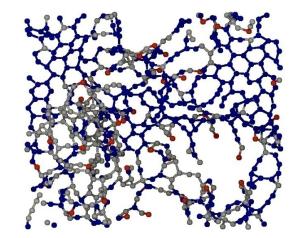
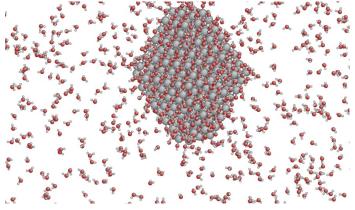
# Development and application of the ReaxFF reactive force field method

Adri van Duin, Murali Raju, Sriram Srinivasan, Jejoon Yeon, Sung-Yup Kim, Thomas Senftle and Kaushik Joshi Department of Mechanical and Nuclear Engineering Pennsylvania State University, 136 Research East Building phone: 814-8636277; E-mail acv13@psu.edu

LAMMPS workshop, August 2013



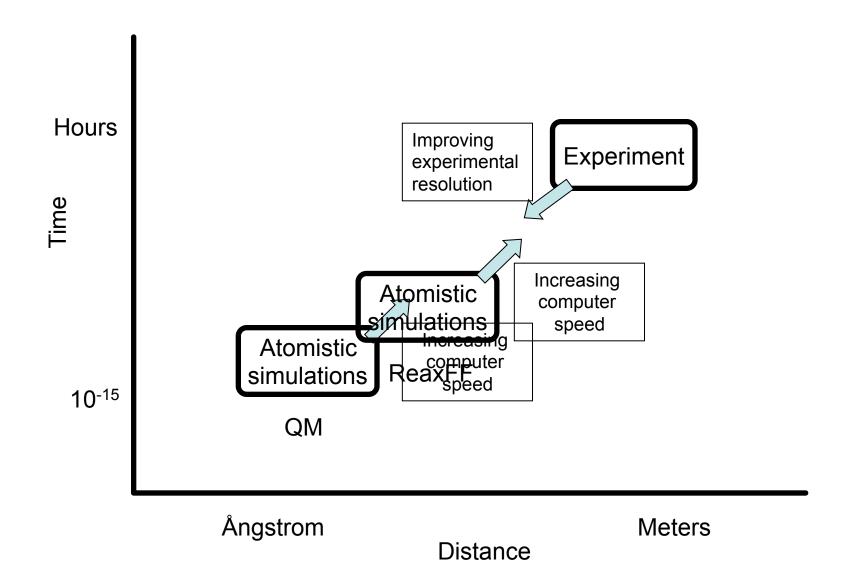
ReaxFF simulation of graphene breakup during oxidation (by Sriram Srinivasan; Srinivasan et al. JPC-A 2011)



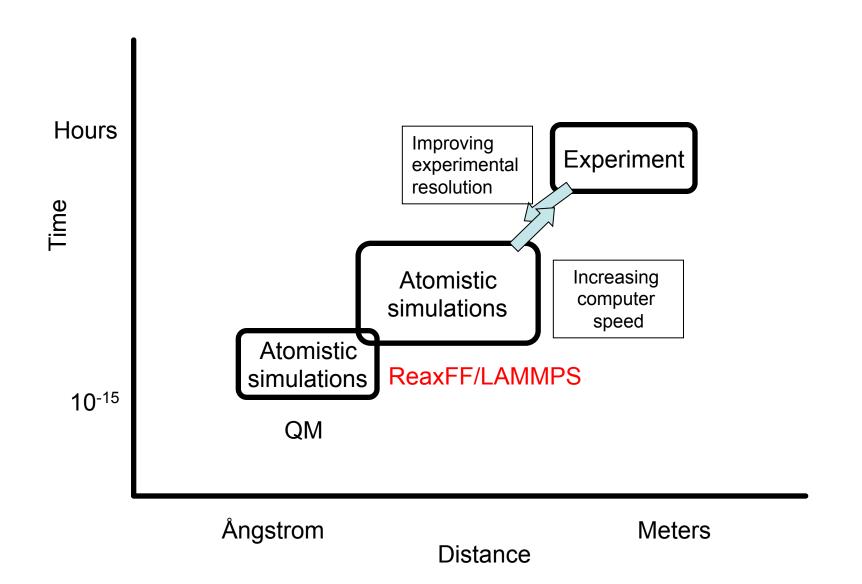
ReaxFF structure of a TiO<sub>2</sub> nanoparticle in water (by Murali Raju; Raju et al. JPC-C 2013)



# Solving the size/time gap between simulation and experiment

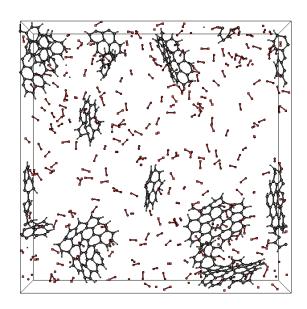


# Solving the size/time gap between simulation and experiment

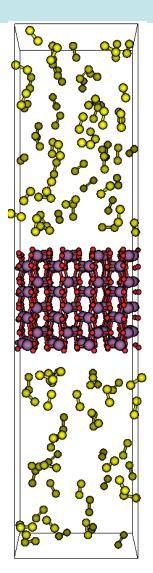


## **Outline**

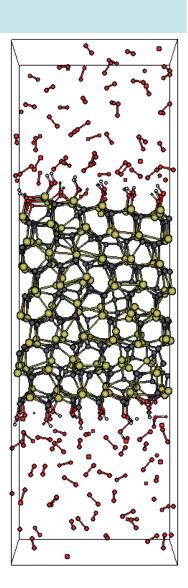
- The ReaxFF reactive force field
- Application to combustion reactions
- Application to aqueous-phase reactions
- New ReaxFF simulation options soon in LAMMPS (?)
- Summary



