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Computational Materials Online Seminar

GPUMD: A highly efficient molecular dynamics code fully implemented on graphics processing units

> Zheyong Fan (樊哲勇) Aalto University





#### Mini CV

- 2005-2010: PhD study at Nanjing University (Zhongzhou Ren) – Nuclear Physics, etc.
- 2010-2012: Postdoc research at Xiamen University (Jin-Cheng Zheng) – Thermoelectric transport
- 2012-now: Postdoc research at Aalto University (Ari Harju) – Heat and charge transport



QMP = quantum many-body http://physics.aalto.fi/en/groups/qmp/

- **GPUMD** = Graphics Processing Units Molecular Dynamics
- Implemented fully on Graphics Processing Units (GPUs)
- Many-body potentials: Tersoff, Stilinger-Weber, EAM, ...
- Highly efficient (you will see)
- Green-Kubo method for thermal conductivity calculations (LAMMPS is wrong for many-body potentials)
- NEMD method for thermal conductivity and thermal conductance calculations, with spectral decomposition
- More...

# Why develop GPUMD, when LAMMPS is available?

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- LAMMPS is not always correct
- LAMMPS is not always fast
- LAMMPS is not always convenient to use
- GPUs become more and more powerful
- LAMMPS cannot explore the full power of GPUs
- Why not create a new one that can get the most from the GPUs?

# **GPU Computing:** A cheap way to achieve high performance



- Price: 3000 RMB to 30000 RMB
- Speed: 3 Tflops DP (K80: 4992 cores)
- Memory: 4 GB to 24 GB
- Programming: CUDA C/C++



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#### Why GPU is faster?





• CPU:

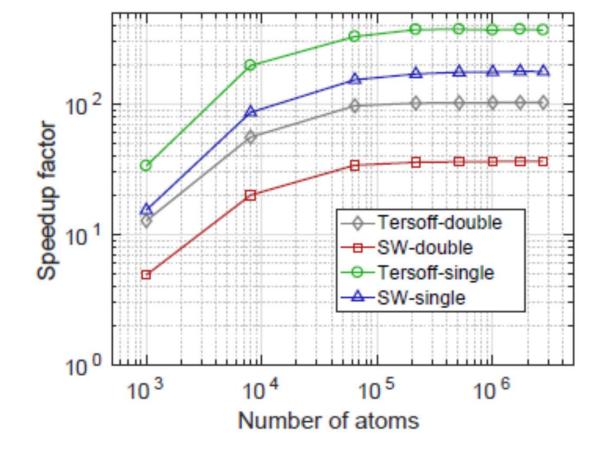
one or a few very fast computational threads

#### • GPU:

tens of thousand (not so fast) parallel threads

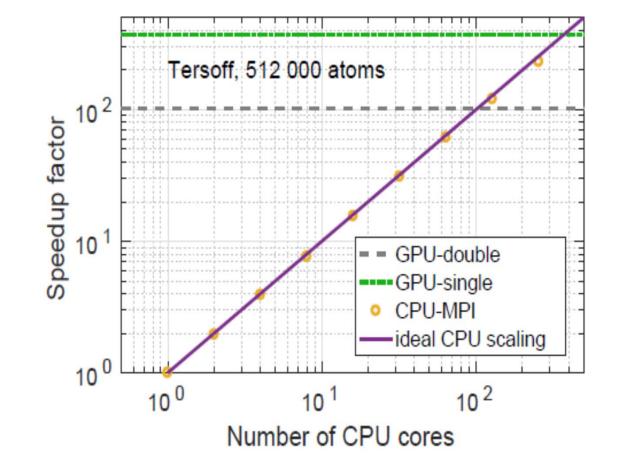
## How fast GPUMD is compared to a single CPU core?





