



# Molecular Dynamics simulations of sulfuric acid cluster collisions

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### Molecular Dynamics of two H<sub>2</sub>SO<sub>4</sub> molecules colliding



• v<sub>rel</sub> = 200 m/s, OPLS-AA force field (sulfuric acid dipole moment = 3.07 Debye)



- Molecules have long-ranged attractive interactions:
  - Charge charge (Coulomb):
  - Charge dipole:
  - Dipole dipole (Keesom):
  - Induced dipole dipole (Debye):
  - Induced dipole induced dipole (London dispersion):

$$U(r) = \frac{q_1 q_2}{4\pi\varepsilon_0 r}$$

$$U(r) = -\frac{q_1^2 \mu_2^2}{6(4\pi\varepsilon_0)^2 kTr^4}$$

$$U(r) = -\frac{\mu_1^2 \mu_2^2}{3(4\pi\varepsilon_0)^2 kTr^6}$$

$$U(r) = -\frac{\mu_1^2 \alpha_2}{(4\pi\varepsilon_0)^2 r^6}$$

$$U(r) = -\frac{4}{3} \frac{h\nu\alpha^2}{(4\pi\varepsilon_0)^2 r^6}$$
Van der Waals interaction

### Motivation: collision coefficients for ACDC code



$$\frac{dc_i}{dt} = \frac{1}{2} \sum_{i' < i} \beta_{i',(i-i')} c_{i'} c_{(i-i')} + \sum_{i'} \gamma_{(i+i'),i'} c_{(i+i')} - \sum_{i'} \beta_{i,i'} c_i c_{i'} - \frac{1}{2} \sum_{i' < i} \gamma_{i,i'} c_i - \operatorname{sinks} (+ \operatorname{sources})$$

- cluster concentrations  $c_i$
- collision coefficients  $\beta_{i, i}$
- evaporation coefficients  $\gamma_{i,j}$  (lifetime =  $1/\gamma_{i,j}$ )
- sinks: coagulation sink to larger particles, wall losses (in the lab)
- sources: for example single vapour molecules can have sources (emissions, air chemistry)

### 1. Force Field validation



### Benchmark set: 22 conformers of [(H<sub>2</sub>SO<sub>4</sub>)<sub>2</sub>,HSO<sub>4</sub>]<sup>-</sup> trimer

- QM structures optmised using B97X-D / 6-31++G(d,p)
- Two force fields tested:
  - Ding et al. [1] : fitted to DFT calculations of clusters containing sulfuric acid, bisulfate and water
  - OPLS-AA [2] : transferable potential for biological systems



C. G. Ding, T. Taskila, K. Laasonen and A. Laaksonen, Chem. Phys. 287, 7 (2003).
 W. L. Jorgensen, D. S. Maxwell and J. Rives-Tirado, J. Am. Chem. Soc. 118(45), 11225 (1996).

### Forcefield benchmark results: energies and structures



- Force field energies agree better between themselves than with QM energies
- Cases of bad structural match appear to be due to bifurcated hydrogen bonds, unfavorable in both FFs

### H<sub>2</sub>SO<sub>4</sub> vibrational spectra from FT of velocity ACF

Force field by Ding et al. (structure fitted to DFT of clusters of sulfuric acid/bisulfate/ water) does a very bad job...





### Forcefield benchmark: summary

- Ding et al.
  - Fitted explicitly to sulphuric acid/bisulphate/water cluster structures from DFT
  - Crude molecular mechanics model no angles and dihedrals fitted, only bonds
  - Gas phase geometries correct; dipole of 3.7 Debye
  - Vibrational spectra horrible
  - No transferability
- OPLS-AA
  - Gas phase geometries correct, dipole of 3.1 Debye
  - QM benchmark a bit worse than Ding et al.
  - Vibrational spectra good
  - Transferable potential
- OPLS-AA will be used for collision simulations

## 2. Collision statistics in molecular dynamics simulation



### Hard sphere diameter vs. dimer PMF from simulation

-0.2

-0.3

0

• Hard sphere model of H<sub>2</sub>SO<sub>4</sub>

*M* = 98.09 g/mol,  $\rho$  = 1.84 g/cm<sup>3</sup> --> *V*/molecule = *M*/( $\rho \cdot N_A$ ) = 8.8516 × 10<sup>-23</sup> cm<sup>3</sup> Volume fraction = 1

--> 
$$R = \sqrt[3]{\frac{1 \cdot V \cdot 3}{4\pi}} = 2.76 \times 10^{-8} \text{ cm} = 2.76 \text{ Å}$$

Random close packing: Volume fraction = 0.64 -->  $R = \sqrt[3]{\frac{0.64 \cdot V \cdot 3}{4\pi}} = 2.38 \times 10^{-8} \text{ cm} = 2.38 \text{ Å}$ 

Hard sphere diameter: 4.8 – 5.5 Å

 PMF from Metadynamics simulation at 300 K, OPLS-AA force field (dipole moment = 3.07 Debye)

 $n^3$  0.4 0.5 0.4 r 0.2 0.2 0.10.1

5

10

Distance r (Å)

15

20

25

PMF has minimum at 4.2 Å, but is attractive to much larger distances!

### Setup for collision simulations

- OPLS-AA force field, LJ cutoff at 14 Å, Coulomb cutoff at 120 Å
- Assign atomic velocities from Maxwell-Boltzmann distribution
- Remove c.o.m. motion of each molecule individually
- Equilibrate/randomise for 50 ps in NVE
- Add translational velocities  $v_{\chi} = \pm \frac{v_r}{2}$  to the molecules
- Run NVE MD for collision
- Sample impact parameter *b* from 0 to 17.5 Å in steps of 0.5 Å
  - 1000 runs for each value of b
- Sample relative velocities  $v_r$  according to Maxwell-Boltzmann distribution, from 50 to 800 m/s in steps of 50 m/s
- Obtain statistics on collision and sticking probability as function of b and  $v_r$



### Collision probability: impact parameter and relative velocity



$$v_r^p = \sqrt{\frac{2kT}{\mu}} = \sqrt{\frac{2 \cdot 2kT}{m}} = \sqrt{2}v^p$$

### Collision coefficients

• Identical hard spheres:

$$\beta_{HS} = \sqrt{\frac{16kT}{\pi m}\pi(2r)^2}$$

- Molecular dynamics:  $\beta_{MD} = \pi \int_0^\infty dv \int_0^\infty db^2 v f(v) P(b)$
- Central field theory:  $\int_{1}^{\infty}$

$$\beta_{CF} = \pi \int_0^\infty v f(v) \ b_{max}^2 \ dv$$

$$\beta_{MD} \approx 2.2 \ \beta_{HS}$$
  
 $\beta_{CF} \approx 2.7 \ \beta_{HS}$ 



"The factor of 2.3 has previously been shown to give good agreement between measured and modeled cluster and particle concentrations for the chemical system of sulfuric acid and DMA" A. Kürten et al., Atmos. Chem. Phys., 18, 845 (2018).

### Summary and Outlook

- Force field benchmark against QM optmised structures of 22 sulfuric acid trimers
  - OPLS-AA suitable for collision simulations
- Simulation of collisions of two sulfuric acid molecules
  - Collisions simulated over a range of impact parameters and relative velocities
  - Need to sample from correct distributions of energies of intramolecular degrees of freedom
- Collision coefficient from simulation is more than twice as large as the current value used in ACDC

#### Outlook

- Need to double (and triple) check that this is not an artefact
- Study collision / fragmentation probabilities of sulfuric acid clusters
- Study statistics of energy transfer between internal degrees of freedom during collisions

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