ReaxFF MD simulation of char combustion at T=2500K



ReaxFF MD simulation of S<sub>2</sub> gas reacting with a MoO<sub>3</sub> slab at T=1000K



ReaxFF MD simulation of the oxidation of a SiC-slab at T=3000K

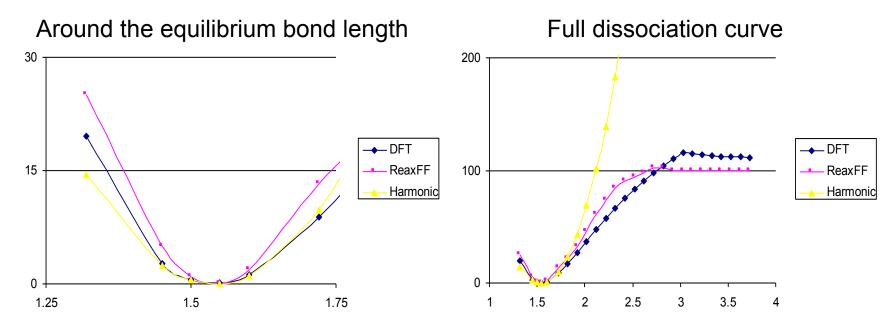
## Force field methods

$$E_{bond-stretch} = \sum_{1.2 \text{ vairs}} K_b (b - b_0)^2$$
 
$$E_{bond-bend} = \sum_{angles} K_\theta (\theta - \theta_0)^2$$
 
$$\Phi$$
 
$$E_{rotate-along-bond} = \sum_{n=1}^\infty K_n (1 - \cos(n\phi))$$

- Empirical, we need to derive values for the force field parameters (intuition, compare to experiment, compare to QM)
- MUCH faster than QM; can be applied to bigger systems

#### Failure of the harmonic model

#### C-C bond stretching in Ethane



- Although the harmonic approximation can describe the bond stretching around the equilibrium it cannot describe the bond dissociation.

#### Key features of ReaxFF

- -To get a smooth transition from nonbonded to single, double and triple bonded systems ReaxFF employs a bond length/bond order relationship [1-3]. Bond orders are updated every iteration.
- -All connectivity-dependent interactions (i.e. valence and torsion angles, H-bond) are made bond-order dependent, ensuring that their energy contributions disappear upon bond dissociation.
- Nonbonded interactions (van der Waals, Coulomb) are calculated between every atom pair, irrespective of connectivity. Excessive close-range nonbonded interactions are avoided by shielding.
- ReaxFF uses EEM, a geometry-dependent charge calculation scheme that accounts for polarization effects [4].
- 1. Brenner, D. W., (1990) Physical Review B 42, 9458-9471
- 2. Tersoff, J., (1988) Physical Review Letters 61, 2879-2882.
- 3. Abell, G. C., (1985) Physical Review B 31.
- 4. Mortier, W. J., Ghosh, S. K., and Shankar, S. (1986) JACS 108, 4315-4320.

#### General rules for ReaxFF

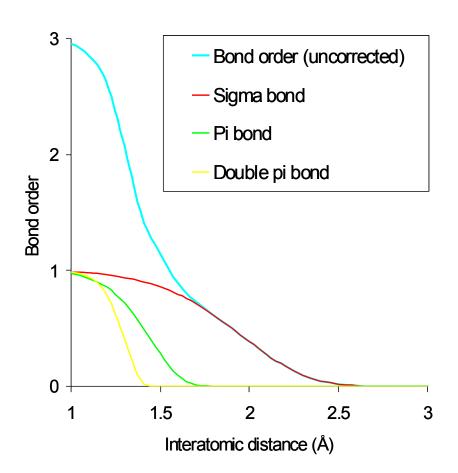
- MD-force field; no discontinuities in energy or forces even during reactions.
- User should not have to pre-define reactive sites or reaction pathways; potential functions should be able to automatically handle coordination changes associated with reactions.
- Each element is represented by only 1 atom type in the force field; force field should be able to determine equilibrium bond lengths, valence angles etc. from chemical environment.

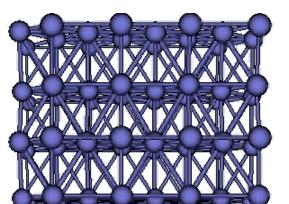
#### Calculation of bond orders from interatomic distances

$$BO'_{ij} = \exp\left[p_{bo,1} \cdot \left(\frac{r_{ij}}{r_o^{\sigma}}\right)^{p_{bo,2}}\right]$$

$$+ \exp\left[p_{bo,3} \cdot \left(\frac{r_{ij}}{r_o^{\pi}}\right)^{p_{bo,4}}\right]$$

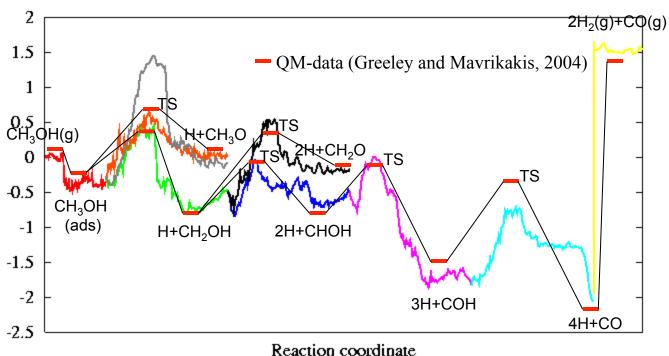
$$+ \exp\left[p_{bo,5} \cdot \left(\frac{r_{ij}}{r_o^{\pi\pi}}\right)^{p_{bo,6}}\right]$$

