

VALPARAISO'S MATHEMATICS AND ITS APPLICATIONS DAYS

Séptimo Encuentro de la Matemática y sus
Aplicaciones

Instituto de Matemáticas, Pontificia Universidad Católica de
Valparaíso

Valparaíso, 12 y 13 de Enero de 2017

PROGRAMA

1. INTRODUCCIÓN

El **Séptimo Encuentro de la Matemática y sus Aplicaciones** ha sido organizado en conferencias secuenciales de **45 y 30 minutos** de duración (40 y 25 minutos de exposición, respectivamente, y 5 minutos para preguntas y comentarios). Todas las charlas se llevarán a cabo en la sala **IMA 2-3** en el segundo piso del Instituto de Matemáticas de la Pontificia Universidad Católica de Valparaíso.

En la siguiente página se detalla la programación correspondiente, incluyendo autor y título de la charla.

La organización agradece al Instituto de Matemáticas y al GEAGAM Network por su gran apoyo para llevar a cabo este evento. Igualmente, extiende su reconocimiento y gratitud a todos los expositores, quienes gracias a su buena voluntad de participar, han hecho posible la realización de este **VMAD 7**.

Comité Organizador

Valparaíso, Enero de 2017

2. JUEVES, 12 DE ENERO

10.10-10.15 BIENVENIDA

10.15-11.00 DAVID PARDO: *Refined Isogeometric Analysis (rIGA)*.

11.00-11.30 SERGIO ROJAS: *An efficient 1.5D Galerkin method for geophysics*.

[Moderador: M. Barrientos]

11.30-12.00 COFFEE BREAK

12.00-12.30 PAULINA SEPÚLVEDA: *A spacetime DPG method for the Schrodinger equation*.

12.30-13.00 TOMÁS BARRIOS: *Analyses of DG approximations for Stokes problem based on velocity-pseudostress formulation*.

[Moderador: I. Muga]

13.00-15.00 ALMUERZO

15.00-15.30 FERNANDO CÓRDOVA: *Impulsive differential equations with dynamical pulse times. Applications towards the biological systems*.

15.30-16.00 KARINA VILCHES: *Modelling the environmental influence in chemotaxis movements*.

16.00-16.30 PATRICIO CUMSILLE: *Classical Mathematical Modeling for Description and Prediction of Experimental Tumor Growth: the case of melanoma*.

[Moderador: D. Paredes]

16.30-17.00 COFFEE BREAK

17.00-17.30 CRISTÓBAL BERTOGLIO: *Estimation of pressure drop in blood flows from MRI*.

17.30-18.00 PABLO MOISSET de ESPANES: *A mathematical insight into intracellular iron regulation: the buffering behavior of ferritin is explained by large capacity and small exchanges*.

[Moderador: M. Barrientos]

3. VIERNES, 13 DE ENERO

10.15-11.00 RYAN McCLARREN: *High Fidelity, Moment-Based Methods for Particle Transport: The confluence of PDEs, Optimization, and HPC.*

11.00-11.30 CARLOS JERES-HANCKES: *Uncertainty Quantification for Electromagnetic Scattering by 1D PEC Gratings.*

[Moderador: D. Paredes]

11.30-12.00 COFFEE BREAK

12.00-12.30 PAUL ESCAPIL: *Wave Diffraction by Random Surfaces: Uncertainty Quantification via Sparse Tensor Boundary Elements.*

12.30-13.00 IGNACIA FIERRO: *Electromagnetic scattering at a telescope mirror.*

[Moderador: M. Barrientos]

13.00-15.00 ALMUERZO

15.00-15.30 ELWIN VAN'T WOUT: *High-performance computing of multiple wave scattering with boundary element methods*

15.30-16.00 CONSUELO FUENZALIDA: *Sparse Tensor Approximations and Smolyak Quadratures for Estimating Second-Order Statistical Moment for Neutron Flux.*

16.00-16.30 CAROLINA URZÚA-TORRES: *Optimal Operator Preconditioning for Boundary Elements on 3D screens.*

[Moderador: I. Muga]

16.30-17.00 COFFEE BREAK

17.00-17.30 GINO MONTECINOS: *A strategy for the treatment of Dirichlet boundary conditions in the context of ADER scheme for one-dimensional conservation laws.*

