

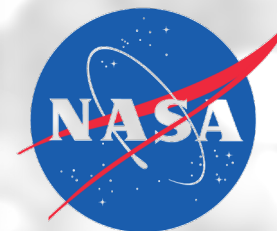
# REACTOR 2.0: Advanced Reaction Constraints and Automated Interaction Typing

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**Langley  
Research  
Center**

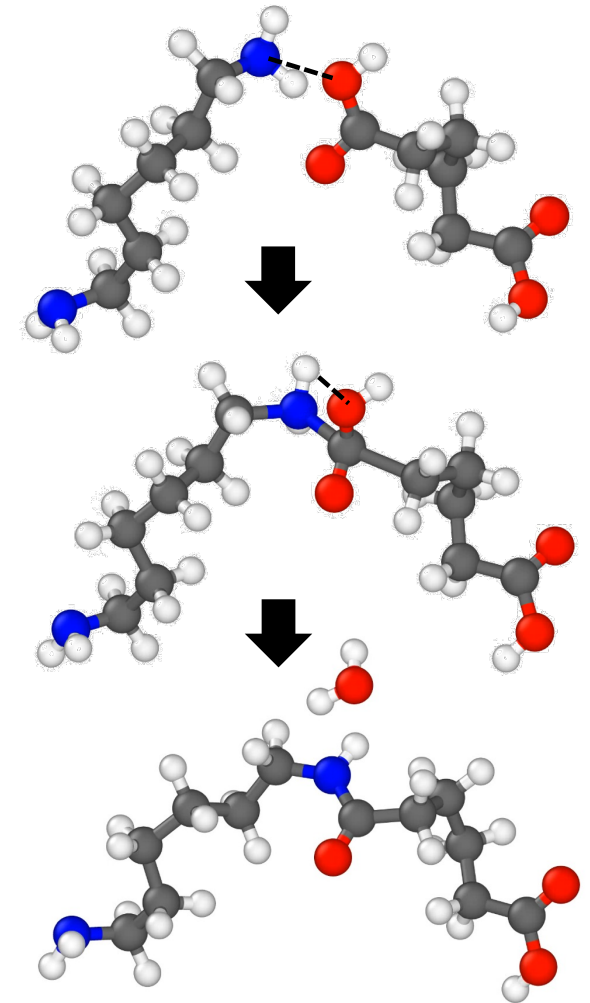
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unless otherwise indicated

# What is REACTER?

- A protocol for adjusting topology during classical MD
  - Unique scale for reaction modeling (fast, atomistic simulations)
  - Add/remove specific bonds, angles, dihedrals, and impropers
  - Modify all force field types as well as atomic charges
  - Supports any fixed-valence force field (PCFF, OPLS, etc.)
  - Reaction stabilization
- Parallel implementation in LAMMPS as ***fix bond/react***
  - User inputs: molecule templates of pre- and post-reaction topology
  - A map file relating atoms before and after the reaction

REACTER: A Heuristic Method for Reactive Molecular Dynamics.  
Gissinger, Jensen & Wise. Macromolecules 53, 22, 9953–9961 (2020).

*www.reacter.org*



# REACTER: Features added since 2019

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## 1) Streamlined input/output formats for reaction templates and data files

- String-based type labels allow easy creation of simulation-ready molecule templates
- Eliminates need to match numeric types between data files and reaction templates

## 2) Automatic interaction typing

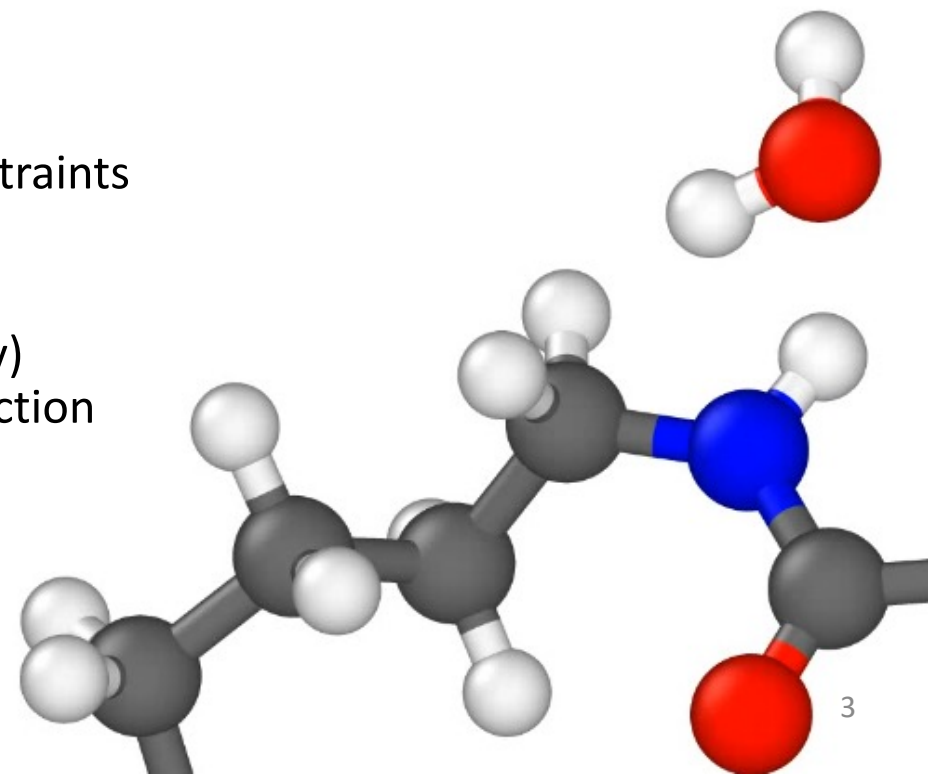
- Enabled by alphanumeric type labels that encode atom types

## 3) Advanced reaction constraints

- Arrhenius, root-mean-square deviation, and custom reaction constraints

## 4) General improvements/features:

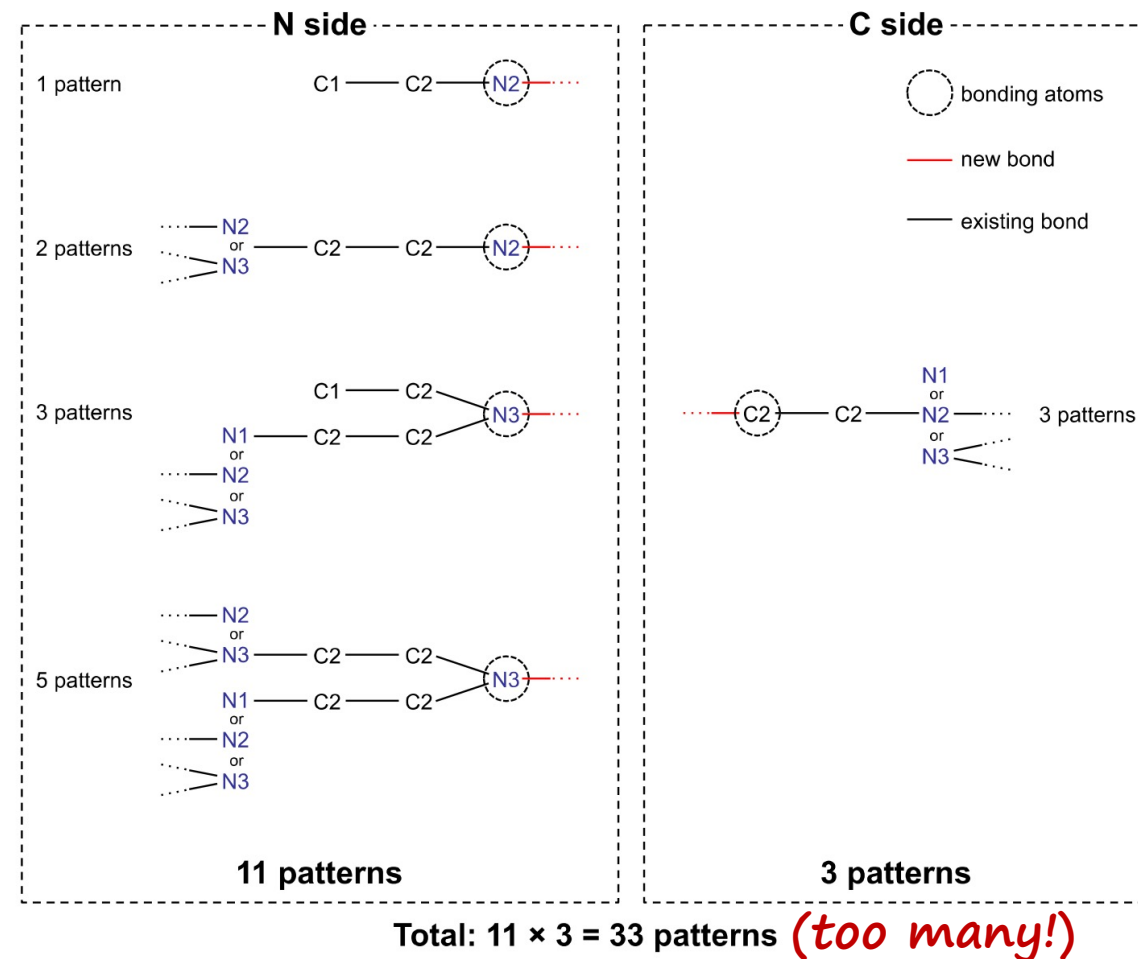
- Use variables for inputs (reaction cutoff, probability and frequency)
  - Use case: define reaction cutoff as a function of extent of reaction
- Specify reaction as inter- or intramolecular
- Update molecule IDs after reaction (default)
  - Makes *fix bond/react* compatible with other fixes, such as *fix gcmc*
- Create new atoms/molecules