- CPU: Intel Xeon
  X5670@2.93GHz
- GPU: K40 (2880
   CUDA cores)
- Model: Si@300K

# How fast GPUMD is compared to many CPU cores?



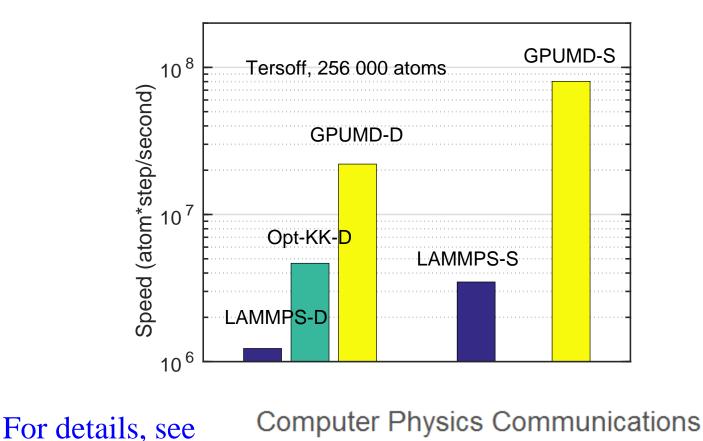
- CPU: Intel Xeon
   <u>X5670@2.93GHz</u>
- GPU: K40 (2880
   CUDA cores)
- Model: Si@300K

# How fast GPUMD is compared to the GPU versions of LAMMPS?

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GPU = Tesla K40

Model = Si@300K



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- It is fully implemented on GPUs
- It uses a set of elegant formulas for any many-body potential:
- Pair (!) force

$$\mathbf{F}_{ij} = -\mathbf{F}_{ji} = \frac{\partial U_i}{\partial \mathbf{r}_{ij}} - \frac{\partial U_j}{\partial \mathbf{r}_{ji}} = \frac{\partial \left(U_i + U_j\right)}{\partial \mathbf{r}_{ij}}$$

Per-atom virial

$$\mathbf{W}_i = -\frac{1}{2} \sum_{j \neq i} \boldsymbol{r}_{ij} \otimes \boldsymbol{F}_{ij}$$

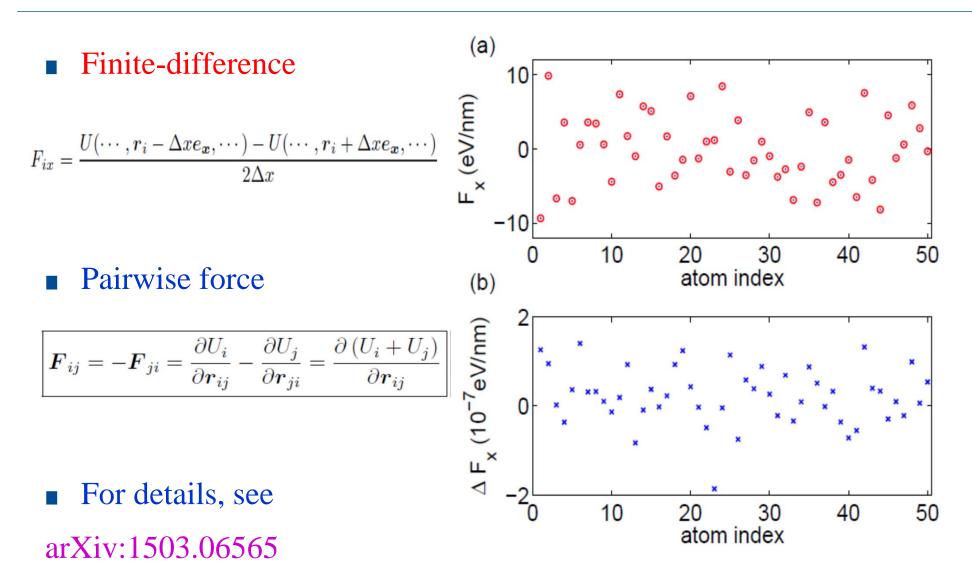
Per-atom heat current

$$oldsymbol{J}_{i}^{ ext{pot}} = \sum_{j 
eq i} oldsymbol{r}_{ij} \left( rac{\partial U_{j}}{\partial oldsymbol{r}_{ji}} \cdot oldsymbol{v}_{i} 
ight)$$

For details, see Phys. Rev. B **92**, 094301 (2015).

