The Process of Computational Science

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My approach to Computational Science is

Holistic

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Holistic

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starting with the numerics of PDEs, and mathematics of the computation,

through the distillation into high quality numerical libraries,

to scientific discovery through computing.

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Community Involvement



Outline

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Collaborators

BIBEE Researchers



Classical DFT Researchers



Dirk Gillespie



Bob Eisenberg

Bioelectrostatics



Induced Surface Charge on Lysezyme

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Bioelectrostatics Physical Model



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Bioelectrostatics Mathematical Model

We can write a Boundary Integral Equation (BIE) for the induced surface charge σ ,

$$\sigma(\vec{r}) + \hat{\epsilon} \int_{\Gamma} \frac{\partial}{\partial n(\vec{r})} \frac{\sigma(\vec{r}') d^2 \vec{r}'}{4\pi ||\vec{r} - \vec{r}'||} = -\hat{\epsilon} \sum_{k=1}^{Q} \frac{\partial}{\partial n(\vec{r})} \frac{q_k}{4\pi ||\vec{r} - \vec{r}_k||} (\mathcal{I} + \hat{\epsilon} \mathcal{D}^*) \sigma(\vec{r}) =$$

where we define

$$\hat{\epsilon} = 2\frac{\epsilon_I - \epsilon_{II}}{\epsilon_I + \epsilon_{II}} < 0$$

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Bioelectrostatics

Mathematical Model

The *reaction* potential is given by

$$\phi^{R}(ec{r})=\int_{\Gamma}rac{\sigma(ec{r}')d^{2}ec{r}'}{4\pi\epsilon_{1}||ec{r}-ec{r}'||}=C\sigma$$

which defines Ges, the electrostatic part of the solvation free energy

$$egin{aligned} \Delta G_{es} &= rac{1}{2} \left\langle q, \phi^R
ight
angle \ &= rac{1}{2} \left\langle q, Lq
ight
angle \ &= rac{1}{2} \left\langle q, CA^{-1}Bq
ight
angle \end{aligned}$$

where

$$egin{aligned} & m{B}m{q} = -\hat{\epsilon}\int_{\Omega}rac{\partial}{\partialm{n}(ec{r})}rac{m{q}(ec{r}')m{d}^3ec{r}'}{4\pi|ec{r}-ec{r}'||} \ & m{A}\sigma = \mathcal{I} + \hat{\epsilon}\mathcal{D}^* \end{aligned}$$

- Boundary element discretizations of the solvation problem (Eq. ??):
 - can be expensive to solve
 - are more accurate than required by intermediate design iterations

BIBEE Approximate \mathcal{D}^* by a diagonal operator

Boundary Integral-Based Electrostatics Estimation

Coulomb Field Approximation: uniform normal field

$$\left(1-rac{\hat{\epsilon}}{2}
ight)\sigma_{CFA}=Bq$$

Lower Bound: no good physical motivation

$$\left(1+rac{\hat{\epsilon}}{2}
ight)\sigma_{LB}=Bq$$

Eigenvectors: BEM $e_i \cdot e_i$ BIBEE/P



BIBEE Approximate \mathcal{D}^* by a diagonal operator

Boundary Integral-Based Electrostatics Estimation

Coulomb Field Approximation: uniform normal field

$$\left(1-rac{\hat{\epsilon}}{2}
ight)\sigma_{CFA}=Bq$$

Preconditioning: consider only local effects

$$\sigma_P = Bq$$

Eigenvectors: BEM $e_i \cdot e_i$ BIBEE/P



BIBEE Bounds on Solvation Energy

Theorem: The electrostatic solvation energy ΔG_{es} has upper and lower bounds given by

$$\frac{1}{2}\left(1+\frac{\hat{\epsilon}}{2}\right)^{-1}\langle q, CBq\rangle \leq \frac{1}{2}\left\langle q, CA^{-1}Bq\right\rangle \leq \frac{1}{2}\left(1-\frac{\hat{\epsilon}}{2}\right)^{-1}\langle q, CBq\rangle,$$

and for spheres and prolate spheroids, we have the improved lower bound,

$$\frac{1}{2}\left\langle q,CBq\right\rangle \leq\frac{1}{2}\left\langle q,CA^{-1}Bq\right\rangle ,$$

and we note that

$$|\hat{\epsilon}| < \frac{1}{2}.$$

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Energy Bounds:

Proof: Bardhan, Knepley, Anitescu, JCP, 130(10), 2008

I will break the proof into three steps,

- Replace C with B
- Symmetrization
- Eigendecomposition

shown in the following slides.

We will need the single layer operator S for step 1,

$$\mathcal{S}\tau(\vec{r}) = \int \frac{\tau(\vec{r}')d^2\vec{r}'}{4\pi||\vec{r}-\vec{r}'||}$$

Energy Bounds: First Step Replace *C* with *B*

The potential at the boundary Γ given by

$$\phi^{\textit{Coulomb}}(\vec{r}) = C^T q$$

can also be obtained by solving an exterior Neumann problem for τ ,

$$\phi^{Coulomb}(\vec{r}) = S\tau$$

= $S(\mathcal{I} - 2\mathcal{D}^*)^{-1}(\frac{2}{\hat{\epsilon}}Bq)$
= $\frac{2}{\hat{\epsilon}}S(\mathcal{I} - 2\mathcal{D}^*)^{-1}Bq$

so that the solvation energy is given by

$$\frac{1}{2}\left\langle q, \mathcal{C} A^{-1} \mathcal{B} q \right\rangle = \frac{1}{\hat{\epsilon}} \left\langle \mathcal{S} (\mathcal{I} - 2\mathcal{D}^*)^{-1} \mathcal{B} q, (\mathcal{I} + \hat{\epsilon} \mathcal{D}^*)^{-1} \mathcal{B} q \right\rangle$$

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Operator Approximation

Energy Bounds: Second Step Quasi-Hermiticity

Plemelj's symmetrization principle holds that

$$\mathcal{SD}^* = \mathcal{DS}$$

and we have

$$\mathcal{S}=\mathcal{S}^{1/2}\mathcal{S}^{1/2}$$

which means that we can define a Hermitian operator H similar to \mathcal{D}^*

$$H = \mathcal{S}^{1/2} \mathcal{D}^* \mathcal{S}^{-1/2}$$

leading to an energy

$$\frac{1}{2}\left\langle q, \mathcal{C} \mathcal{A}^{-1} \mathcal{B} q \right\rangle = \frac{1}{\hat{\epsilon}} \left\langle \mathcal{B} q, \mathcal{S}^{1/2} (\mathcal{I} - 2\mathcal{H})^{-1} (\mathcal{I} + \hat{\epsilon} \mathcal{H})^{-1} \mathcal{S}^{1/2} \mathcal{B} q \right\rangle$$

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Energy Bounds: Third Step Eigendecomposition

The spectrum of \mathcal{D}^* is in $\left[-\frac{1}{2}, \frac{1}{2}\right)$, and the energy is

$$\frac{1}{2}\left\langle q, CA^{-1}Bq\right\rangle = \sum_{i}\frac{1}{\hat{\epsilon}}\left(1-2\lambda_{i}\right)^{-1}\left(1+\hat{\epsilon}\lambda_{i}\right)^{-1}x_{i}^{2}$$

where

$$H = V \wedge V^T$$

and

 $\vec{x} = V^T \mathcal{S}^{1/2} B q$

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Energy Bounds: Diagonal Approximations

The BIBEE approximations yield the following bounds

$$\frac{1}{2} \left\langle q, CA_{CFA}^{-1} Bq \right\rangle = \sum_{i} \frac{1}{\hat{\epsilon}} \left(1 - 2\lambda_{i}\right)^{-1} \left(1 - \frac{\hat{\epsilon}}{2}\right)^{-1} x_{i}^{2}$$
$$\frac{1}{2} \left\langle q, CA_{P}^{-1} Bq \right\rangle = \sum_{i} \frac{1}{\hat{\epsilon}} \left(1 - 2\lambda_{i}\right)^{-1} x_{i}^{2}$$
$$\frac{1}{2} \left\langle q, CA_{LB}^{-1} Bq \right\rangle = \sum_{i} \frac{1}{\hat{\epsilon}} \left(1 - 2\lambda_{i}\right)^{-1} \left(1 + \frac{\hat{\epsilon}}{2}\right)^{-1} x_{i}^{2}$$

where we note that

$$|\hat{\epsilon}| < \frac{1}{2}$$

BIBEE Accuracy

Electrostatic solvation free energies of met-enkephalin structures



Snapshots taken from a 500-ps MD simulation at 10-ps intervals. Bardhan, Knepley, Anitescu, JCP, 2009.

