# Bio-molecular Simulations on Future Computing Architectures

Oak Ridge, Tennessee



Location: JICS Auditorium, Building 5100, Oak Ridge National Laboratory

|                        | Day 1: September 16 <sup>th</sup>   |
|------------------------|---|
| 8:00 – 8:30 am         | Meet and Greet (Breakfast provided)   |
| Session I: Emerging A  | Architecture Trends for Scientific Computing                                    |
| Chair: Pratul Agarwal, | Oak Ridge National Laboratory   |
| 8:30 – 8:40 am         | Arthur Barney Maccabe   |
|                        | Welcome   |
| 8:40 – 9:20 am         | Al Geist, Oak Ridge National Laboratory   |
|                        | "The Co-Design Challenges of going from Petascale to Exascale"                  |
| 9:20 – 9:50 am         | Duncan Poole, NVIDIA  |
|                        | "GPU architecture and applications supporting life science development"         |
| 9:50 – 10:00 am        | Coffee Break  |
| 10:00 – 10:30 am       | Eric Stahlberg, Wittenberg University   |
|                        | "Confidence in Computing –Interfaces, Portability, Reliability and Performance" |
| 10:30 – 11:00 am       | Gilad Shainer, Mellanox Technologies  |
|                        | "The Development of Mellanox-NVIDIA GPUDirect over InfiniBand – a New Model for |
|                        | GPU to GPU Communications"  |
| 11:00 – 11:30 am       | Robert Hinde, University of Tennessee, Knoxville                                |
|                        | "GPU-accelerated quantum Monte Carlo simulations"                               |
| 11:30 – 12:15 pm       | Round Table Discussion A  |
| 12:15 – 1:00 pm        | Lunch (Provided)  |

### Session II: Exploiting GPUs and FPGAs for Biomolecular Simulations

Chair: Qiang Cui, University of Wisconsin

| 1:00 – 1:40 pm | John Stone, University of Illinois at Urbana-Champaign                            |
|----------------|---|
|                | "High Performance Molecular Simulation, Visualization, and Analysis on GPUs"      |
| 1:40 – 2:10 pm | Gianni De Fabritiis, Universitat Pompeu Fabra, Spain                              |
|                | "High-throughput molecular dynamics simulations using ACEMD on GPUs"              |
| 2:10 – 2:40 pm | Joshua Anderson, University of Michigan   |
|                | "Simulating nano-materials, polymers, and complex fluids on GPUs with HOOMD-blue" |
| 2:40 – 3:00 pm | Coffee Break  |
| 3:00 – 3:30 pm | Martin Herbordt, Boston University  |
|                | "FPGA Acceleration of Molecular Dynamics and Docking"                             |
| 3:30 – 4:00 pm | Scott LeGrand, NVIDIA   |
|                | TENTATIVE: "Adapting AMBER particle mesh Ewald and generalized Born molecular     |
| 4.00 5.00 mm   | dynamics to GPUs"   |
| 4:00 – 5:00 pm | Round Table Discussion B  |

# Day 2: September 17<sup>th</sup>

7:45 – 8:00 am Meet and Greet (Breakfast provided)

**Session III: Challenges and Solutions for Biomolecular Simulations on Future Architectures** Chair: Paul Crozier, Sandia National Laboratories

| John Levesque, Cray Inc.  |
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| <i>"Future Exascale architectures and the challenges they impose on the application developers"</i> |
| Dave Norton, High Performance Fortran Associates  |
| "Programming GPUs with PGI's Compilers and Tools"   |
| Erik Lindahl, Stockholm University, Sweden  |
| "How can molecular simulation reach the exascale? Challenges in performance and parallelism"        |
| Coffee Break  |
| Laxmikant (Sanjay) Kale, University of Illinois at Urbana-Champaign                                 |
| "Strong scaling issues for classical molecular dynamics"  |
| Pratul Agarwal, Oak Ridge National Laboratory   |
| "Optimal biomolecular simulations on future computing architectures"                                |
| Round Table Discussion C  |
| Lunch (Provided)  |
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#### Session IV: Novel methods and better simulations

Chair: Kennie Merz, University of Florida

| 1:00 – 1:40 pm | Kennie Merz, University of Florida   |
|----------------|--|
|                | "Quantum Chemical Insights into the Computation of Absolute Interaction Energies"                                  |
| 1:40 – 2:10 pm | Imran Haque, Vijay Pande's Lab., Stanford University   |
|                | "Hybrid Vigor: Using Heterogeneous HPC to Accelerate Chemical Informatics"   |
| 2:10 – 2:40 pm | David Hardy, University of Illinois at Urbana-Champaign  |
|                | "Simulating Biomolecules on GPUs with the Multilevel Summation Method"   |
| 2:40 – 3:00 pm | Coffee Break   |
| 3:00 – 3:30 pm | Jaydeep Bardhan, Rush University Medical Center  |
|                | "Simultaneous Evolution of Physical Models and Computing Architectures: An   |
|                | Engineering Perspective"   |
| 3:30 – 4:00 pm | Chakra Chennubhotla, University of Pittsburgh  |
|                | "Automatic discovery of energetically coherent sub-states and rare events in a protein's conformational landscape" |
| 4:00 – 4:30 pm | Round Table Discussion D & Wrap-up   |

## **NOTES:**

- Shuttle service for pick-up and drop-off will be provided from Oak-Ridge hotels (Double-tree & Staybridge Suites) to the location.
- Please remember to bring suitable government issued identification to enter Oak Ridge National Laboratory. If you are NOT a US citizen, please bring your passport (and Green-card, if applicable).