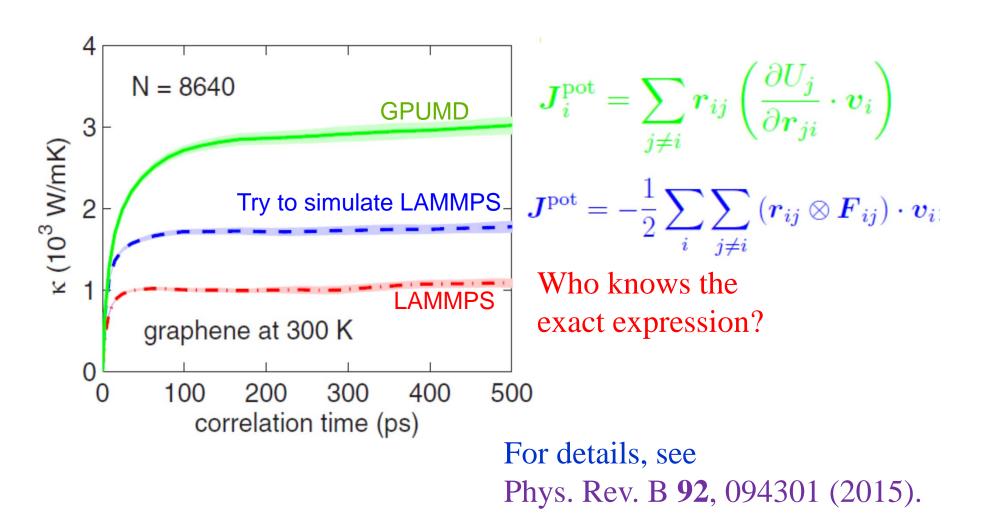
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#### **Doubt about the pairwise force expression?**



Zheyong Fan–Aalto University: zheyongfan@163.com

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#### The Green-Kubo method in GPUMD

Running thermal conductivity with the in-out decomposition

### **GPUMD input script:**

potential	potentials/c_tersoff_fan_2017.txt	
ensemble	npt_ber 300 300 0.01 0 0 0.0005	
time_step	1	]
dump_thermo	1000	]
run	100000	
ensemble	nve	
compute_hac	20 50000 10	
run	1000000	

For details, see Phys. Rev. B **95**, 144309 (2017).

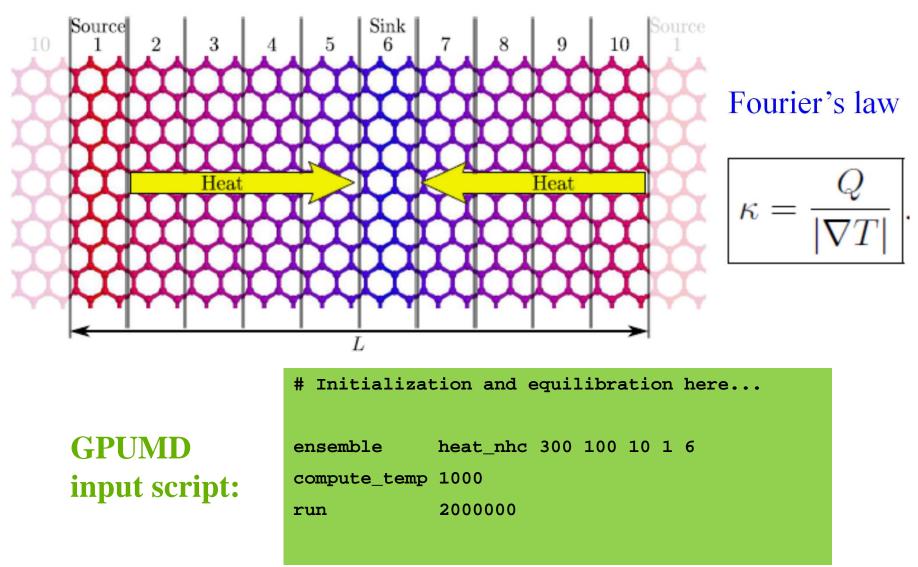
 $\kappa_{xx}^{\rm in}(t) = \frac{1}{k_B T^2 V} \int_0^t dt' C_{xx}^{\rm in}(t');$ 