# Beyond explicit reaction templates

- ‘Combinatorial explosion’ of templates needed when atoms surrounding reaction site are not constant
  - E.g., simple substitution of second and third neighbors required Kawagoe et al. to use 33 templates
  - Better solution needed than building a template for each possibility

*Three-stage solution: Labels, wildcards, and autotyping*

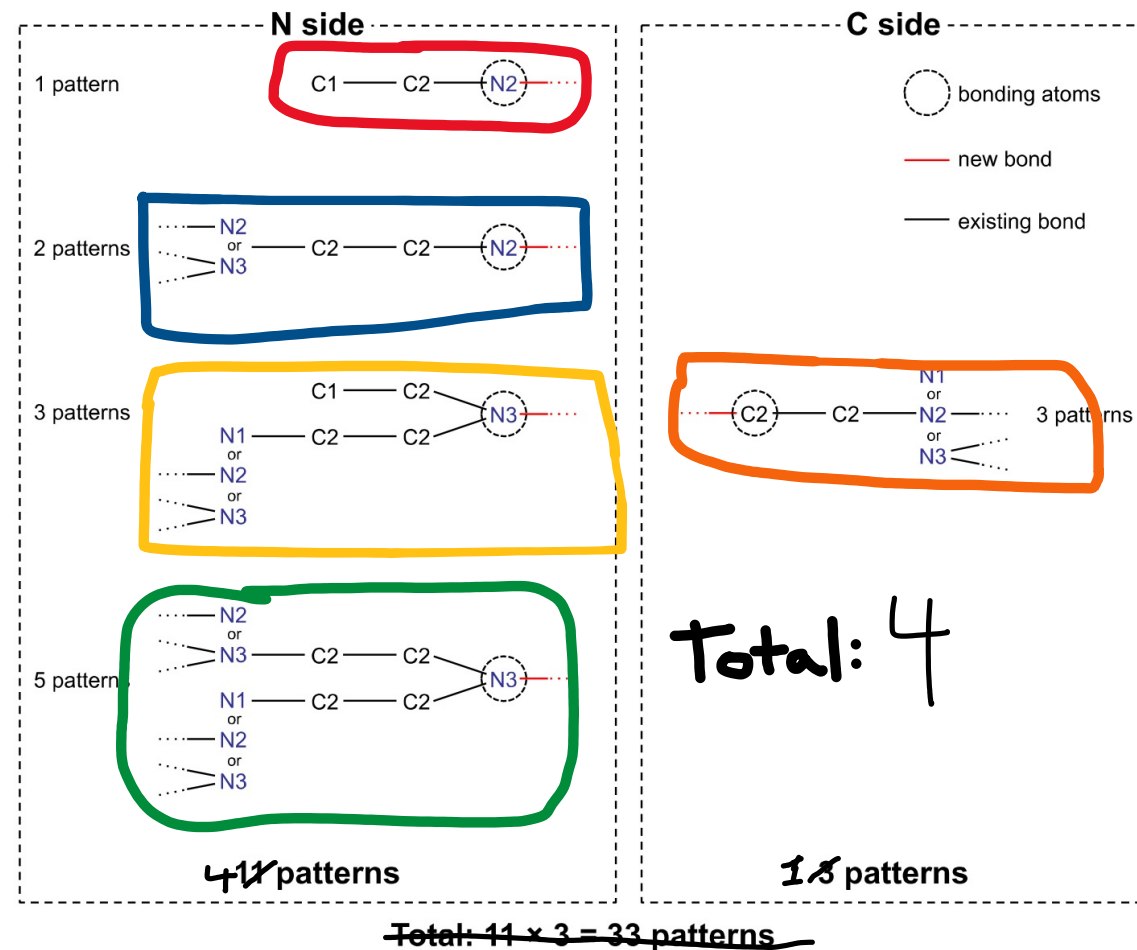


**Fig. S2.** Schematic of reaction patterns on the N (Nitrogen) side in the left panel and on the C (Carbon) side in the right panel.



# Labels, wildcards, and autotyping

- Kawagoe et al. example can be reduced to four templates.
- Type labels: mapping of each numeric type to an alphanumeric type
  - E.g., LAMMPS knows which atoms are 'N1'
  - Similarly, LAMMPS knows which bonds are [N1][C2], etc.
- Wildcards and autotyping:
  - E.g., use N\* for {N1, N2 or N3}
  - Automatically determine the types of new bonds, angles, etc. that involve wildcard atoms



**Fig. S2.** Schematic of reaction patterns on the N (Nitrogen) side in the left panel and on the C (Carbon) side in the right panel.

# Type labels: conventions and benefits

- Extra 'Type Labels' section in data file listing the type label mapping
  - Atom types in 'Atoms' section are not replaced directly, to minimize file size for large data files
- Molecule templates are small and benefit more from being human-readable
  - Atom, bond etc. types in 'Types', 'Bonds' etc. section are replaced directly
- Greatly simplifies/accelerates process of creating simulation-ready templates:
  - Automates task of coordinating types between templates and simulation

*In data file*

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Atom Type Labels

1 cp  
2 hc  
3 ct  
4 c=2  
...

