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Biophysical Study of X-ray Scattering in Biological Samples

Presented by

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To
My Father and
Mother,

To
My Brothers,

To
My Sisters

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Abstract

Abstract

Wide angle x-ray scattering (WAXS) from protein in solution and dry (lyophilized) proteins is characterized by the presence of two weak broad scattering peaks. The first peak corresponds to a d-spacing around 10 Å and is attributed to inter-helix packing. The second peak is at about 4.5 Å and is attributed to α -helix backbone. For a protein with β -sheets as the main secondary structure, these peaks correspond to inter-sheet packing and hydrogen bonding distance between β -strands, respectively.

Since the scattering from water dominates over the scattering from protein in solution, one have to use a highly intense x-ray beam (probably from synchrotron) in order to be able to observe the scattering from protein. Otherwise, it would still be possible to unveil the protein scattering peaks through the lyophilization of protein solution into a powder form and obtaining the WAXS profile using a conventional x-ray diffractometer. The latter method is adopted in this study.

Since it has been reported that wide-angle x-ray scattering would provide a means to identify induced changes in the secondary, tertiary and quaternary structure of protein, this work aims to evaluate the potential of WAXS as a probe of induced conformational changes in insulin. For such purpose, native (control) insulin is forced to unfold and breakdown either using thermal denaturation alone or in the presence of thiol (cysteine) catalysts via disulfide scrambling. Denatured products are acid-trapped and monitored using WAXS in addition to FTIR, gel

filtration chromatography and Transmission Electron Microscopy (TEM) as supportive techniques.

Results show that the WAXS peak at 10 Å is sensitive towards the α -helix content of insulin. A reduction in the intensity of such peak (due to thermal denaturation or denaturation using thiol catalysts) is proven to be directly linked to the reduction of native insulin having normal α -helix content. It has been shown that the decrease in α -helix content of insulin is accompanied by the appearance of β -sheet structures that in turn rearrange into a fibrous form (amyloid fibers).

Chapter 1

Introduction

1. Introduction:

Since the discovery of x-rays by Roentgen in 1895, wide range of applications in the biological and medical fields had been developed. Among these applications are x-ray radiography, x-ray crystallography, x-ray florescence analysis, x-ray computed radiography, Dual Energy X-ray Absorptiometry (DEXA) and possibly many other useful developments to come in the future.

Among the recent interesting applications of x-ray in the biological and medical fields are the x-ray scattering techniques. These techniques make use of photons scattered from biological entities to gain characteristic information about the scattering tissue or molecule.

While Compton scattering has shown remarkable sensitivity toward the density of the scattering medium, coherent (elastic or Rayleigh) scattering has shown a potential sensitivity toward the molecular structure of biomolecules. This is a major interest in the present study.

Kosanetzky et al (1987) presented x-ray diffraction patterns of some plastics and several biological samples. Their scattering profiles showed that the investigated biological samples had characteristic scattering distribution with one or more forward scattering peaks. These peaks are attributed to the interference of photons coherently scattered from molecules of the medium (molecular interference effects) and considered as a fingerprint of the investigated sample.