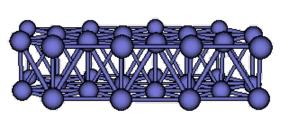




#### QM/ReaxFF methanol reaction pathways

#### Methanol reaction on Pt[111]surface

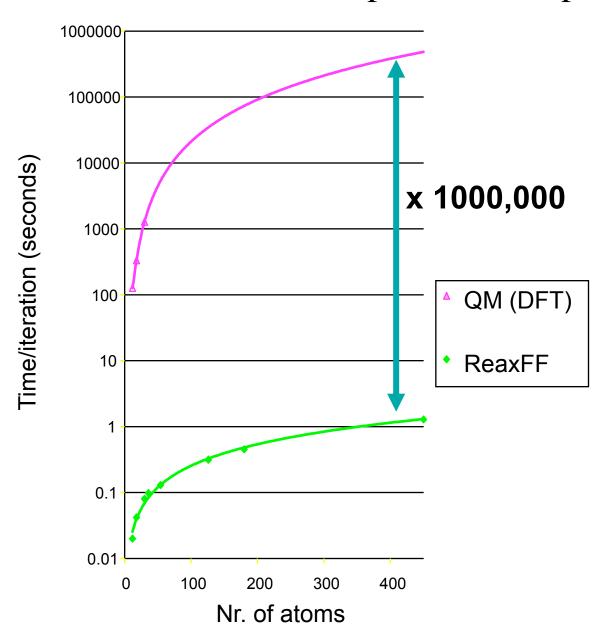




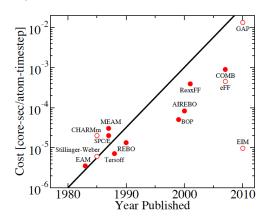


- Good agreement between ReaxFF and QM for entire reaction path

## ReaxFF Computational expense



- -ReaxFF allows for reactive MD-simulations on systems containing more than 1000 atoms
- ReaxFF is 10-50 times slower than non-reactive force fields
- Better scaling than QMmethods (NlogN for ReaxFF, N<sup>3</sup> (at best) for QM

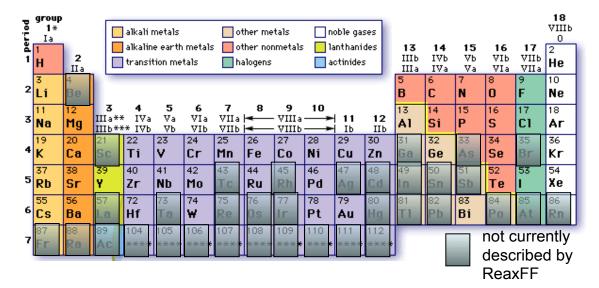


Steve Plimpton and Aidan Thompson, MRS review 2012

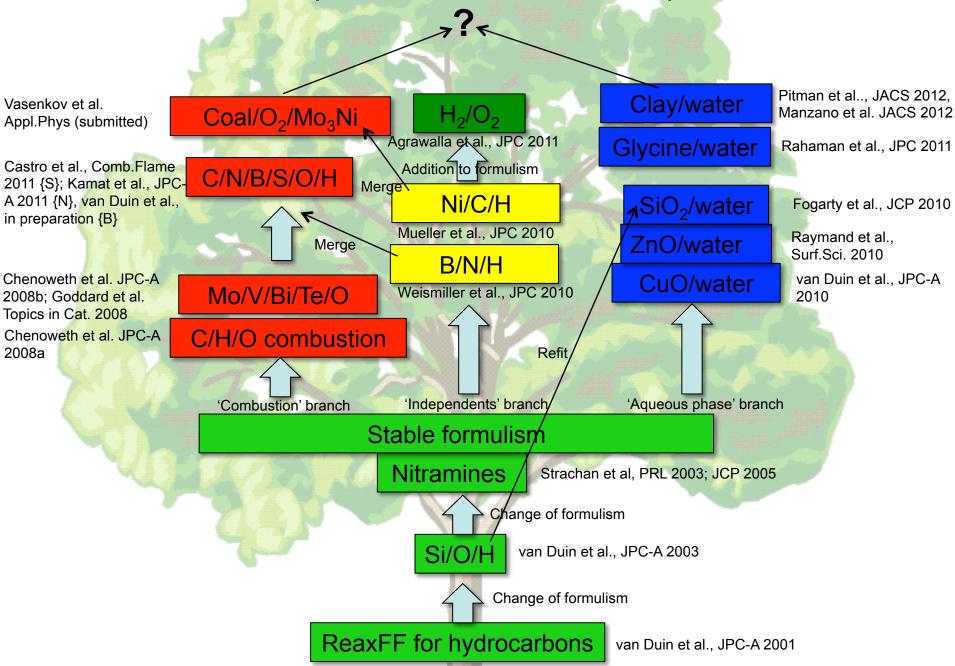
### Current development status of ReaxFF

- ReaxFF combines covalent, metallic and ionic elements allowing applications all across the periodic table
- All ReaxFF descriptions use the same potential functions, enabling application to interfaces between different material types
- Code has been distributed to over 250 research groups
- Over 150 publications in peer-reviewed journals
- Parallel ReaxFF (LAMMPS/ReaxFF) available as open-source
- Integrated in ADF/BAND graphical user interface

#### ReaxFF transferability



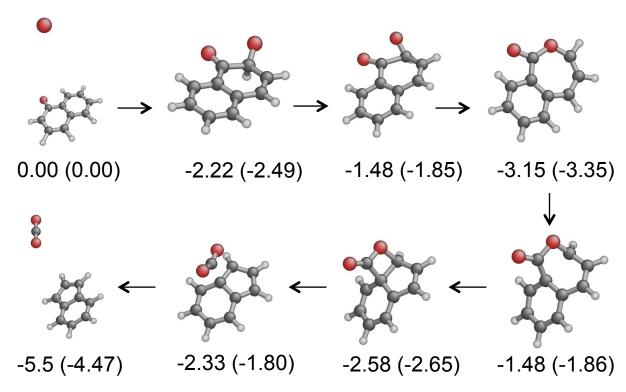
## ReaxFF development tree: towards complex materials