17.30-18.00 CRISTÓBAL CASTRO: *Numerical methods for hyperbolic equation. Applications to shallow water and linear elasticity.*

[Moderador: M. Barrientos]

20:30 CENA DE CAMARADERIA

Analyses of DG approximations for Stokes problem based on velocity-pseudostress formulation

TOMÁS P. BARRIOS * ROMMEL BUSTINZA † FELIPE SÁNCHEZ ‡

Abstract

In the present paper we propose two families of local discontinuous Galerkin methods for the numerical approximation of Stokes equations in their (nonsymmetric) velocity-pseudostress formulation. Because of a characterisation of the discrete kernel of the relevant bilinear form is not straightforward in the present setting, the well-posedness of the first kind of problems could follow from the Babuska-Brezzi theory only in the case of high-order approximations. For low-order methods (which may be more attractive in view of the associated computational complexity of tensor field-based discretisations), one Raviart-Thomas projection in combination with the Fredholm alternative allows to conclude the bijectivity of the solution operator. Error estimates are then derived, always under the assumption that the divergence of the approximation space of stresses is contained in the discrete velocity space, and that the gradient of the velocity space is contained in the pseudo-stress approximation space. Next, we relax these last restrictions by proposing two stabilised schemes: one incorporating a div-div term, and a second one bearing a Galerkin least-squares residual. The analysis of the stabilised methods is also presented, and the behaviour of the three schemes are illustrated via a few numerical examples.

References

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Estimation of pressure drop in blood flows from MRI

CRISTÓBAL BERTOGLIO* DAVID NOLTE AXEL OSSES

Abstract

Blood flow velocity, hemodynamic pressure gradients, arterial stiffness, etc are very important clinical indexes of the cardiovascular condition. To date, clinical gold standard are invasive procedures or imaging techniques requiring exogenous contrast agents or x-rays. Recent advancements in inverse problems and numerical hemodynamics have shown their potential to use coupled fluid-solid models and medical images in order estimate non-invasively hemodynamic quantities from medical images [1, 2, 3].

In this talk we will focus on strategies to estimate pressure gradients from velocity data. We will detail two approaches: (a) efficient methods for the pressure gradient computation assuming the velocity is measured everywhere in space (at a certain spatial resolution), as it can be obtained by 3D Phase-Contrast MRI [4], and (b) an optimization-based approach with the incompressible Navier-Stokes equations as constraints, that may allow a reduced amount of data (e.g. 2D Phase-Contrast MRI).

References

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Numerical methods for hyperbolic equations. Applications to shallow water and linear elasticity

CRISTÓBAL CASTRO *

Abstract

One of the main characteristic of hyperbolic equations is the development of discontinuous solution even for smooth initial data. In order to address this condition a robust and accurate Riemann solver is necessary. In most cases, the exact solution is not available or not desirable due to computational complexity. If we consider Euler equations as example, there are number of approximate Riemann solver in the literature [3]. In this presentation we show recent developments of numerical method for solving hyperbolic equations. In particular we present an approximate, general equation of state, Riemann solver for Euler equations [4]. Another interesting aspect of hyperbolic equations is the wave propagation capability. In geophysical problems the propagation of small amplitude for long distance and time pose a mayor challenge. We show applications of tsunami [2] and seismic wave propagation [1].

References

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V-MAD 7

Séptimo Encuentro en Aplicaciones de la Matemática

Instituto de Matemáticas, Pontificia Universidad Católica de Valparaíso, 12 y 13 Enero, 2017

Impulsive differential equations with dynamical pulse times. Applications towardas the biological systems.

FERNANDO CÓRDOVA-LEPE *

Abstract

A type of impulsive differential equations (IDE) with dynamical pulse times is presented. These equations were introduced by Córdova-Lepe et al in the year 2012, [1]. In addition, some applications in mathematical modeling of biological systems are given. More precisely, some models in epidemiology (pulse vaccination), bioeconomics (pulse fishery) and bioprocess (pulse control) are discussed, see [2, 3, 4].