*In molecule template*

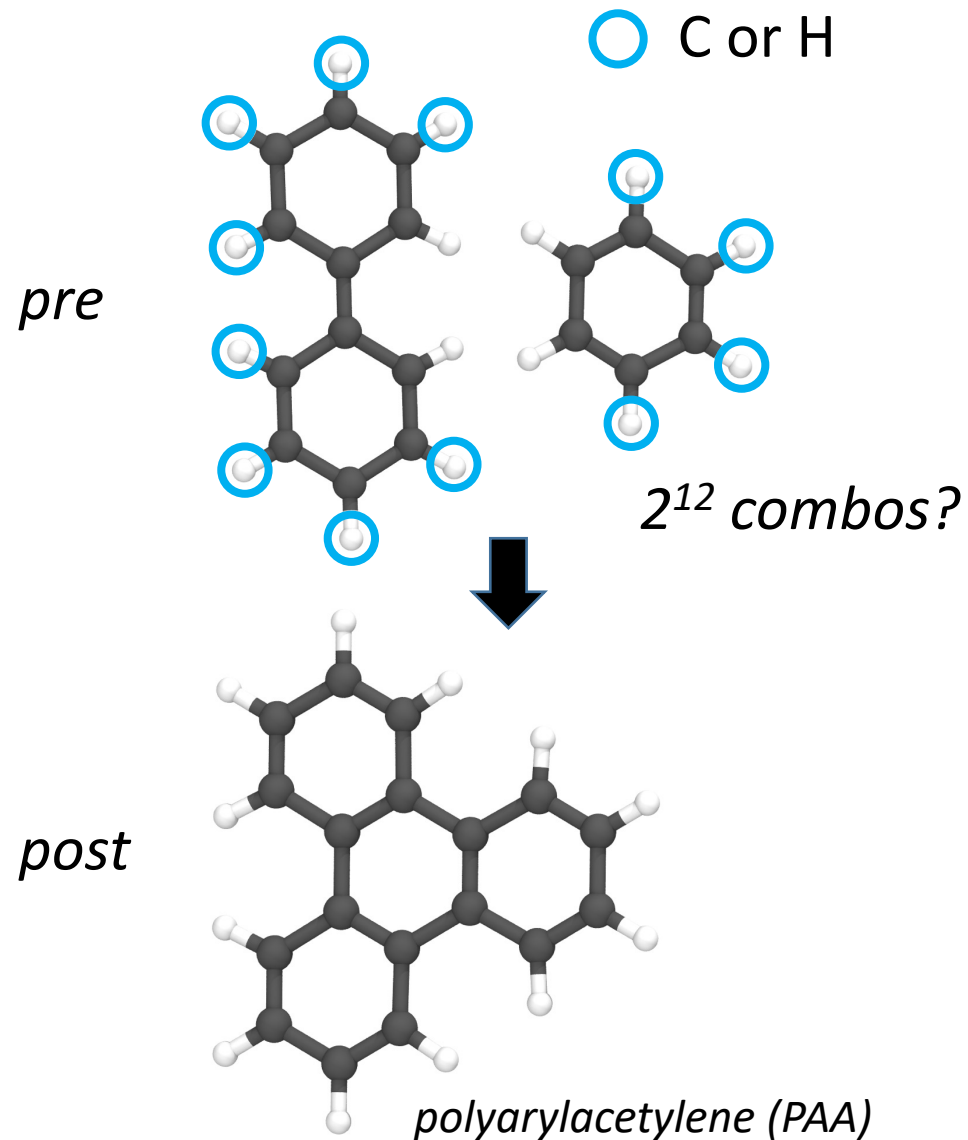
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Types

1 cp  
2 hc  
3 cp  
4 cp  
5 hc  
...

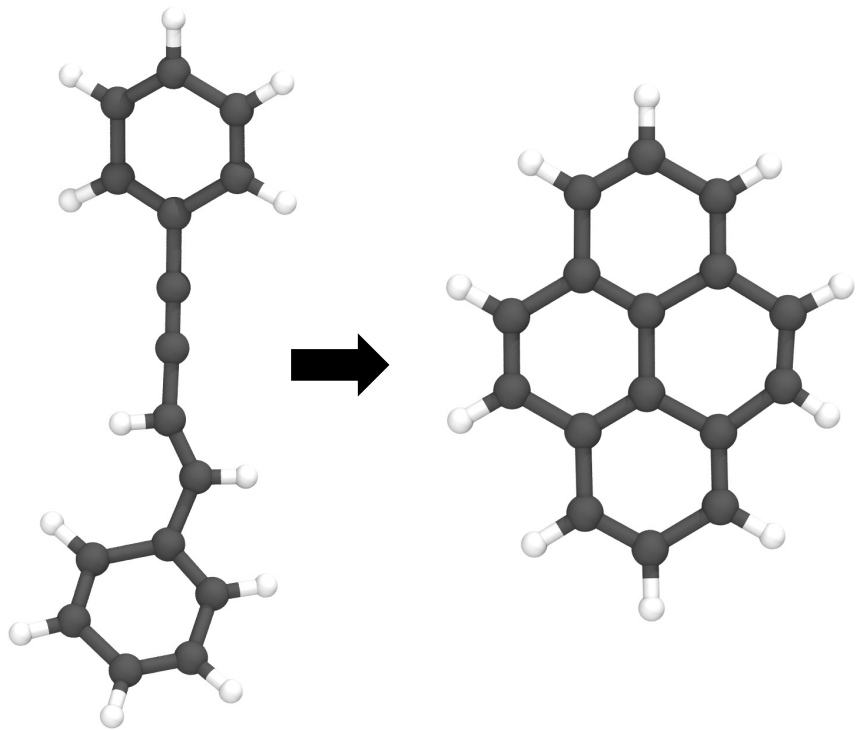
# Case study: Carbonization reactions

- Starting point: diethynylbenzene polymerized via linear and cyclotrimerization reactions
- End goal: Approximate model of the carbonization and/or graphitization process
- The problem is not tractable using explicit reaction templates
  - Combinatorial explosion when atoms connected to ring may be C or H
- Only one template needed when atoms near edge of template are defined as wildcards

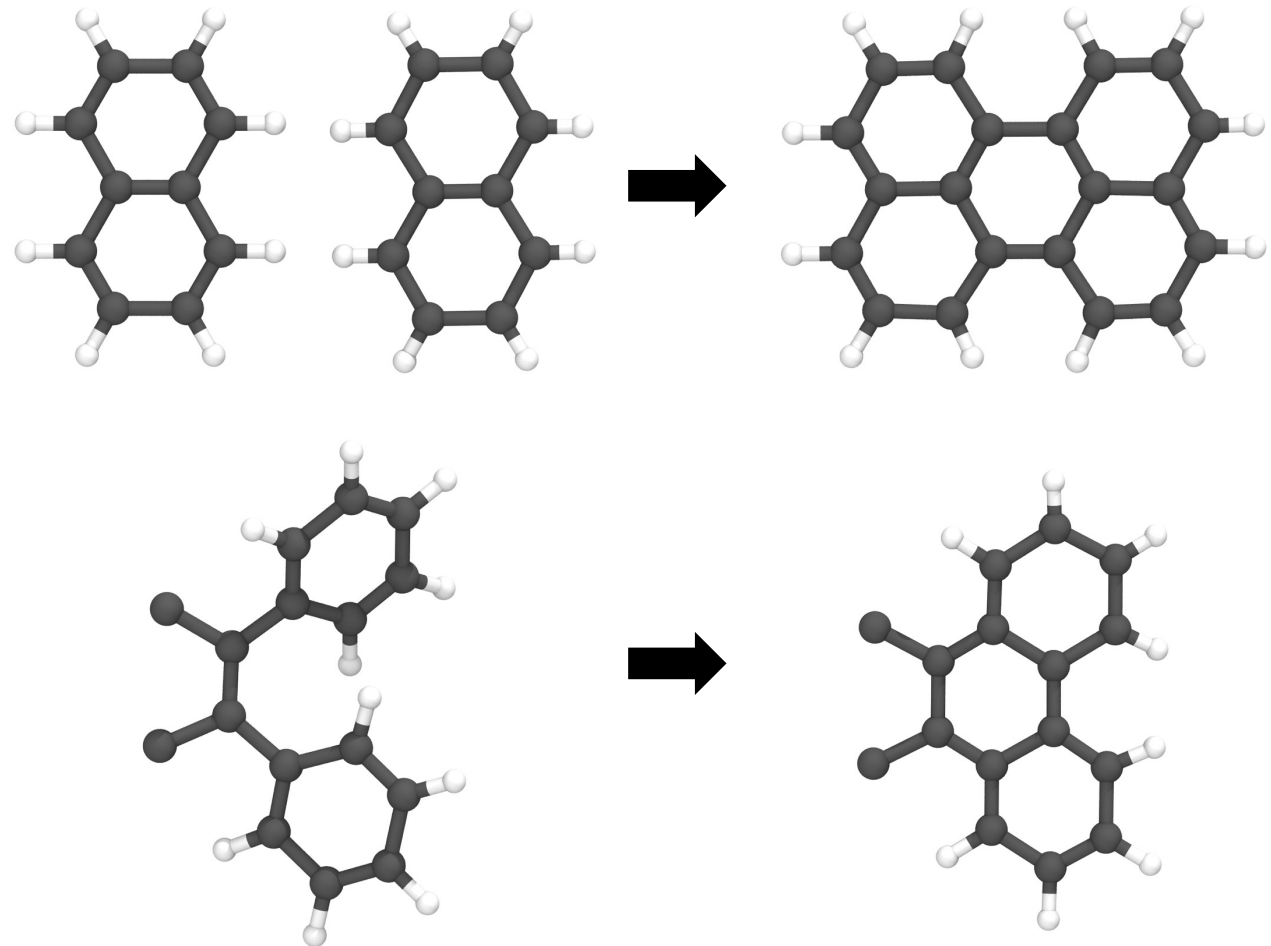


# Case study: Nanoribbon formation

- Define carbonization moves from polymerized PAA motifs, as well as ring-closing and Diels-Alder reaction