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BIBEE Scalability



Yokota, Bardhan, Knepley, Barba, Hamada, CPC, 2011.

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Boundary element discretizations of the solvation problem:

- can be expensive to solve
 - Bounding the electrostatic free energies associated with linear continuum models of molecular solvation, JCP, 2009
- are more accurate than required by intermediate design iterations
 - Accuracy is not tunable

Evolution of BIBEE

- Sharp bounds for solvation energy
- Exploration of behavior in simplified geometries
 - Mathematical Analysis of the BIBEE Approximation for Molecular Solvation: Exact Results for Spherical Inclusions, JCP, 2011
 - Represent BIBEE as a deformed boundary condition
 - Fully developed series solution
 - Improve accuracy by combining CFA and P approximations
- Application to protein-ligand binding
 - Analysis of fast boundary-integral approximations for modeling electrostatic contributions of molecular binding, Molecular-Based Mathematical Biology, 2013

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Future of **BIBEE**

- Framework for systematic exploration
 - Both analytical and computational foundation
- Reduced-basis Method with analytic solutions
 - Tested in protein binding paper above
 - The spatial high frequency part is handled by BIBEE/P topology is not important
 - The spatial low frequency part is handled by analytic solutions insensitive to bumpiness
 - Computational science and re-discovery: open-source implementations of ellipsoidal harmonics for problems in potential theory, CSD, 2012.
- Extend to other kernels, e.g. Yukawa
- Extend to full multilevel method

Outline

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Collaborators

PETSc Developers





Barry Smith



Jed Brown



Andy Terrel



Peter Brune

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Problem

Traditional PDE codes cannot:

Compare different discretizations

- Different orders, finite elements
- finite volume vs. finite element

• Compare different mesh types

- Simplicial, hexahedral, polyhedral, octree
- Run 1D, 2D, and 3D problems

Enable an optimal solver

• Fields, auxiliary operators

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Problem

Traditional Mesh/Solver Interface is Too General:

- Solver not told about discretization data, e.g. fields
- Cannot take advantage of problem structure
 - blocking
 - saddle point structure
- Cannot use auxiliary data
 - Eigen-estimates
 - null spaces

Problem

Traditional Mesh/Solver Interface is Too Specific:

- Assembly code specialized to each discretization
 - dimension
 - cell shape
 - approximation space
- Explicit references to element type
 - getVertices(faceID), getAdjacency(edgeID, VERTEX), getAdjacency(edgeID, dim = 0)
- No interface for transitive closure
 - Awkward nested loops to handle different dimensions

Mesh Representation

We represent each mesh as a Hasse Diagram:

- Can represent any CW complex
- Can be implemented as a Directed Acyclic Graph
- Reduces mesh information to a single *covering* relation
- Can discover dimension, since meshes are ranked posets

We use an abstract topological interface to organize traversals for:

- discretization integrals
- solver size determination
- computing communication patterns

Mesh geometry is treated as just another mesh function.

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Sample Meshes



Sample Meshes Optimized triangular mesh



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Sample Meshes Interpolated quadrilateral mesh



Sample Meshes Optimized quadrilateral mesh



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Sample Meshes Interpolated tetrahedral mesh



Mesh Abstraction

By abstracting on the key topological relations, the interface can be both concise and quite general

- Single relation
- Enables dimension-independent programming
- Dual is obtained by reversing arrows
- Can associate function(al)s with DAG points
- Dual operation gives the support of the function

Mesh Algorithms for PDE with Sieve I: Mesh Distribution, Knepley, Karpeev, Sci. Prog., 2009.

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Basic Operations Support





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