References

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Classical Mathematical Modeling for Description and Prediction of Experimental Tumor Growth: the case of melanoma

PATRICIO CUMSILLE * ANDRÉS RODRÍGUEZ †

Abstract

In this work we study classical mathematical models of tumor growth (see [2, 5]) in order to describe and predict melanoma development (melanoma is the most serious type of skin cancer). To do so, firstly an experimental methodology will be carry out: human melanoma cells will be grown in a culture medium RPMI-1640 with a pH of 7.2, supplemented with serum at 5% in order to stimulate division. Once confluent, cells will be suspended in a sterile medium. Next, 200,000 of these melanoma cells will be injected in the skin of two group of mice, from which daily growth will be observed starting from the 7th day post injection, by using a Caliper, measuring the length and the width of melanoma tumor for each group, control and experimental, data which will be used in order to make parameters estimation of mentioned models, thereby describing and forecasting tumor growth. Control group is composed by wild-type mice, whereas experimental group consists of mice to which Adenosine receptor of type A2 has been knocked out, with the aim to investigate the influence of Adenosine in tumor growth, since this proteine has a role as an angiogenesis modulator by enhancing vascular endothelial growth factor (VEGF)¹ activity (see [1]).

The numerical methodology required in order to make parameters estimation of the classical mathematical models, based upon ordinary differential equations (ODE), consists in employing Trust-Region-Reflective Algorithm (see [3]), a non linear optimization method specially designed to solve parameters estimation problems by non linear least squares criterion. This algorithm is implemented in Matlab© through lsqnonlin function (see [7]). In order to use it, we will give as input data of this function: the experimental data (the length and the width measured at certain times, in days), the ODE solution, if an explicit formula is available (in the case where this is not available, we use the ode45 solver of Matlab© in order to numerically solve the ODE for each

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¹VEGF is the most important pro-angiogenic molecule triggered under hypoxic conditions.

model), an initial guess of the optimal value of the parameter vector (which in general is unknown, but exploring the relative square error, in evaluating it as a function of parameters space, we achieve an initial guess to start the optimization procedure), the Jacobian matrix of the residuals (when an explicit solution of the ODE model is available), and lower and upper bounds for the parameters (obtained also by the exploratory method mentioned before). Next, we will apply statistical methods based upon non linear least squares regression tools (see [4, 6]) in order to assess goodness of fit for parameters estimated, as well as, performance of prediction to each model studied.

The expected results of this work are: 1) establish the descriptive power of each model, using several goodness-of-fit metrics and a study of parametric identifiability, 2) assess the models ability to forecast tumor growth, this for both groups of individuals, under experimental conditions described before, and 3) conclude that both groups come from different populations, that is to say, the fact of knocking out the type A2 Adenosine receptor influences tumor growth, which will be achieved through validation of mathematical models with experimental data.

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Wave Diffraction by Random Surfaces: Uncertainty Quantification via Sparse Tensor Boundary Elements

PAUL ESCAPIL-INCHAUSPÉ ^{*} CARLOS JEREZ-HANCKES [†]

Abstract

We consider the numerical solution of time-harmonic scattering of acoustic and electromagnetic waves from obstacles with uncertain geometries. Using first-order shape derivatives, we derive deterministic boundary integral equations for the mean field and the two-point correlation function of the random solution for a soft-obstacle Dirichlet problem. Sparse tensor Galerkin discretizations of these equations are implemented with the so-called *combination technique*. Similar discretization errors for the covariance is achieved with $\mathcal{O}(N \log N)$ degrees of freedom instead of $\mathcal{O}(N^2)$. Performance comparison of our approach to classic Monte-Carlo Galerkin formulation is given for different shapes. Finally, we verify the robustness of the sparse tensor approximation and compare it to low-rank approximations techniques.

References

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V-MAD 7

Séptimo Encuentro en Aplicaciones de la Matemática

Instituto de Matemáticas, Pontificia Universidad Católica de Valparaíso, 12 y 13 Enero, 2017

Electromagnetic scattering at a telescope mirror

MARÍA IGNACIA FIERRO PICCARDO *

Abstract

This presentation is part of a ModEM project, which has as a main objective finding out out how defects on a telescope mirror could cause noise its observations. The hipotesis is that gaps on a telescope surface produce changes on the polarizarion of the field scattered by the mirror. This problem is usually attacked by high frequency methods, as Physical Optics (PO) or Physical Theory of Diffraction (PTD), but in order to validate the results obtained by those methods, we propose to use the Boundary Elements Method on a common problem with the PO/PTD methods. For this reason, in this presenttation we show some of the preliminar results that will allow us to decide about the effect of the defects on the telescope mirror and, in a future, to validate or refute the results obtained by PO/PTD.