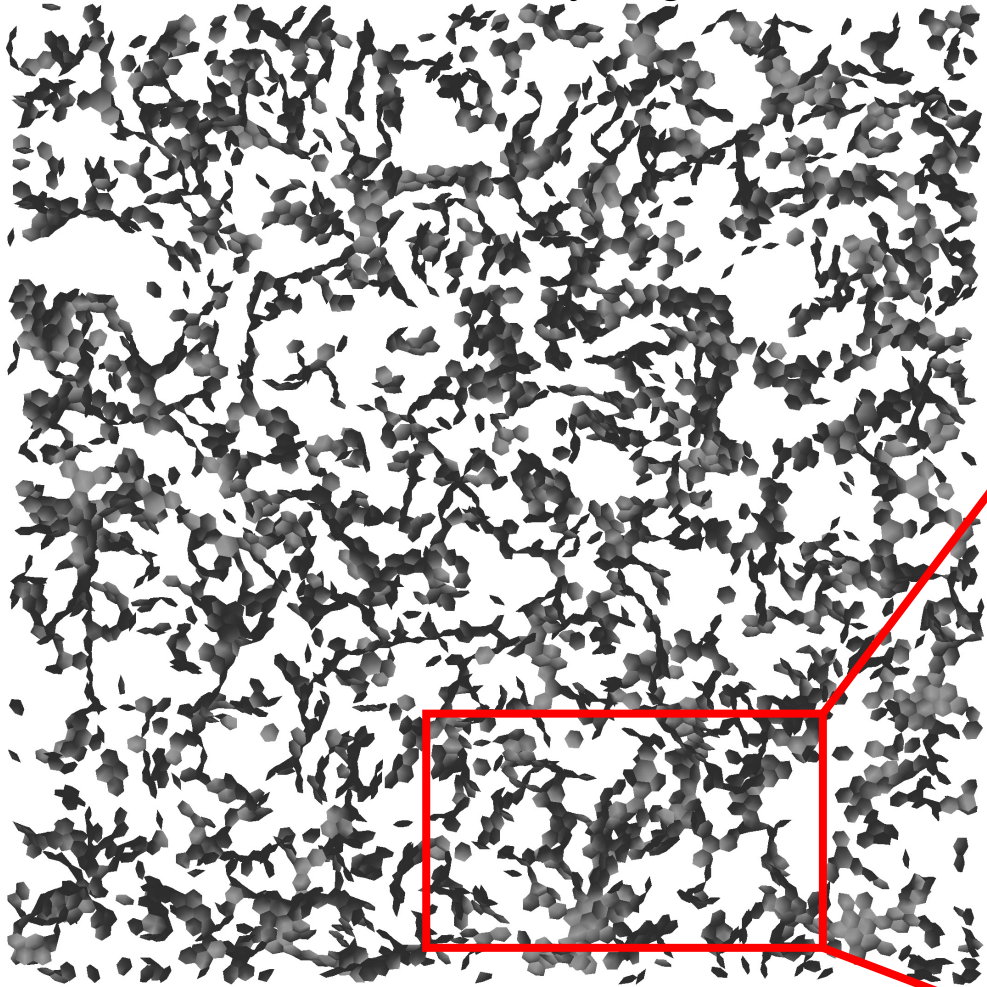
Ring creation/fusing rearrangement  
move for the linear Strauss coupling





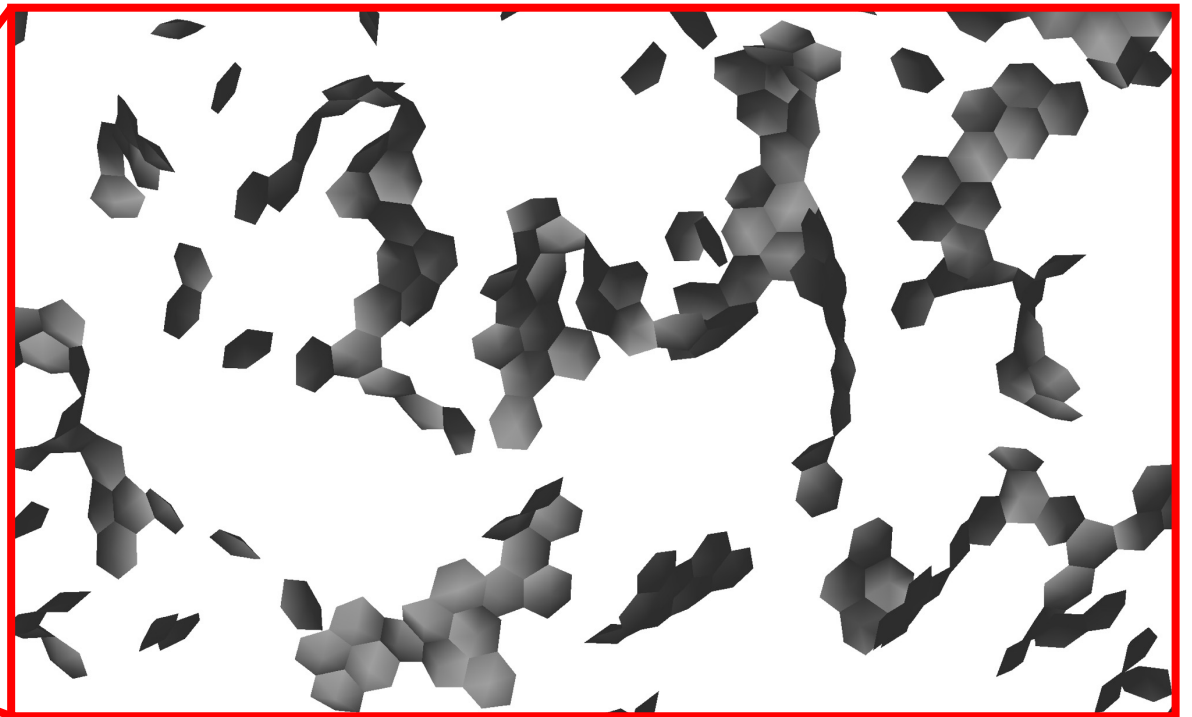
# Case study: Nanoribbon formation

Slice is 7 Å thick; only rings shown

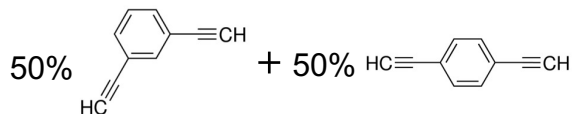


After ~99% polymerization and ~14.5K carbonization reactions

Expedient large-scale model for early-stage carbonization of polyarylacetylene (200K atoms)

