UK and Republic of Ireland SIAM Section Annual Meeting
Held 6th January, 2006 in Cardiff.

This was the first annual meeting of the section held in Wales and

was very well attended with over 60 participants.

The meeting was hosted by the mathematics department at Cardiff University

and thanks to Marco Marletta who acted as a local organizer to the meeting.

The meeting was opened by the section president, Nick Trefethen, who gave

a brief overview of on the role of Europe, the UK and Republic of Ireland for SIAM.

The first speaker was Ben Leimkuhler (Leicester) who spoke on enhancing

thermal exchange in molecular dynamics. This work has applications in

biochemical systems and atomisitic materials models. He discussed dynamics

based approaches and numerical methods.

Hinke Osinga (Bristol) was the next speaker and she gave an entertaining talk

on computing global stable and unstable manifolds. The algorithm was described

and she showed it possible to construct the stable and unstable manifolds for

the Lorenz euqations, not only on a computer but also by crocheting. The talk

included an interview for Channel 4 news with Jon Snow.

The business meeting and an excellent lunch (supported by an anonymous donor) followed.

Peter Clarkson (Kent) then spoke on special polynomials associated with

(PI--PVI) are six nonlinear ordinary differential equations that have

been the subject of much interest in the past thirty years, which have

arisen in a variety of physical applications such as soliton equations

which are solvable by the inverse scattering method, including the

Korteweg-de Vries, modified Korteweg-de Vries, Boussinesq and nonlinear

Schr\"odinger equations.

Des Higham spoke on a new model for protein-protein interaction networks.

These are typically viewed as large unweighted, undirected graphs that, when

analyzed appropriately, may reveal important biological information.

Researchers have considered high-level questions, such as ``can we describe

these networks in terms of a few parameters?'' and low-level questions such

as ``can we identify interesting groups of proteins?''. He described a new

graph model that aims to contribute at both levels and presented results for

real biological data sets.

Marty Golubitsky rounded off an enjoyable day with a talk on coupled cell

models. These are a collection of interacting dynamical systems. Coupled

cell models assume that the output from each cell is important and that

signals from two or more cells can be compared so that patterns of synchrony

can emerge. He asked: How much of the qualitative dynamics observed in coupled

cells is the product of network architecture and how much depends on

the specific equations? The ideas were illustrated with a number of examples and theorems.

Business meeting :

The UKIE section president, Nick Trefethen, gave a brief overview of

the past year and reminded the members that the post of secretary/treasurer becomes vacant at the end of March 2005 and $\,$

nominations were solicited. The finances and activities funded in

the year including the student prizes presented were reported on.

The next annual meeting will be in January 2007, probably Jan 5, at

a venue to be decided although an offer was made from ${\tt Oxford}$ university.

Finally the SIAM President, Marty Golubitsky, reported from SIAM and in

particular called for participation for nodes on the web on "Why Do Maths ?".

Gabriel Lord