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Sparse Tensor Approximations and Smolyak Quadratures for Estimating Second-Order Statistical Moment for Neutron Flux

MARÍA CONSUELO FUENZALIDA * CARLOS JEREZ * RYAN McCLARREN †

Abstract

In this work, we develop a novel method to compute the second order statistical moment of neutron flux inside a reactor by solving the neutron transport equation for its moments. Randomness comes from the lack of precise knowledge of external sources as well as of cross-section parameters. Thus, the flux is itself a random variable and we are interested in computing its moments directly instead of using Monte-Carlo simulations. This approach, however, entails the appearance of the so-called curse of dimensionality. By assuming as given both the second moment of sources and probability distributions of cross-section parameters, we present an efficient method based on a sparse tensor finite-element method approximation as well as the use of Smolyak quadratures.

References

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Uncertainty Quantification for Electromagnetic Scattering by 1D PEC Gratings

R. AYLWIN* C. JEREZ-HANCKES * G. SILVA *,† P. FAY †

Abstract

We present a deterministic numerical method able to calculate efficiently first and second statistical moments of the scattered field by a periodic perfect electric conductor surface with stochastic perturbations on its surface. The electric field integral equation (EFIE) was solved using boundary elements method (BEM), and constant hierarchical bases, called Haar wavelets. To validate the BEM implementation the method was compared to COMSOL Multiphysics. On the other hand, to validate the stochastic calculations, the algorithm was compared to the Monte-Carlo simulation, obtaining good agreement in both cases. The proposed deterministic approach converges faster than Monte-Carlo simulation with $O(N \log N)$ computational effort.

References

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V-MAD 7

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High Fidelity, Moment-Based Methods for Particle Transport: The confluence of PDEs, Optimization, and HPC

RYAN G. McCLARREN *

Abstract

The calculation of the transport of particles is important in many applications including rarefied gas dynamics, plasma physics, and the nuclear energy systems. In this talk I will motivate the choice of moment-based methods for solving particle transport problems, and discuss the difficulties such approaches have. To obtain physically-meaningful solutions the discretization of the original integro-differential equation can depend on the solution to an optimization problem. I will show how these optimization problems arise, what methods perform best in terms of cost and accuracy, and how these problems can be well-suited for high performance computing.

References

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V-MAD 7

Séptimo Encuentro en Aplicaciones de la Matemática

Instituto de Matemáticas, Pontificia Universidad Católica de Valparaíso, 12 y 13 Enero, 2017

A mathematical insight into intracellular iron regulation: the buffering behavior of ferritin is explained by large capacity and small exchanges

P. MOISSET DE ESPANS ^{*} A. OLIVERA-NAPPA [†] J. SOTO [‡]

Abstract

Ferritin is a highly conserved ubiquitous iron-storage protein present in archaea, bacteria and eukaryotic cells and is also thought to be the major iron concentration buffer inside cells. Ferritin is a hetero-24-mer formed by two different peptide chains that assemble in a hollow sphere multimeric structure. This molecular structure is strikingly different from all other intracellular metal ion concentration buffers and determines two unique ferritin functional features: ferritin fer- rooxidase activity sequesters soluble Fe^{2+} ions as insoluble Fe^{3+} oxides inside the protein cavity, where they remain isolated from the external medium, and they can only be exchanged with the exterior through small channels that allow only a few iron atoms to pass simultaneously. The latter is exacerbated by the enzymatic activity itself, which imposes a kinetic limiting step on the amount of iron atoms that can be processed in a given period, in stark contrast with most common intracellular buffers that do not possess catalytic activity. Using a simple model based on mass action kinetics with minimal mechanistic hypothesis, we show that the isolation of iron atoms inside the protein and the exchange of few atoms at a time are sufficient to qualitatively determine ferritins dynamic behavior and to satisfy the stringent free iron intracellular concentration requirements. The main characteristics of this control profile are an almost constant free iron 1profile for a broad range of total iron concentrations, a single and globally stable equilibrium, a fast control velocity when faced to total iron concentration perturbations and an energy-efficient operation. This sophisticated and robust control response is radically different from those of other intra and extracellular buffer systems and has been achieved by using a surprisingly simple structurally and catalytically based strategy that do not require feedback control loops or sensing devices.