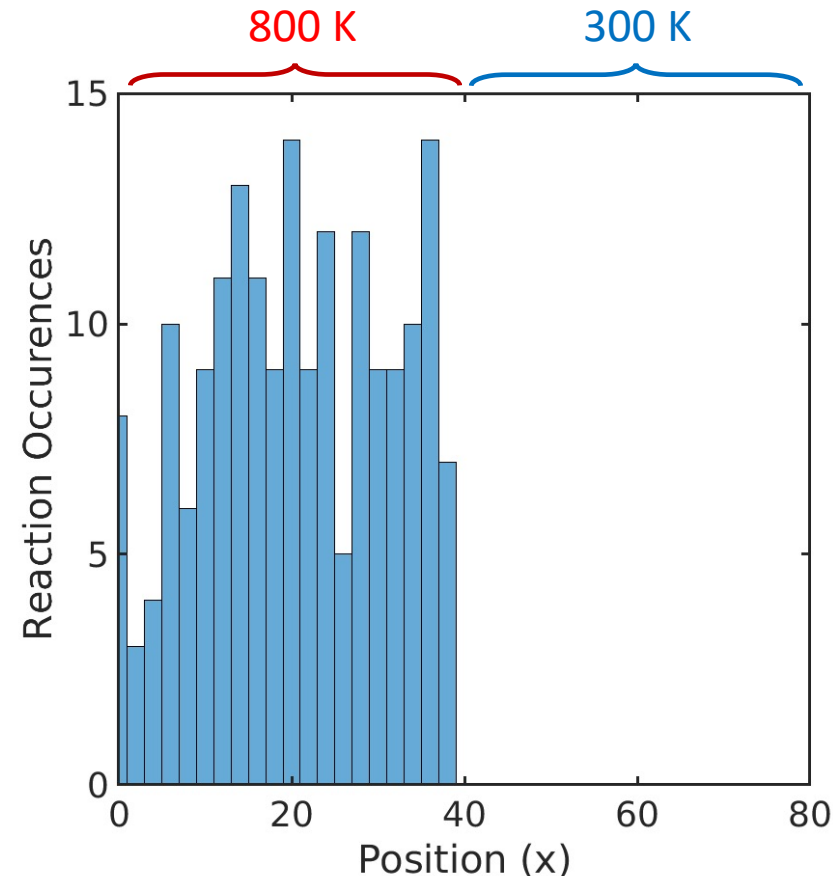
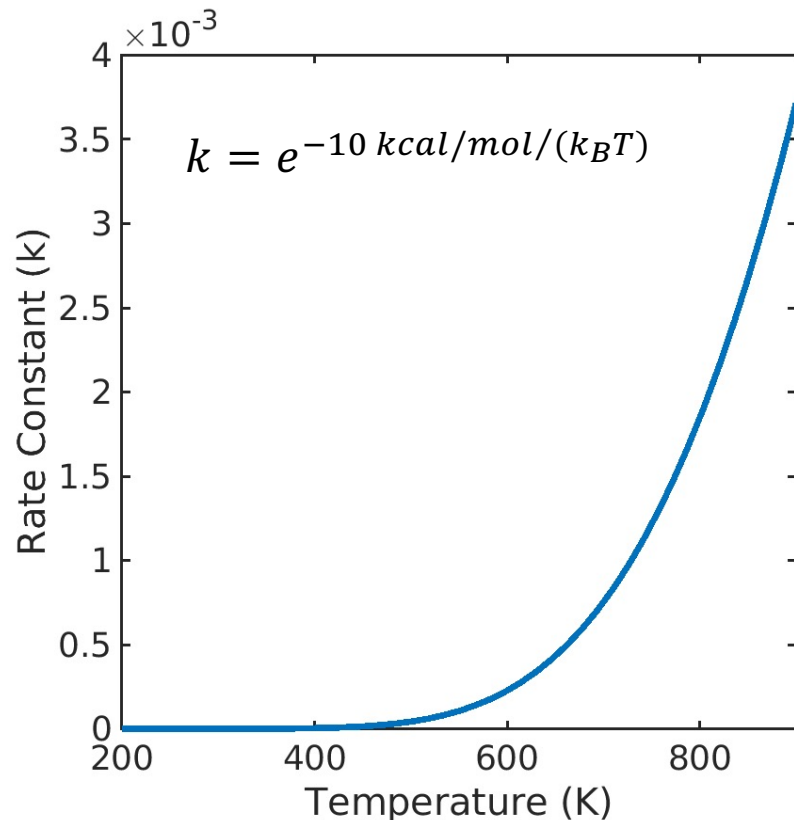


Density: 1.0 g/cm<sup>3</sup>  
Box Size: 13.8 nm



# Arrhenius Reaction Constraint

- Uses local temperature to enforce effective activation energy
  - Temperature-dependent Arrhenius equation:  $k = AT^n e^{\frac{-E_a}{k_B T}}$
  - Reaction probability is proportional to calculated rate constant

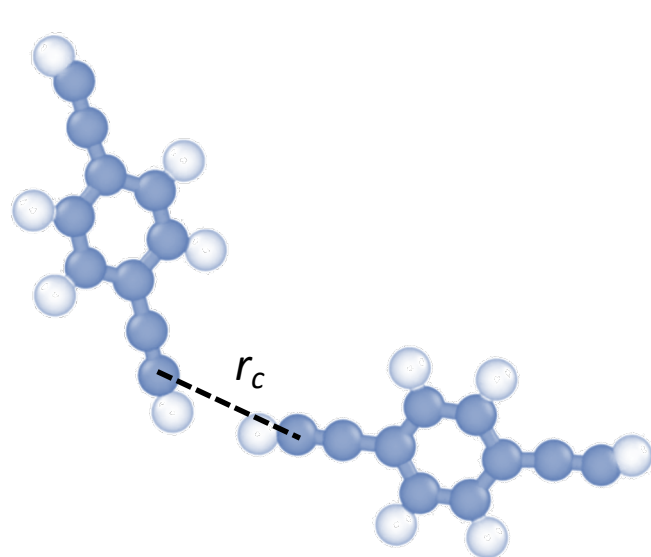


Nylon melt with  
Arrhenius constraint:  
80x80x80 Å  
35,000 atoms  
10,000 fs  
Reaction cutoff: 3.5 Å

Simulation box  
thermostated with a  
hot half and a cold half

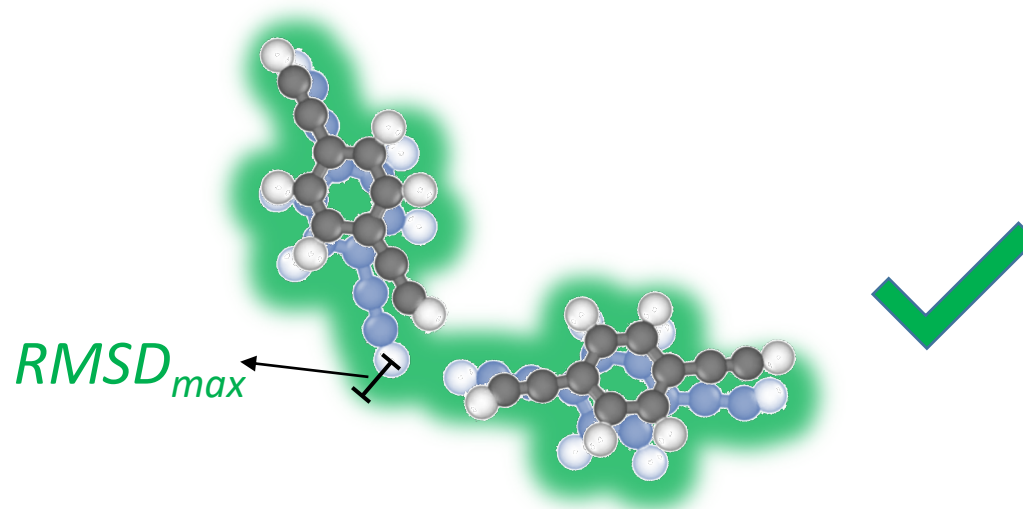
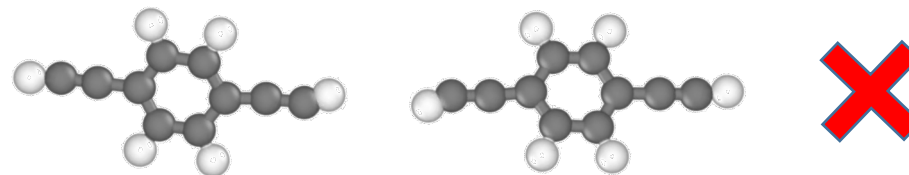
# RMSD Reaction Constraint