#### Graphene oxidation simulations

With Sriram Srinivasan and Tom Schwartzentruber

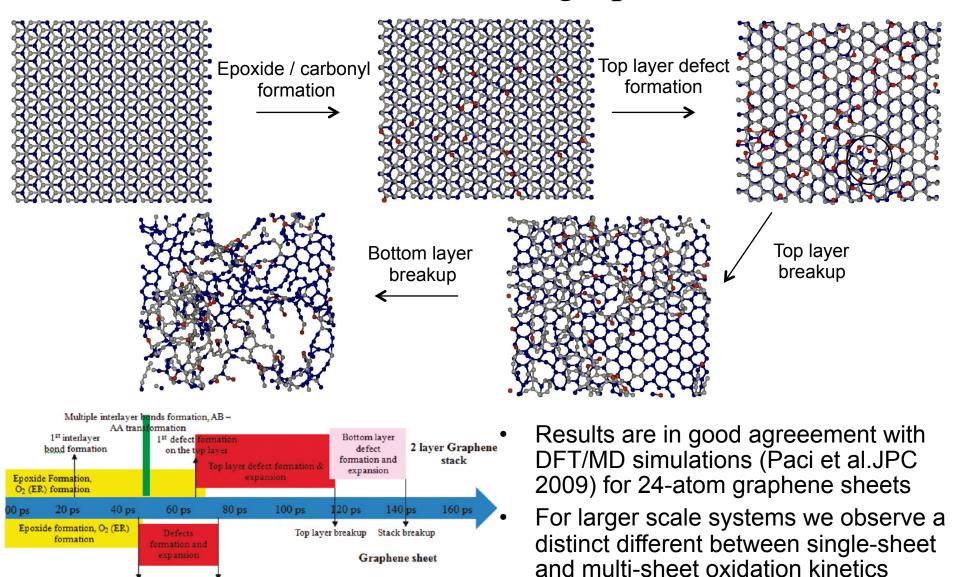
1. Force field development



dE in eV/mol; DFT [2] results in brackets

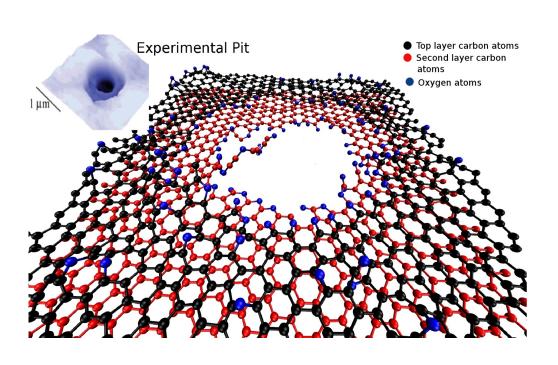
[1] Chenoweth K., van Duin A. C. T., Goddard W. A., *J. Phys. Chem. A* **2008**, 112, 1040–1053 [2] Orrego J. F., Zapata, F., Truong T. N., Mondragón F., *J. Phys. Chem. A* **2009**, *113* (29), 8415-8420

# 2. Validation: Process of graphite erosion

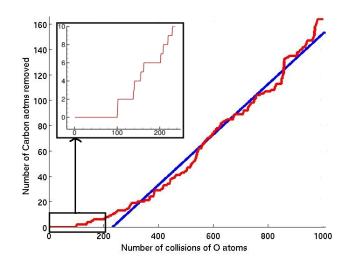


Appearance of the first defect Sheet Breakup

# 3. Application: Process of HOPG erosion, comparison with experiment and links to coarse-grain simulations

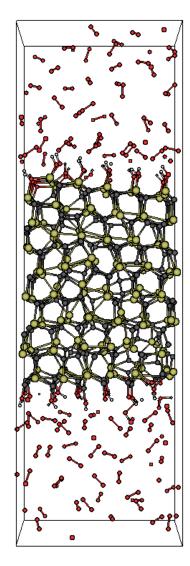


Nicholson K.T., Minton T.K., Seibner S.J., *J. Phys. Chem. B*, **2005**, 109, 8476 Poovathingal S., Schwartzentruber T.E., Goverapet Srinivasan S., van Duin A.C.T., *J. Phys. Chem. C.***2013**, 117, 2692



- Molecular beam experiments on HOPG sample showed a pitting behavior. Large shallow pits were observed on the surface of the sample.
- Rate of removal of carbon atoms from the graphite specimen: 1 carbon atom for every 5 oxygen atom collisions

### Increasing complexity: Silicon carbide oxidation [1,2]

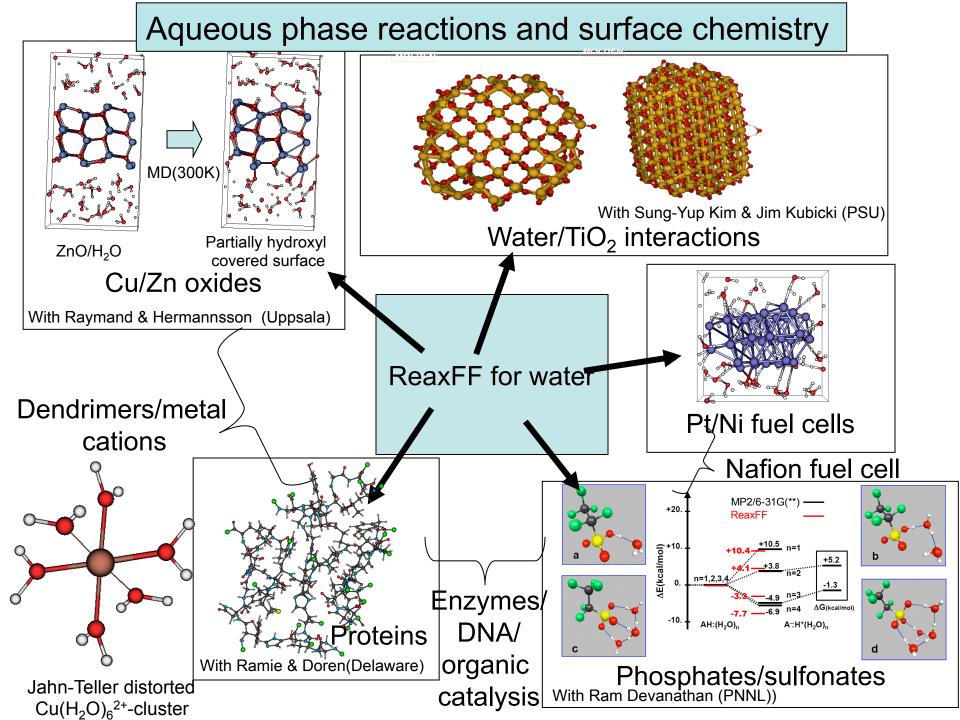


ReaxFF MD simulation of the oxidation of a SiC-slab at T=3000K

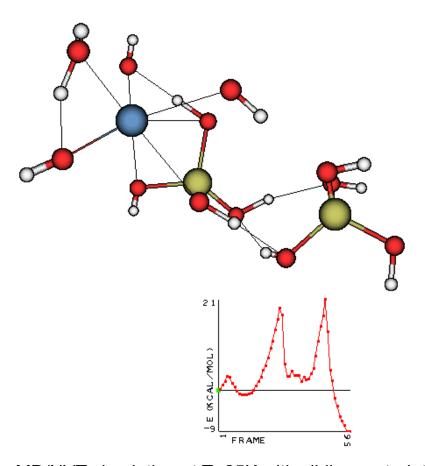
- Oxidation of a SiC-slab results in the formation of a protecting SiO<sub>2</sub> layer and a simultaneous phase separation, results in the formation of an amorphous graphitic phase
- Gas phase contains early carbon oxidation products (CO, CO<sub>2</sub>), but graphitic phase remains unoxidized for the duration of the MD-simulations

<sup>[1]</sup> Newsome, D., Sengupta, D., and van Duin, A.C.T., **2013**. Journal of Physical Chemistry C accepted for publication.

<sup>[2]</sup> Newsome, D., Sengupta, D., Foroutan, H., Russo, M. F., and van Duin, A. C.T. Journal of Physical Chemistry **2012**, *116*, 16111-16121.



#### Applications to water/silica chemistry



Si(OH)<sub>4</sub> dimerization catalyzed by a Ca(OH)<sub>2</sub>.2H<sub>2</sub>O complex

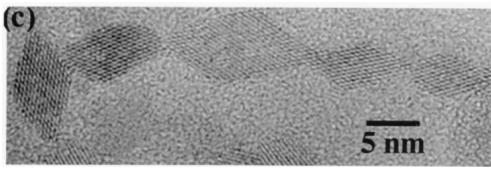
- Ca-ion assists by coordinating to the silanol-group, facilitating proton transfer to a hydroxyl coordinated to the Ca-cation
- Barrier with Ca: 21 kcal/mol; without Ca: 28 kcal/mol (DFT/B3LYP/6-311G\*\* [1])
- Ca/O/H parameters from Manzano et al., Langmuir 2012 [2]

MD/NVT-simulation at T=25K with sliding restraint to drive reaction 5000 iterations
CPU-time (this laptop) 35 seconds

## Oriented attachment in Titania Nanoparticles

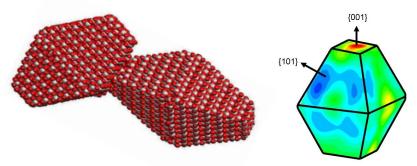
With Murali Raju and Kristen Fichthorn

- Oriented attachment occurs most commonly on {112}, occasionally on (001), and rarely on {101}.
- This mechanism effectively serves to reduce overall surface energy by eliminating the surfaces at which the crystallites join

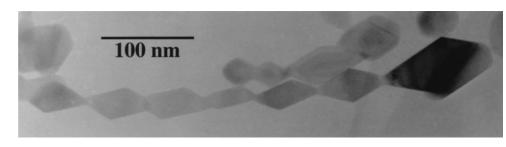


five primary crystallites forming a single crystal via oriented attachment

#### Preferential Alignment in vacuum



M. Alimohammadi and K. Fichthorn, *Nano Lett.* 9, 4198 (2009).



TEM micrograph of a single crystal of anatase that was hydrothermally coarsened in 0.001 M HCl.

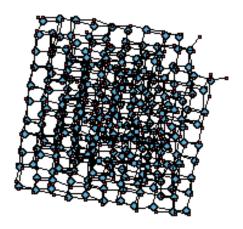
R. Penn and J. Banfield, *Geochim. Cosmochim. Acta* 63, 1549 (1999).

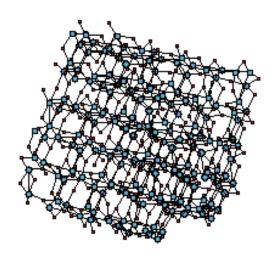
A Review on Oriented Attachment:

20

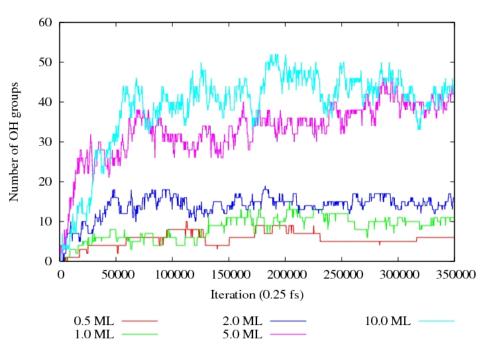
M. Niederberger and H. Cölfen, Phys. Chem. Chem. Phys. 8, 3271 (2006).