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A strategy for the treatment of Dirichlet boundary conditions in the context of ADER scheme for one-dimensional conservation laws

G. I. MONTECINOS *

Abstract

ADER schemes are high-order numerical methods, which nowadays are very popular. The so called reconstruction procedure is one of their building block, see [1] for details. ADER method for general conservation laws, may lose accuracy if boundary conditions are not deal properly. In this work the treatment of Dirichlet boundary conditions, is concerned. The reconstruction procedure, near boundaries, demands for information outside the computational domain, which is carried out in terms of numerical solutions of auxiliary problems. These problems are hyperbolic and they are constructed from the conservation laws and the information at boundaries. The evolution of these problems, unlike to the usual manner, is done in space rather than in time due to that, these are named, *reverse problems*. The methodology can be seen as a numerical version of the inverse Lax-Wendroff procedure, [2]. Expected orders of accuracy for solving conservation laws by using the proposed strategy at boundaries, are obtained up to fifth-order in both space and time, [3] for details.

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Séptimo Encuentro en Aplicaciones de la Matemática

Instituto de Matemáticas, Pontificia Universidad Católica de Valparaíso, 12 y 13 Enero, 2017

Refined Isogeometric Analysis (rIGA)

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MACIEJ PASZYŃSKI § NATHAN COLLIER ¶ VICTOR CALO ||

Abstract

Isogeometric Analysis (IGA) is a discretization method commonly employed nowadays to solve numerical problems governed by partial differential equations (PDEs). This method employs Computed-Aided Design (CAD) functions as basis functions to build the discretization of the governing PDEs [1]. This choice avoids to define a secondary set of functions for the engineering analysis, which drastically reduces the time and implementation efforts of the pre-processing step. In addition, highly continuous CAD functions provide better approximation properties than traditional Finite Element method on a per-degree of freedom basis, suggesting IGA is an accurate and robust scheme to approximate the solution of PDEs [2, 3]. Unfortunately, a high degree in the continuity also degrades the performance of direct solvers, increasing the computational cost per degree of freedom, resulting in larger solution time and memory requirements [4].

In this work, we propose an approach to dramatically reduce the computational time employed to approximate the PDEs solution when using highly continuous IGA discretizations. This method adjusts locally the continuity of the basis functions according to how the solver of linear equations operates. Starting from a highly continuous C^{p-1} IGA, the method enriches the space by decreasing the continuity over certain hyperplanes. These hyperplanes correspond to: (a) the separators used during the elimination of the degrees of freedom (dof) when using direct solvers, and (b) the skeleton obtained after static condensation of certain macro-element unknowns for the case of iterative solvers. We denote this method as “refined Isogeometric Analysis” or rIGA [5].

Despite the fact that rIGA spaces are richer than those coming from highly continuous C^{p-1} IGA systems, the computational time required to solve the resulting system of linear equations for a given mesh and fixed polynomial order is reduced both in the

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cases of direct and iterative solvers with respect to their highly continuous C^{p-1} IGA counterpart.

For the case of direct solvers, this reduction factor is approximately p^2 , being p the polynomial order. For instance, in a $2D$ quadrilateral mesh with four million elements and $p = 5$, the system resulting from rIGA is solved in almost *one* minute, while the system obtained from IGA requires approximately 22 minutes to be solved. In $3D$, when the mesh size is two million elements and $p = 3$, the rIGA system is solved in *one* hour, while the system obtained from IGA requires almost 15 hours to be solved.

For the case of iterative solvers, the savings of rIGA with respect to C^{p-1} IGA are more moderate and show an increment as the polynomial degree grows. For a mesh with four million elements and polynomial degree $p = 9$, rIGA delivers a reduction factor of approximately 3 with respect to C^{p-1} IGA.

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Séptimo Encuentro en Aplicaciones de la Matemática

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An efficient 1.5D Galerkin method for geophysics.

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Abstract

To characterize hydrocarbon (oil and gas) bearing formations, usually maps of the Earth subsurface are estimated by interpreting resistivity measurements obtained from various electromagnetic impulses through the media. In logging-while-drilling (LWD) operations, it is required to estimate the resistivities ideally in real (logging) time (see [2, 5]).

