RECSEM REU 2017- Molecular Dynamics of Epoxy Resin Systems

Stephen Wu and Lam Tran Dr. Lonnie Crosby



Molecular Dynamics

• Molecular Dynamics

- Simulations that numerically solve the equations of motion for a system of particles as a function of time given a force field describing particle interactions.
- Microscopic length and time scales (nm/angstrom, ps/ns). Results "can be as accurate as needed".
- Relative to actual experiments, MD simulations can save time and money.
 Furthermore, they can oftentimes more easily reveal physical properties of a system (structure, microscopic interactions, etc.)





Epoxy Resins

• Epoxy Resins

- First developed in the 1930s and used today in a wide range of applications.
- High strength/durability, low shrinkage and excellent adhesive and insulative properties.
- Composed of a base resin (BADGE, cycloaliphatic, biobased, etc.) and a hardener/curative (amine type, alkali, catalytic) that confers additional properties.

Epoxy – Amine Crosslinking Reaction (ReaxFF_N/C/O/H)





Project Purpose and Goals-Why?

- Use MD simulations to examine the physical properties of the ProSet M1002 base resin/M2046 amine hardener system as a function of the components and the extent of crosslinking.
 - \circ ∂ V terms- Thermal expansion coefficient α (~ ∂ V/ ∂ T), Isothermal compressibility κ (~ ∂ V/ ∂ P)
- Are the adhesive properties of the system appropriate for usage in an industrial setting with aluminum surfaces?





Overview of Steps to Take

• Parameterization

- Are numerical values (like force constants) associated with charges, bonds, angles, torsions/dihedrals present and correct?
- Nonreactive Force Field Simulations-GROMACS
 - Are individual component molecules of the epoxy resin system behaving as expected? For example, is a molecule remaining compact and its atomic distances staying relatively constant?
 - Are the atoms with positive dipoles in polar solvents attracted to the oxygen with nonbonded electrons?
- Reactive Force Field Simulations-LAMMPS
 - Is the epoxy-amine crosslinking reaction dominant over other unwanted side reactions?
 - How do the properties of the system change depending on the extent of temperature-dependent crosslinking?

Past Progress

- Parameterization on each individual molecules
 - OPLS-AA topology of molecule
- Use quantum mechanics to optimize molecule structure with NWCHEM
 - Spatial coordinates Solution to Schrödinger equation
 - Partial charge Lowdin population analysis
- GROMACS sample test to calibrate and visualize the simulation system:
 - Water-solvated oxirane simulation
 - Methane-solvated oxirane simulation



Past Progress

- Parameterization on each individual molecules
 - OPLS-AA topology of molecule
- Use quantum mechanics to optimize molecule structure with NWCHEM (density function theory)
 - Spatial coordinates Solution to Schrödinger equation
 - Partial charge Lowdin population analysis
- GROMACS sample test to calibrate and visualize the simulation system:
 - Water-solvated oxirane simulation
 - Methane-solvated oxirane simulation



Past Progress

- Parameterization on each individual molecules
 - OPLS-AA topology of molecule
- Use quantum mechanics to optimize molecule structure with NWCHEM
 - Spatial coordinates Solution to Schrödinger equation
 - Partial charge Lowdin population analysis
- GROMACS sample test to calibrate and visualize the simulation system:
 - Water-solvated oxirane simulation
 - Methane-solvated oxirane simulation

Current Stages

- Parameterization CGenFF
 - VMD plugin fftoolkit bond, angle, dihedral parameters
- GROMACS sample test data analysis
 - Data analysis on free energy for sample simulation ensemble
- GROMACS Small-scale MD simulation
 - Single epoxy and Single hardener
 - Single epoxy and multiple hardener
 - Multiple epoxy and multiple hardener



Current Stages

- Parameterization CGenFF
 - VMD plugin fftoolkit bond, angle, dihedral parameters
- GROMACS sample test data analysis
 - Data analysis on free energy for sample simulation ensemble
- GROMACS Small-scale MD simulation
 - Single epoxy and Single hardener
 - Single epoxy and multiple hardener
 - Multiple epoxy and multiple hardener



Current Stages

- Parameterization CGenFF
 - VMD plugin fftoolkit bond, angle, dihedral parameters
- GROMACS sample test data analysis
 - Data analysis on free energy for sample simulation ensemble
- GROMACS Small-scale MD simulation
 - Single epoxy and Single hardener
 - Single epoxy and multiple hardener
 - Multiple epoxy and multiple hardener



Next Steps

- Small Molecule LAMMPS system construction
 - Translation from GROMACS to LAMMPS
 - Reaction between oxirane and methylamine (N/C/O)



- LAMMPS system simulation
 - Epoxy resin polymer reaction simulation
 - Volume change w.r.t Temperature T, pressure p and molecule number N

Next Steps

- Small Molecule LAMMPS system construction
 - Translation from GROMACS to LAMMPS
 - Reaction between oxirane and methylamine (N/C/O)



- LAMMPS system simulation
 - Epoxy resin polymer reaction simulation
 - Volume change w.r.t Temperature T, pressure p and molecule number N