## Behavior in Vacuum: No Oriented Attachment





### Water Dissociation on Anatase Nanoparticles



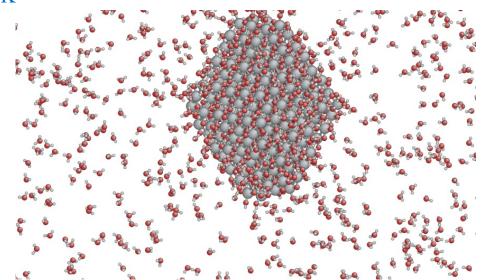
#### Hypothesis

Solvent mediates growth by oriented attachment through adsorbed surface groups; OH groups and adsorbed H<sub>2</sub>O.

Investigate amount of solvent needed to saturate the nanoparticle surface with surface groups.

Hydroxyl coverage on an anatase nanoparticle at 1100K

- The nanoparticle surface is saturated with OH groups around a coverage of 5 ML.
- Run simulations with a coverage of 5 ML to aid in faster diffusion of nanoparticles.

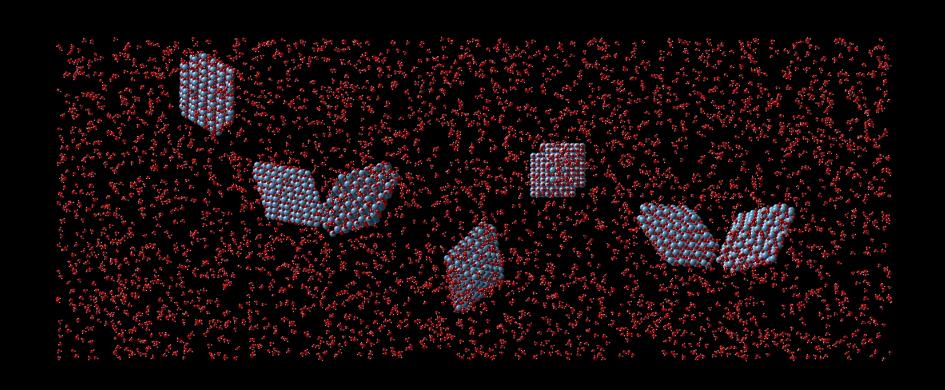


#### Behavior in Water

Simulation Cell: 125 Å x 325 Å x 125 Å

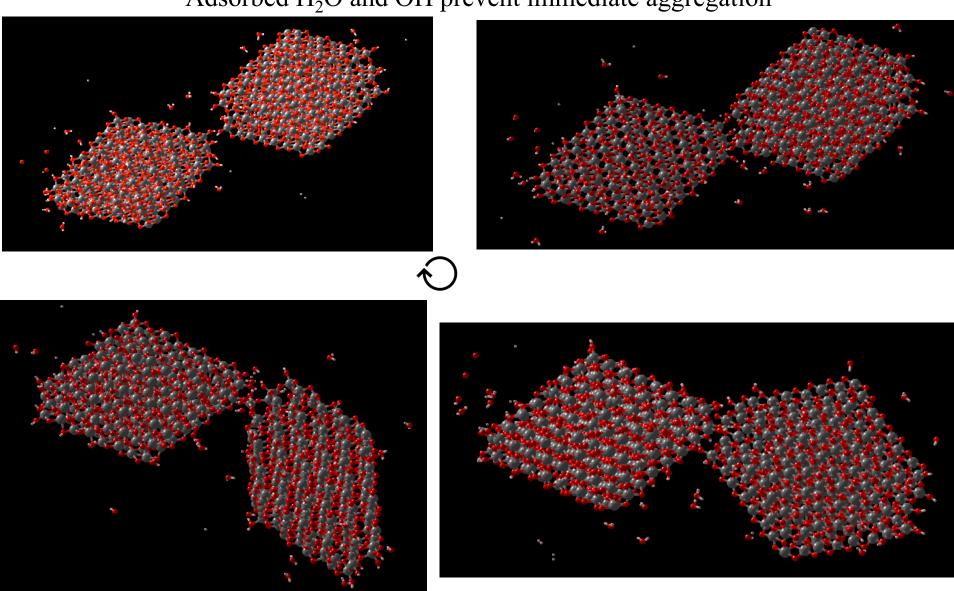
NVT at 1100 K (aid in faster diffusion of nanoparticles)