 $\kappa_{xx}^{\text{out}}(t) = \frac{1}{k_B T^2 V} \int_0^t dt' C_{xx}^{\text{out}}(t');$ 

 $\kappa_{xx}^{\rm cross}(t) = \frac{1}{k_B T^2 V} \int_0^t dt' C_{xx}^{\rm cross}(t').$ 

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#### The NEMD method in GPUMD



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### The spectral-decomposition method in GPUMD

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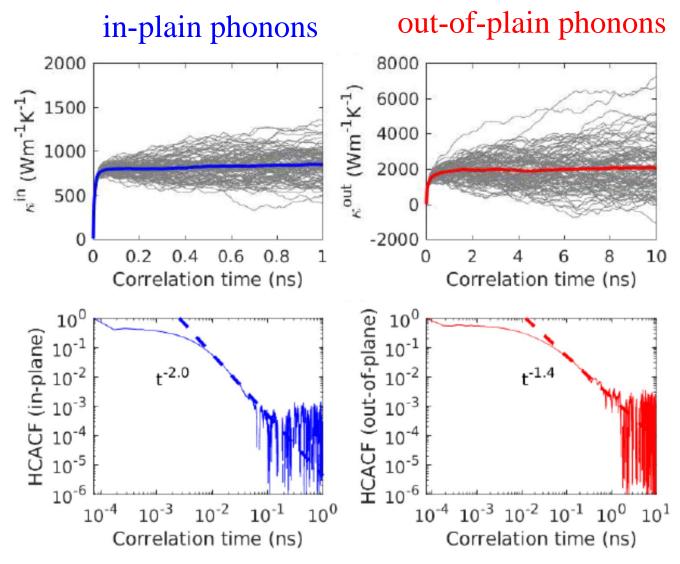
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Zheyong Fan–Aalto University: <u>*zheyongtan@163.com*</u>

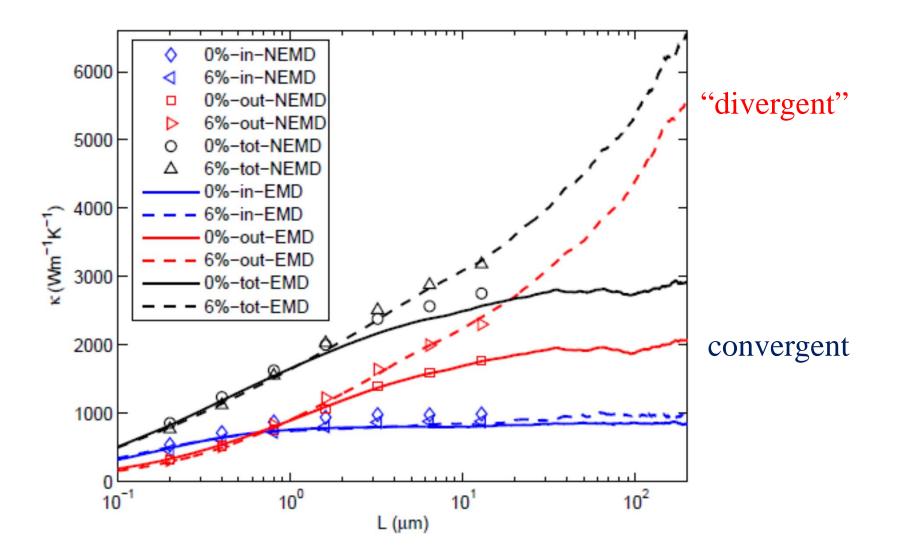
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#### **Application (1) – Thermal conductivity components of pristine graphene**





# Application (2) – Thermal conductivity $\mathcal{CMOS}$ "divergence" of graphene under tensile strain $2\ 0\ 1\ 7$



#### In summary,

- There is a fast MD code called GPUMD
- And I want to share it with you all
- <u>https://github.com/brucefan1983/GPUMD</u>
- I want to make it more powerful
- And your comments will be helpful to me
- The current version of GPUMD can already do many things
- And collaborations are welcome

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#### Last words:

