

# ***Tinker Software Workshop 2018***

**The University of Texas at Austin, June 2-4, 2017**

**All meetings are in BME 3.204**

( BME Building on Google map: <https://goo.gl/maps/DgoBYwGehqA2> )

## **● June 2nd, Saturday, (9:00-11:45 am & 13:30-16:30 pm)**

### **1) Code Bases & Programming Models (Ponder)**

*(Approximately 10 minute presentation per topic, plus discussion)*

**Topics: Current status of various code bases and what have been changed since the last meeting**

- a) Tinker: Ponder
- b) Tinker-HP: Piquemal
- c) Tinker-OpenMM: Ren
- d) OpenMM: Eastman
- e) Amber & PMEMD: Case
- f) CHARMM: Brooks
- g) FFX: Schnieders
- h) Q-Chem/EFP: Head-Gordon
- i) MolSSI Intro and Database project: Jessica Nash

### **2) Algorithms & Methodology (Schnieders)**

**Topics: Very brief description of the method, applications, and current & future implementations**

#### **a) Polarization**

- iELSCF: Head-Gordon
- OPT: Simmonett
- TCG and beyond: Aviat
- DC-JI/DIIS: Greg Beran or D. Nocito
- Simplification of “d” and “p”: Simmonett

#### **b) MD**

- Constant pH: Schnieders

Integrators: Lagardere  
Integrators: Tuckerman  
Best practice in implementing/choosing thermo & barostat

### **c) Sampling**

Sampling OSRW: Wei Yang  
Parallel-in-Time: Maday  
Accelerated MD: Eastman  
Steered MD: Piquemal

### **d) Models**

Implicit solvent MIB-PB: Guowei Wei  
ddCOSMO: Stamm  
GK: Schnieders

QM/MM: Cisneros  
QM/MM: Yihan Shan  
QM/MM: Daniele Loco (AMOEBA embedding and QM/MD)  
Filippo Lipparini (Gaussian interface)  
PME for dispersion: Andy Simmonett

Metal ions/Sammer Varma

Automated Parameterization: Chris Ho  
POLTYPE: Brandon Walker, torsion/implicit solvent  
Coarse-grained RNA model: Sara Cheng

### **e) New Force Fields**

Open Force Field: Michael Schaperl  
SIBFA: Piquemal/Gresh  
GEM: Cisneros  
MPID: Brooks  
AMOEBA2: Rackers/Ponder  
AMOEBA+: water model Chengwen/Ren

## **3) Development over the next year (Part A)**

### **a) Hardware (Piquemal)**

CPU Vectorization: AVX512 speedups (Jolly)  
CPUs/hybrid  
GPUs: NVIDIA vs. AMD, CUDA vs. OpenCL vs OpenACC  
Volta/Tensor core

~~IBM OpenPOWER~~ Something else? Swope/Rice

Programming Models: MPI 3.x, OpenMP 4.x  
Hybrid CPU / GPU Code Multi-Platform Support

**b) Software (Ponder)**

Tinker (C++ vs. Fortran)

Tinker-HP

Tinker-OpenMM

Amber

Charmm

QM/MM (Gaussian, Q-chem)

**Priorities for next year:**

Support for Current Versions in Tinker of Existing Models/Force Fields (Amber, CHARMM, OPLS, *etc.*)

Support for New and Emerging Force Fields, esp. Polarizable Methods Algorithmics & new Mathematics / Physics

Support for free energy methods

● **June 2<sup>nd</sup>, Saturday Dinner**

**(6:30 – 8:30 pm; dinner at the [AT&T center in room PDR 1&2.](#))**

**4) Discussion & Social Hour(s)**

Taco Bar 16

Smoked Brisket, Pork Carnitas, Tinga Chicken, Portobello Mushroom, Flour Tortillas, Lettuce, Tomato, Jalapeño, Cheddar, Sour Cream, Guacamole, Fire Roasted Salsa

● **Sunday, June 3<sup>rd</sup>**

**(9-11:45 am; 1:30-4:30 pm)**

**5) Development over next year (Part B) (Ponder/Ren/Piquemal)**

a) TINKER as the reference code; put new methods / code there first, and then move to TINKER-HP, OpenMM, *etc.*