8 - (112) truncated nanoparticles and 4270 - water molecules

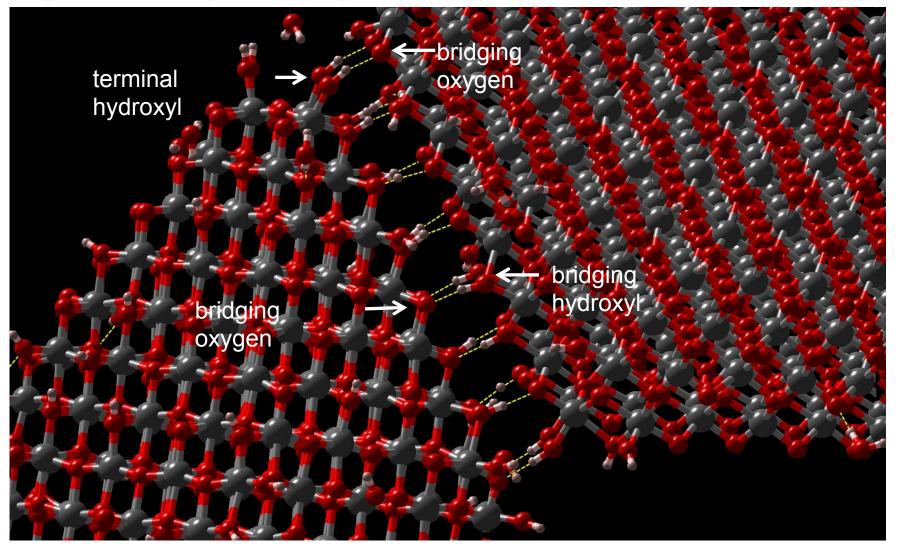


### Behavior in Water

Role of Water -1 Adsorbed H<sub>2</sub>O and OH prevent immediate aggregation



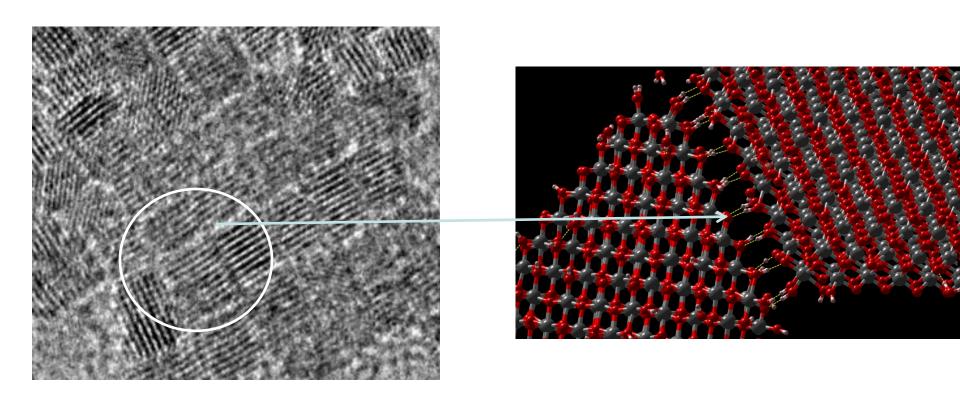
Hydrogen bond



- The nanoparticles slide along each other, making and breaking hydrogen bonds until they align
- Aligning favors the formation of a network of hydrogen bonds between the terminal or bridging hydroxyls on one nanoparticle and the surface oxygens on the other nanoparticle.

#### Comparison with Experiment

## How does it compare with experiment?



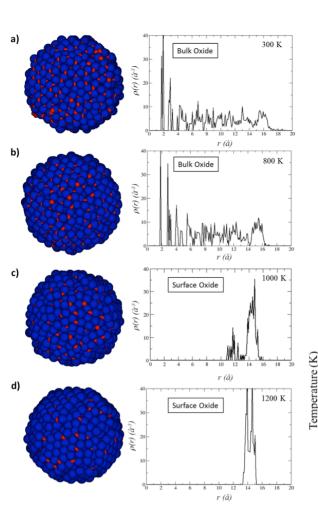
TEM image of 'wet' SnO<sub>2</sub> nanoparticles

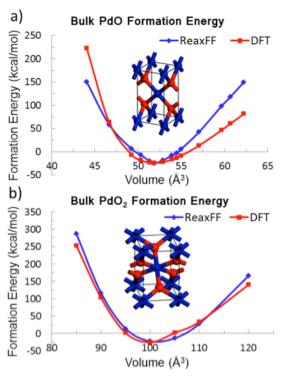
Image Courtesy: Wei Wang, Hsiu-Wen Wang at ORNL

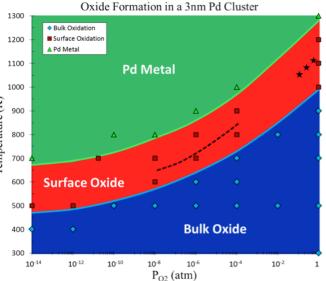
#### New ReaxFF simulation options – soon in LAMMPS (?)

# Hybrid Grand Canonical Monte Carlo/MD ReaxFF [1] With Thomas Senftle and Michael Janik

- -Enables prediction of thermodynamic end states at various P/T conditions - Application to PdO<sub>x</sub> shows high-T reduction, in good agreement with experiment [2,3]
- Hybrid GCMC/MD allows crystal morphology change during oxidation/reduction
- Highly transferable tool







[1] Senftle, Janik, Meyer and van Duin, Journal of Chemical Physics 2013, in print

[2] Ketteler, G.; Ogletree, D. F.; Bluhm, H.; Liu, H. J.; Hebenstreit, E. L. D.; Salmeron, M. J. Am. Chem. Soc. 2005, 127, 18269.

[3] Zhang, H.; Gromek, J.; Fernando, G.; Marcus, H.; Boorse, S. Journal of Phase Equilibria and Diffusion 2002, 23, 246.

# Parallel Replica Dynamics with ReaxFF in LAMMPS (Kaushik Joshi & Sumathy Raman(EMRE)

• PRD¹ parallelizes rare event simulation in time domain and accelerate time

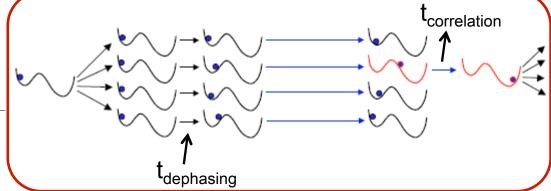
scale of MD

Infrequent events

Transitions can be detected

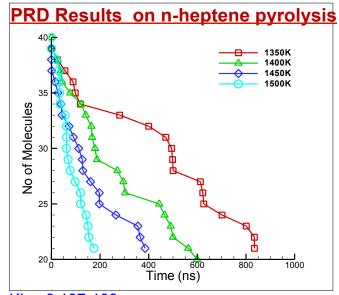
 Exponential distribution of firstescape times