In multiple reservoirs, it is customary to assume homogeneous materials along two spatial directions, making possible the dimensional reduction of the problem through a Hankel transform, obtaining a so called 1.5D formulation. Under this assumption, semi-analytical methods can be employed to solve a single forward problem in a fraction of a second (see [1, 3, 8], and more recently [4, 6, 7, 9]). However, these methods have several limitations, including: (a) they can only consider piecewise constant materials, and (b) it is difficult (and sometimes impossible) to obtain semi-analytical derivatives, forcing the use of expensive finite difference schemes. When considering Galerkin-type formulations for the 1.5D problem, more general models can be considered. Moreover, accurate approximations of the derivatives can be obtained through the resolution of an adjoint problem. Unfortunately, standard Galerkin methods are relatively slow when compared to the semi-analytical solution, as it occurs in practical applications.

In this talk, we present a novel Hankel-Galerkin multi-scale method that drastically reduces the total computational time. The method employs multi-scale basis functions. The resulting procedure is computationally expensive when comparing with a single forward resolution; however, it becomes competitive when solving multiple forward problems. In the presentation, we introduce the method for the acoustic and electromagnetic cases, and show numerical results validating its accuracy.

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Séptimo Encuentro en Aplicaciones de la Matemática

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A spacetime DPG method for the Schrödinger equation

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SRIRAM NAGARAJ [‡] PAULINA SEPÚLVEDA [§]

Abstract

The aim of this talk is to describe a spacetime Discontinuous Petrov Galerkin (DPG) method for the linear time-dependent Schrödinger equation. We show that it is possible to lose solutions if the second order Schrödinger equation is reformulated into a first order formulation. This makes spacetime methods based on a second order formulation particularly attractive for capturing irregular solutions. We show some numerical experiments motivated by pulse propagation in dispersive optical fibers.

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Optimal Operator Preconditioning for Boundary Elements on 3D screens

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Abstract

The discretization of first-kind boundary integral operators (BIOs) by low-order Galerkin BEM leads to ill-conditioned linear systems on fine meshes. Consequently, iterative solvers become prohibitively slow and preconditioning is necessary.

The standard “Calderón preconditioning” technique breaks down when considering open boundaries as when modeling screens. In this case, the double layer operator and its adjoint vanish. Moreover, the associated weakly singular and hypersingular operators no longer map fractional Sobolev spaces in a dual fashion.

We propose new Calderón-type preconditioners for the hypersingular and weakly singular operators arising from the Laplacian on screens. For their construction, we use operator preconditioning [1] and the bilinear forms induced by their recently found inverse BIOs over the disk [2, 4].

In addition, we are able to apply our preconditioning technique to non-uniform meshes [3]. This property poses a great advantage, as solutions of boundary integral equations on screens Γ feature a square-root type singularity at $\partial\Gamma$, which can be resolved by refining towards the boundary.

Numerical examples illustrate the optimality of our preconditioners when applied to different screens.

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Séptimo Encuentro en Aplicaciones de la Matemática

Instituto de Matemáticas, Pontificia Universidad Católica de Valparaíso, 12 y 13 Enero, 2017

Modelling the environmental influence in chemotaxis movements

KARINA VILCHES PONCE *

Abstract

In this presentation, we review the state of the art about the incorporation of the environmental influence in the chemotaxis classical modelling. The mathematical results obtained by different authors will be summarized and the remaining challenges in this subject will be given.

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High-performance computing of multiple wave scattering with boundary element methods

ELWIN VAN 'T WOUT *

Abstract

Computational simulation of wave scattering at multiple objects is of interest to different areas in engineering and physics. Two such applications are the prediction of acoustic scattering at rib cages for the planning of medical treatment procedures and the electromagnetic scattering at telescope mirrors for astronomical signal processing. Furthermore, both problems require the simulation of high-frequency wave fields at large-scale objects. Since the geometry is embedded in free space, the Boundary Element Method (BEM) is the prime choice of numerical algorithm. Because of the surface integral representation, the number of degrees of freedom scales favourably compared to volumetric methods. However, solving the dense set of linear equations poses severe limitations on the maximum frequency that can be used on present-day computing platforms. Here, a combination of parallelisation, preconditioning, and compression techniques will be used to achieve large-scale BEM simulations.

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