- b) Integration of the three Tinker code bases (merge at "TINKER 9" ?) Multiple versions of polarizable electrostatics, or a single version – ?? Not happening
- c) Tinker 8, Tinker-HP, QI; Speed Comparisons Refactoring to enable MPI
- d) PCG and TCG vs. DIIS on GPUs
- e) Updated OSRW Versions; Lambda Derivatives,  $dU/d\lambda$ ; Soft-Core vdw & Electrostatics
- f) QI implementation of multipoles
- g) Refactoring to enable Domain Decomposition Tests of importance of dispersion PME (?)
- h) Stay in sync with "Canonical" OpenMM Tinker-OpenMM branch with periodic merges
- i) Develop new GPU code (CPU/GPU hybrid, AMD, Intel Phi) Free Energy Methods and Sampling
- j) Free Energy Calculations; dual topology. Alchemical Perturbation (extensions of existing code) Thermodynamic Integration / Metadynamics
- k) Double Precision (CPUs) vs. Mixed / Single Precision (GPUs) Does it matter? for which Problems?
- l) QM/MM (Gaussian, LChem, CHARMM, Q-Chem, others)
- m) Trajectory Storage, Transport and Archiving; Binary file format(s) Compatibility with ptraj / Amber

## 6) Applications & Tutorials (Nerenberg/Schnieders)

- a) UI/FFX (Schneiders/Ponder)
- b) Classroom & Laboratory Exercises Wiki and /or User Forum for Questions
- c) System set ups
- d) Free Energy Simulations  
FEP, BAR, MBAR,  $\Delta H / \Delta S$  QM / MM
- e) pKa Calculations
- f) Protein & Nucleic Acid
- g) Small Molecule Crystal Simulations
- h) Heavy Metal Ions

● **June 3rd, Sunday Dinner**  
**(Downtown Austin; 18:00-21:00 pm)**

Parkside <http://parkside-austin.com/>  
Roaring Fork <http://roaringfork.com/>

● **June 4<sup>th</sup>, Monday (BME 3.204, 9:00-11:30 am)**

## 7) GitHub / WebSites & License Issues

**TinkerTools.org Site**  
**Documentation, Regression Testing Tutorials**

**GitHub Site for each Tinker Code**

**Github usage/best practice: Josh/Andy**

Version N-1 will be Publically Available

Public GitHubs will allow Free Download (including Companies). Private GitHubs with access for Developers

Free license/open source for academic and non commercial

Commercial license between Washington Univ., Univ. Texas-Austin and Sorbonne

## 8) Organization of Software Development

### **Software Developers**

Core Developers Ponder / Ren / Piquemal, and parts of our groups Developers

Head-Gordon, Brooks, Cisneros, Schnieders, Yang, their groups, and other from this meeting

Contributors People from Outside this Meeting

### **Who is Going to Do What?**

Development Targets Release Dates / Plans Publications and Reference

Tinker papers:

Tinker 8 publication by tinker/tinker-HP key developers

Tinker 9 publication, including all developers?

## Participants:

Greg Beran	University of California, Riverside
Bernie Brooks	National Institutes of Health, NHLBI
Dave Case	Rutgers University
Sara Cheng	University of Texas, Austin
Andres Cisneros	University of North Texas
Tom Darden	OpenEye Scientific Software
Akshaya Das	University of California, Berkeley
Omar Demerdash	Oak Ridge National Laboratory
Peter Eastman	Stanford University
Nohad Gresh	Sorbonne Universite
Matthew Harger	University of Texas, Austin
Teresa Head-Gordon	University of California, Berkeley
Chris Ho	Drug Design Methodologies
Luc-Henri Jolly	Sorbonne Universite
Preston Keller	Kingdom Capital
Louis Lagardere	Sorbonne Universite
Filippo Lipparini	University of Pisa
Jacob Litman	University of Iowa
Chengwen Liu	University of Texas, Austin
Marie Laury	Washington University in St. Louis
Daniele Loco	Sorbonne Universite
Yvon Maday	Sorbonne Universite
Jessica Nash	MolSSI
Paul Nerenberg	California State University, Los Angeles
Dominique Nocito	University of California, Riverside
Xiaoliang Pan	University of Oklahoma
Cong Pan	Hong Kong University of Science & Technology
Jean-Philip Piquemal	Sorbonne Universite
Jay Ponder	Washington University in St. Louis
Rui Qi	University of Texas, Austin
Josh Rackers	Washington University in St. Louis
Julia Rice	IBM, Almaden
Pengyu Ren	University of Texas, Austin
Michael Schnieders	University of Iowa
Yihan Shao	University of Oklahoma
Michael Schauerl	University of California, San Diego
Andrew Simmonett	National Institutes Health, NHLBI
Benjamin Stamm	RWTH Aachen University
Ernest Smiley	Kingdom Capital
Bill Swope	IBM, Almaden
Mark Tuckerman	New York University
Sameer Varma	University of South Florida
Brandon Walker	University of Texas, Austin
George Whitwell	North Carolina Wesleyan College
Wei Yang	Florida State University
Zhi Wang	Washington University in St. Louis
Guowei Wei	Michigan State University