- Root-mean-square deviation (RMSD): a metric for similarity of molecular structures
- Enforce a maximum RMSD between template and local reaction site atoms
  - Calculated after optimal translation/rotation of pre-reaction template
- Orientation that allows reaction can be empirical or based off QM results
- A step toward more complex energy surfaces for reactive sites



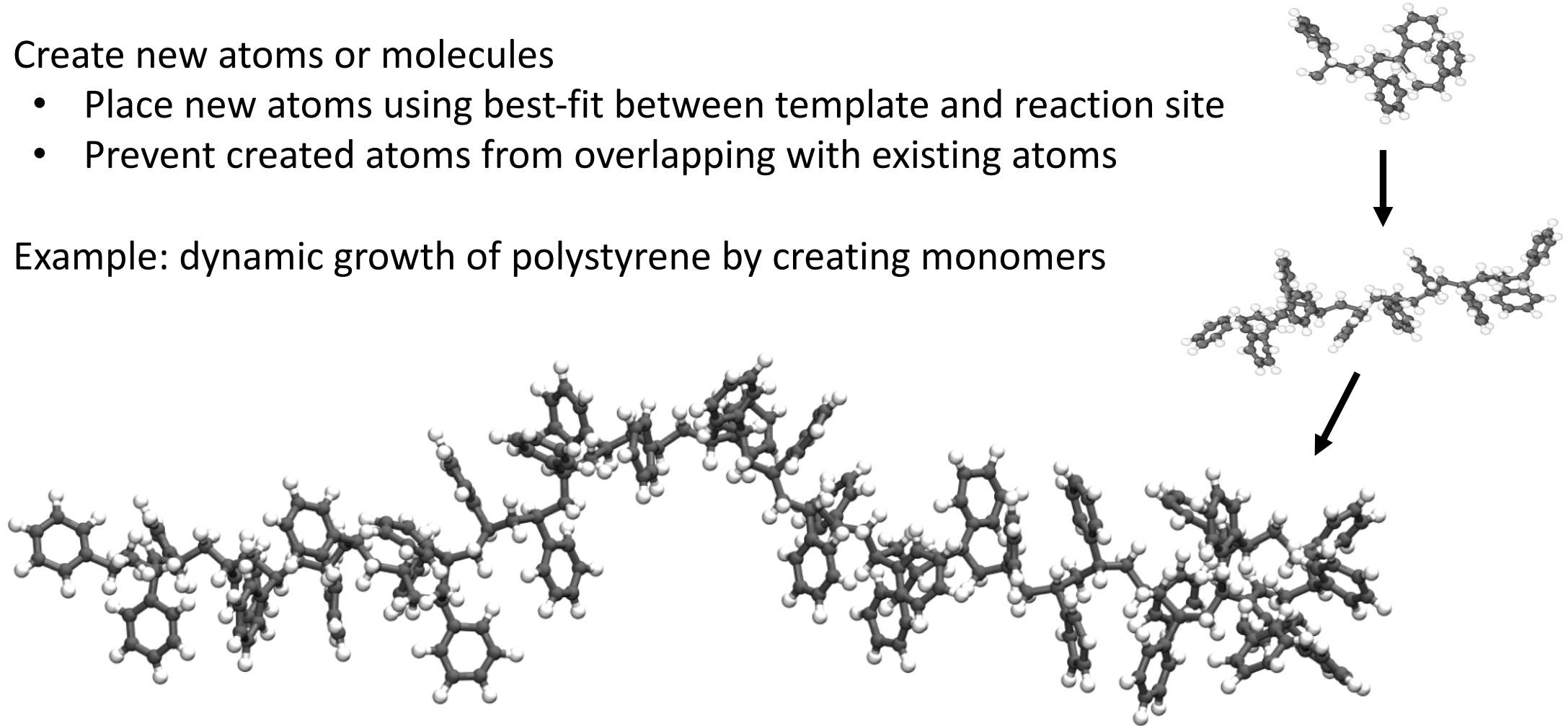
Pre-reaction Template

Simulation  
Atoms



# Create Atoms (and their Bonds, etc.)

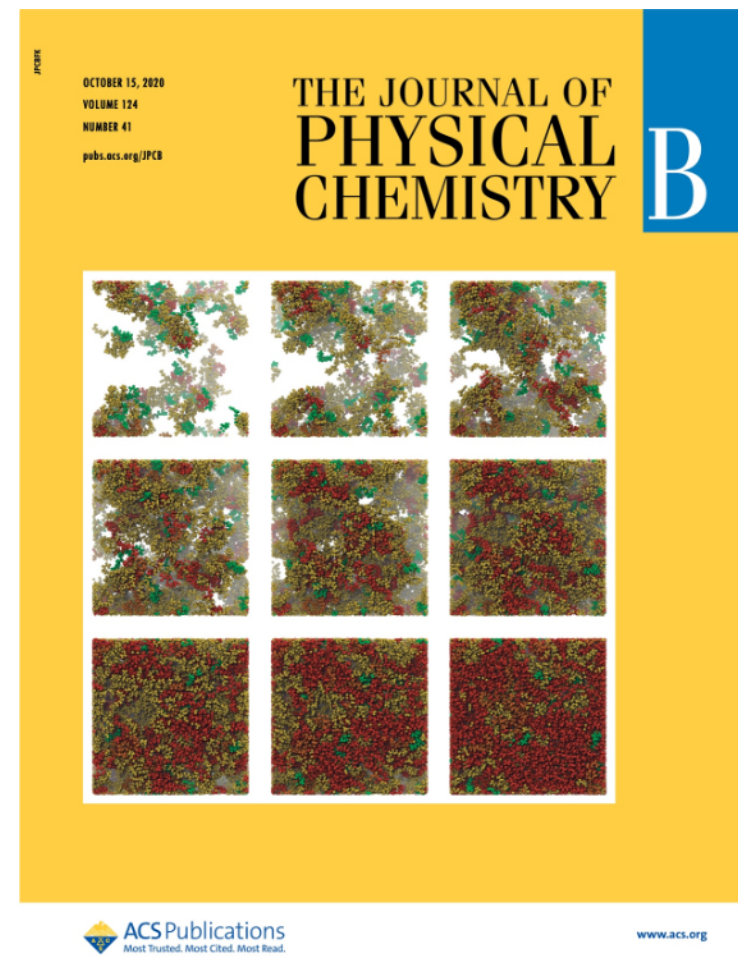
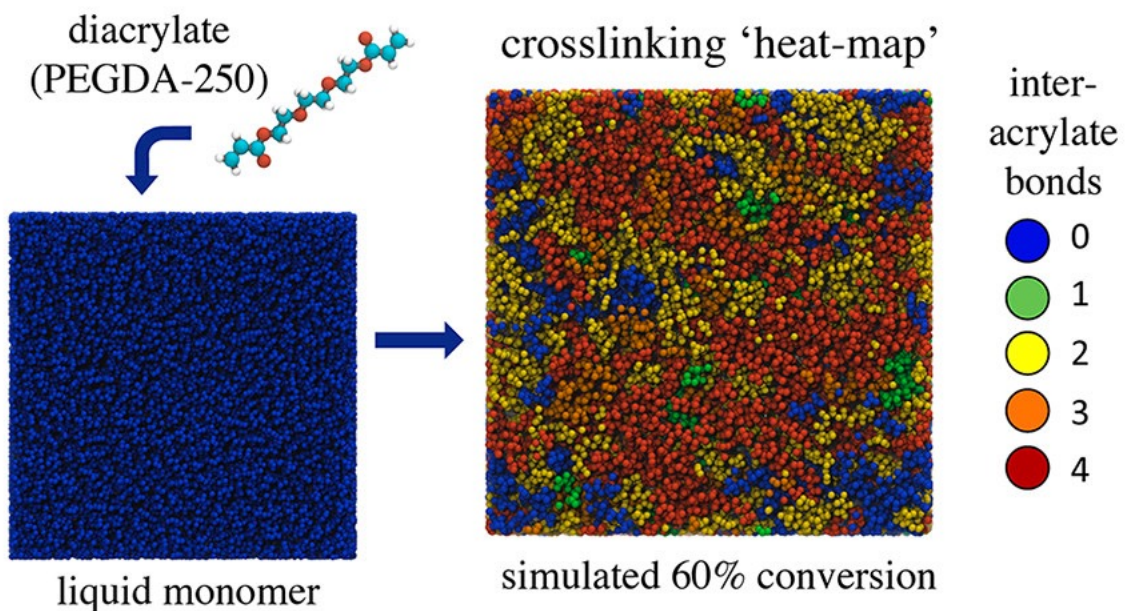
- Create new atoms or molecules
  - Place new atoms using best-fit between template and reaction site
  - Prevent created atoms from overlapping with existing atoms
- Example: dynamic growth of polystyrene by creating monomers





