an essential reference for biochemists biological chemists bioanalytical chemists and students 1990 0 471 61303 7 396 pp **Techniques in Protein Modification** Roger L. Lundblad, 1994-11-17 This exciting laboratory support book encompasses the encyclopedic coverage of the most frequently used methods for the site specific chemical modification of proteins In a concise manner it presents

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