Correlation time known



 PRD is coupled with REAXFF in LAMMPS by implementing a connectivity-based event detection

	PRD	Experiment <sup>7</sup>
E <sub>a</sub> (kcal/mol)	50.01	54
Pre exponential factor, A	$7.17 \times 10^{13}$	$6.3 \times 10^{11}$



<sup>1</sup>Phys. Rev. B, 57, R13985 (1998); <sup>7</sup> Blades A.T., Sandhu H.S. (1971) Int, J. Kin., 3 187-193

### EEM upgrade: ACKS2

#### Condensed form for the 'electronic energy'

$$E_{\text{EEM}} = \left(\sum_{A} \chi_{A} q_{A} + \frac{1}{2} \eta_{A} q_{A}^{2}\right) + \left(\frac{1}{2} \sum_{A,B \neq A} \frac{q_{A} q_{B}}{|\mathbf{r}_{A} - \mathbf{r}_{B}|}\right)$$

Variables:  $q_i$ 

Parameters:  $\chi_i, \eta_i, \bar{r}_i$ 

# In collaboration with Toon Verstraelen (U. Gent)

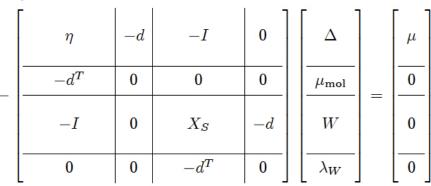
#### Charges minimize E<sub>EEM</sub> (with total charge constraint)

$$\frac{\partial E_{\text{EEM}}}{\partial q_A} = \sum_{A} \left( \chi_A + \eta_A q_A + \sum_{B \neq A} \frac{q_B}{|\mathbf{r}_A - \mathbf{r}_B|} \right) = \chi_{\text{mol}}$$

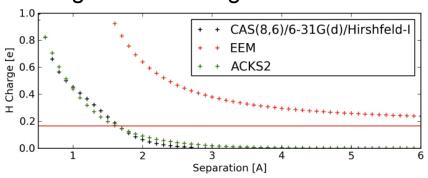
#### EEM equations in block matrix notation

$$- \left[ egin{array}{c|c} \eta & -d \ \hline -d^T & 0 \end{array} 
ight] \left[ egin{array}{c|c} q \ \hline \chi_{
m mol} \end{array} 
ight] = \left[ egin{array}{c} \chi \ \hline q_{
m tot} \end{array} 
ight]$$

#### ACKS2 equations in block matrix notation



#### Charges on H during HF-dissociation



- -EEM gives non-zero charges after bond dissociation
- ACKS2 enforces integer charges
- ACKS2 solves polarization issues (EEM: metallic)

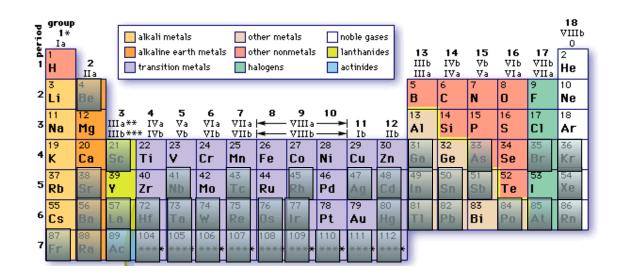
Verstraelen et al. JCP 2013<sup>29</sup>

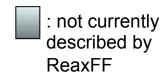
#### ReaxFF/GPU developments

- ReaxFF/GPU published by Zheng, M., Li, X., and Guo, L., 2013. Algorithms of GPU-enabled reactive force field (ReaxFF) molecular dynamic. Journal of Molecular Graphics and Modelling 41. Single precision? Not very impressive speed-up.
- Ananth Grama-group has recently finished an initial ReaxFF/GPU version, based on C++/ReaxFF version integrated in LAMMPS. Double precision, gcc 4.5.4, CUDA 5.0.
- Purdue/GPU/ReaxFF looks very promising 4 times faster then best current ReaxFF method for 2000 atoms; looks even better for larger systems (5000-10,000 atoms).
- Needs further development (memory management, parallel GPU)

## Summary

- ReaxFF has proven to be transferable to a wide range of materials and can handle both complex chemistry and chemical diversity. Specifically, ReaxFF can describe covalent, metallic and ionic materials and interactions between these material types.
- The low computational cost of ReaxFF (compared to QM) makes the method suitable for simulating reaction dynamics for large (>> 1000 atoms) systems (single processor). ReaxFF/LAMMPS allows reactive simulations on >>1000,000 atoms, enabling close interaction with experiment





## Acknowledgments

#### Collaborators:

- Jonathan Mathews, Jim Kubicki, Deborah Levin, Rich Yetter and Mike Janik (Penn State)
- Kimberley Chenoweth, Vyacheslav Bryantsev and Bill Goddard (Caltech)
- Aidan Thompson, Steve Plimpton (Sandia), Ananth Grama (Purdue), Metin Aktulga (Purdue) (parallel MD)

Funding: - PSU/KISK startup grant #C000032472



- NSF (TiO<sub>2</sub>/water, PdO/Ceria)

- NETL/RUA (Fuel catalysis)

DoE/NETL (Refractory materials)

AFRL/SBIR (Hydrocarbon cracking)

- AFOSR/MURI (O-resistant materials)

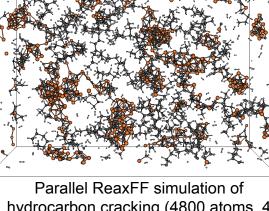
- DoE/EFRC FIRST-center

- Exxon (Software development, catalysis)

- British Royal Society (initial ReaxFF funding)







hydrocarbon cracking (4800 atoms, 4 processors)

More information:

Website: http://www.engr.psu.edu/adri

Office: 136 Research East Phone: 814-863-6277 E-mail: acv13@psu.edu

