

First Molecular Graphics and Modelling Society Electronic Conference

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Abstract

In this article we describe activities at the First Molecular Modelling and Graphics Society Electronic Conference (MGMS EC-1) which was held on the Internet and World Wide Web in October 1996.¹ MGMS EC-1 involved the presentation and discussion of scientific research results in a virtual conferencing environment which incorporated virtual replicas of many activities usually observed at a physical conference in addition to features unique to the electronic medium. Highlights of the scientific programme and technical developments in the design and use of these facilities are briefly described. A second electronic conference is planned for October 1997.²

Introduction

At the MGMS EC-1 event¹ presenters were able to display research results using a World Wide Web (WWW) presentation of text and graphics and could discuss the results within the context of an interactive software environment running on the Internet. Simply put, a scientist was able to sit at his/her computer and view presentations and exchange comments with other scientists situated in their own offices or labs in other parts of the world. A variety of activities was pos-

sible including the ability to manipulate molecular structures, to submit and review resumés, to review exhibitor product descriptions and to carry out online searches of the proceedings.

MGMS EC-1 was held on the Internet 7-18 October 1996¹ with entry to the site via an extensive set of WWW pages. The conference was sponsored by Elsevier, the Molecular Graphics and Modelling Society (MGMS), Oxford Molecular, MSI, Springer-Verlag, Wyeth-Ayerst, NASA-Ames and GlaxoWellcome. The event attracted close to a hundred presentations and over two hundred participants. The conference was served both from a UK location (The University of Oxford) and the US (S.U.N.Y. and Wesleyan University) with continuous real-time mirroring of the interactive environment. A similar event is planned for October 1997.² In this article we give an overview of the features, events and news of the first conference whereas a more detailed analysis of the software environment and its use is given elsewhere.³ A hypertext version of this article and accompanying full color graphics are available at <http://www.vei.co.uk/mgmsec1/article/>.

Overview

MGMS EC-1 was a fully international event open to all members of the scientific community and covered a broad range of disciplines related to molecular modelling, graphics and simulation methods and applications including section areas and conveners in the following topics:

Protein Structure

(Rod Hubbard, University of York, UK);

Membranes and Membrane Proteins

(Alan Robinson, University of Oxford, UK);

Computational Nanotechnology

(Al Globus, NASA-Ames, USA);

Protein Folding

(Jeffrey Skolnick, Scripps Institute, USA);

Modelling of In Vivo Activity

(Edward Hodgkin, Wyeth-Ayerst Research, USA);

Knowledge-based Library Design

(Mike Hann, GlaxoWellcome, UK);

Surface Science (Donald Brenner, North Carolina State, USA);

Chemical Design Automation News

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