# What else can REACTER be used for?

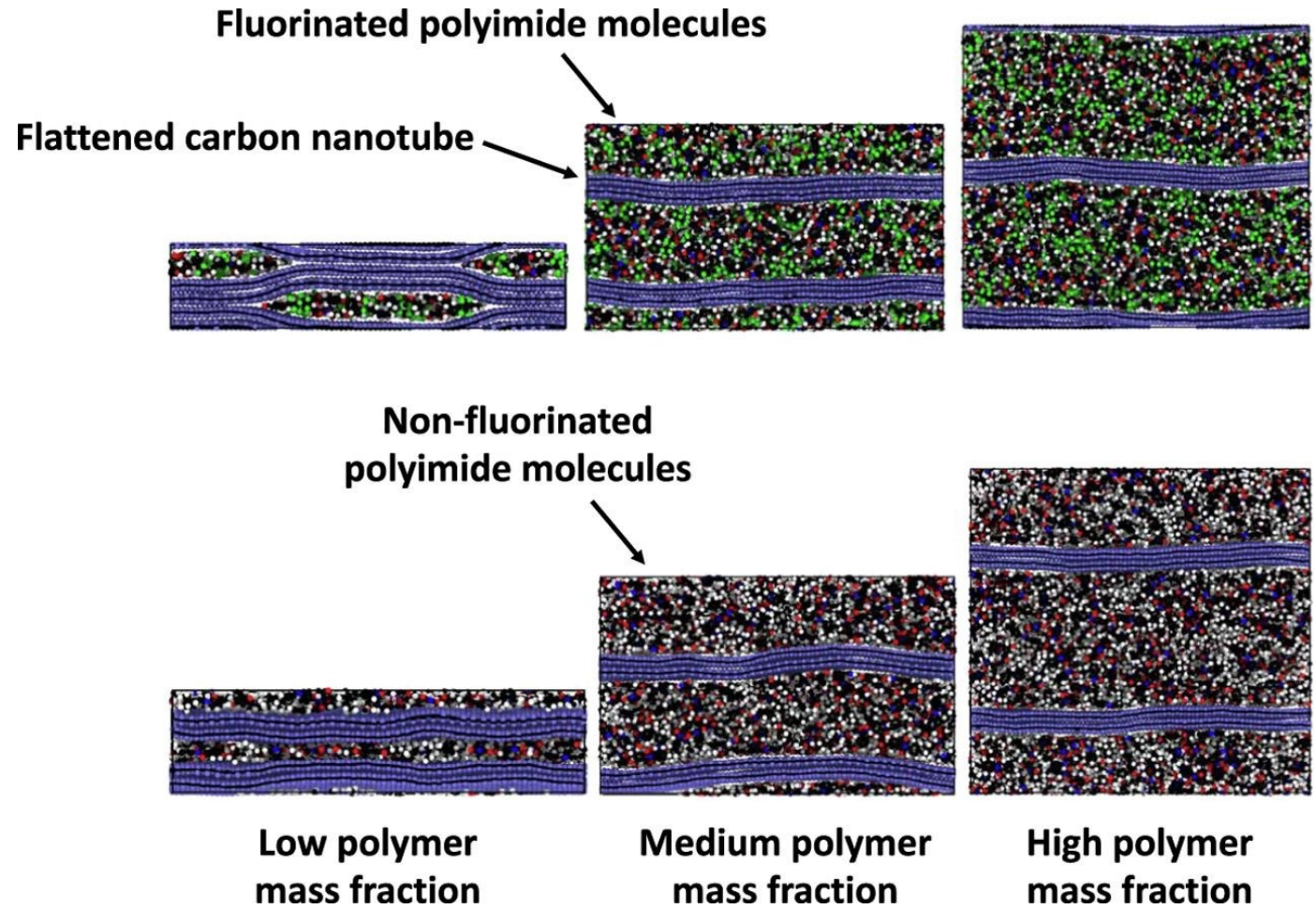
- Modeling polymerization of photopolymer resins
- Reactive MD revealed the dependence of gel points and cyclic content on various poly- and difunctional acrylates



Karnes *et al.* On the network topology of cross-linked acrylate photopolymers: A molecular dynamics case study. *The Journal of Physical Chemistry B*, 124(41), 9204-9215 (2020).

# What else can REACTER be used for?

- *Fix bond/react* can act on a subset of the simulation, when nonreactive phases are present
- E.g., modeling the *in situ* polymerization of the matrix in nanocomposites
- The properties of the interface between polyimides and flattened CNTs depend on the polyimide chemistry (fluorinated vs non-fluorinated)



Patil *et al.* Interfacial characteristics between flattened CNT stacks and polyimides:  
A molecular dynamics study. *Computational Materials Science*, 185, 109970 (2020).



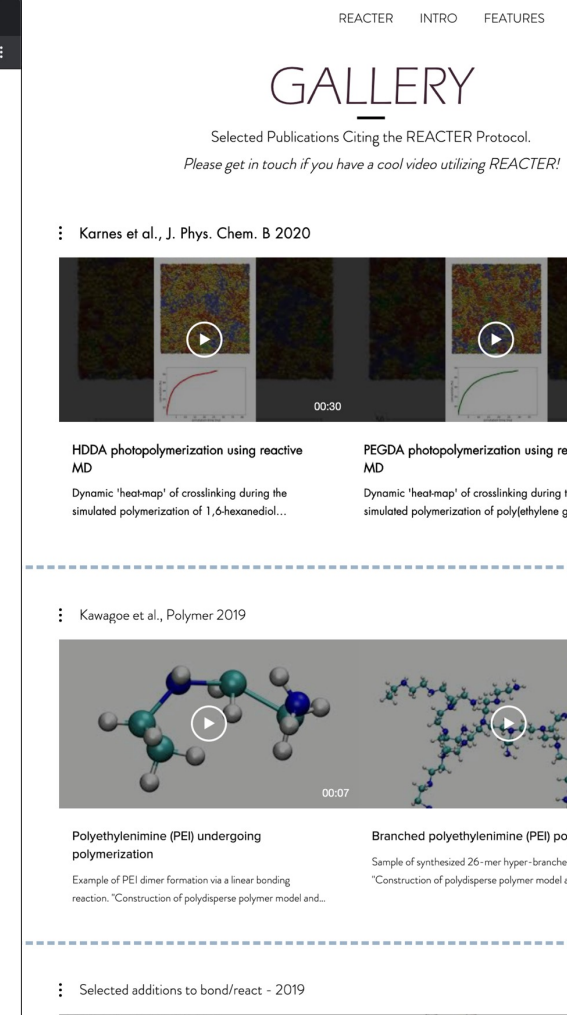
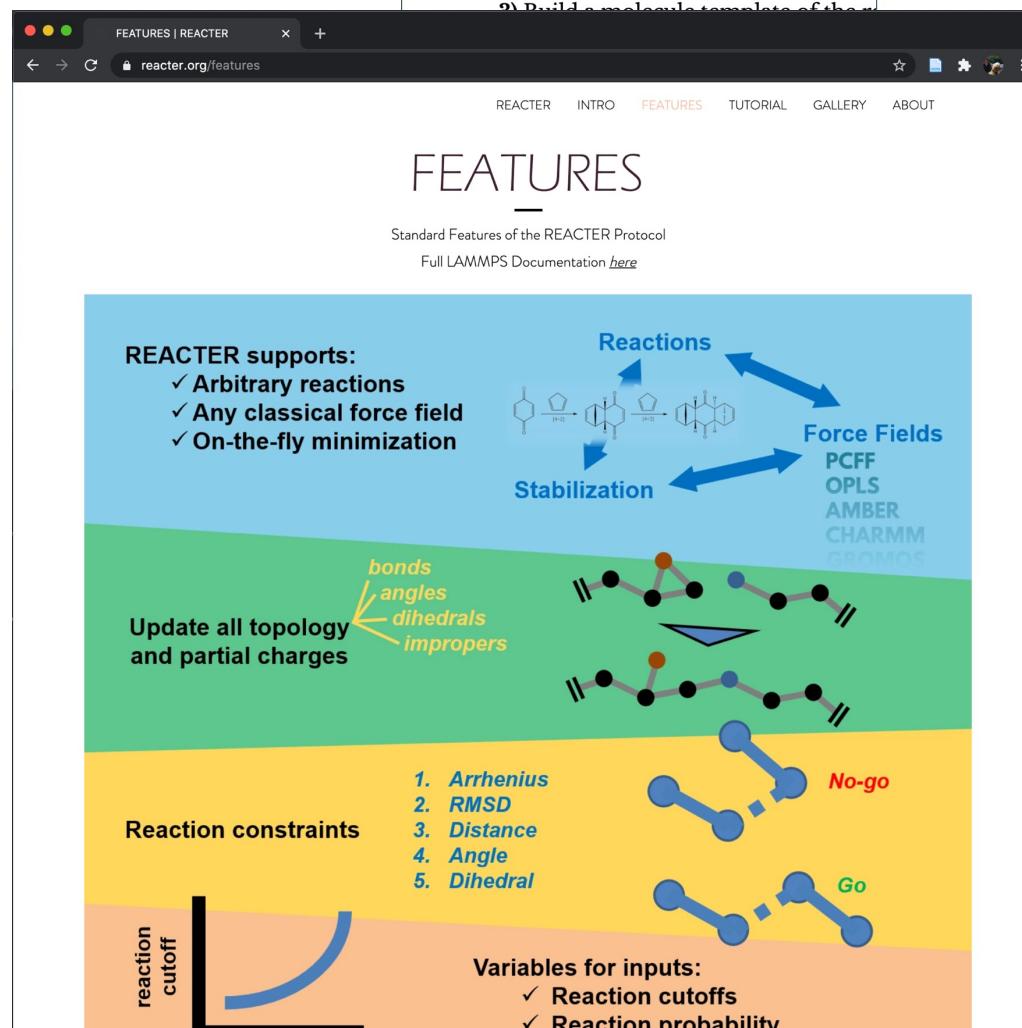
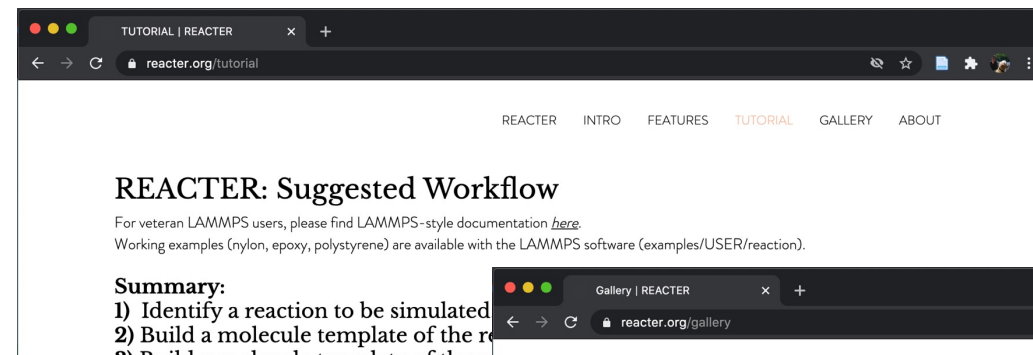
# Where to learn more?

Visit ***www.reacter.org*** for:

- Up-to-date features
  - Tutorials
  - Videos
- Links to related material

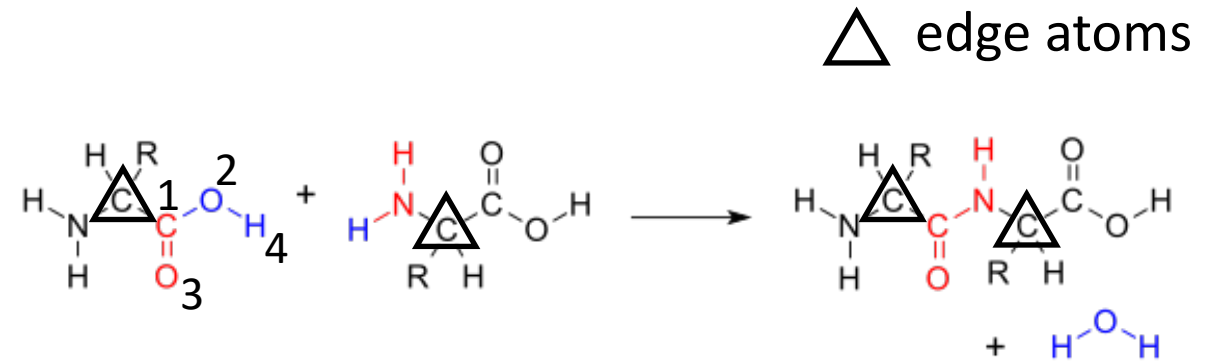
If you have a cool video or image that utilizes REACTER, send me an email to be added to the gallery

Also, check out the new AutoMapper tool (Matt Bone, University of Bristol) for a streamlined way to create reaction templates



# Outlook: The potential of type labels

- Type labels are invaluable for rapid creation of simulation-ready data files and reaction templates
- Library of reactions or classes of reactions
  - Templates could be applied directly to many force fields via type labels, wildcards and autotyping
- Existing databases, often expressed in SMILES or SMARTS format, can be converted into templates
- Incorporate reaction constraints such as RMSD if known (e.g., via DFT)



*Molecule template in library: aa.pre*

Types

1 C\*  
2 O\*  
3 O\*  
4 H\*

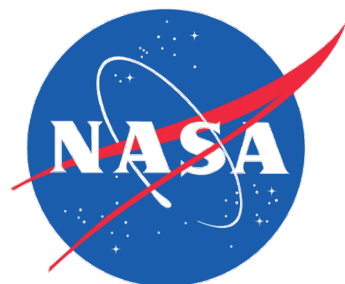
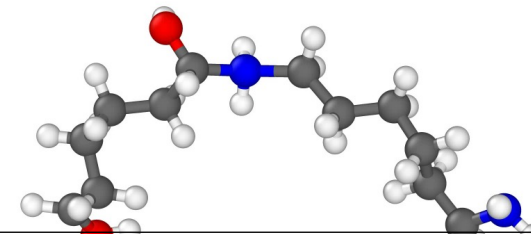
...

*In LAMMPS input script*

fix 1 all fix bond/react &  
react rxn1 all 1 0.0 3.0 aa.pre aa.post aa.map



*Thank You!*



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