

A Quantum Chemistry Simulation Strategy for Dynamic Network Polymers Multiscale Mechanochemistry and Mechanobiology (2019) Michael Buche¹, Zachary Sparrow², Yuval Vidavsky³, Robert DiStasio², Meredith Silberstein^{1,3} ¹Theoretical and Applied Mechanics, Cornell University, Ithaca, NY, USA

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-Ni

2

1.5

Tensile Force [nN]

-Ni-2Im

–Ni-2Py

—Ni-2Pipe

-Ni-2DMA

2.5

Ni-2MeIm

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Organometallic Crosslinking Compounds

Dynamic crosslinking structures such as organometallic compounds can be used in place of covalent bonds in order to retain considerable crosslinking strength, while enabling improved material properties such as increased toughness and processability. In order to tailor or predict mechanical properties of the bulk material, the dynamic crosslinks must be understood at a fundamental level.

Differing Ligand Amount



Presented here is a step-by-step density functional theory simulation-based strategy we use to quantify the mechanics of a particular organometallic complex, nickel(II) acetate hydrate, that can be used as a polymer crosslinker. The complex is analyzed before and after adding ligands, and while varying the ligand number and type. The mechanical stiffness and free energy barrier to breaking the complex change dramatically when differing the ligand amount, but the two change less when varying the ligand type.



Reaction Pathway (FSM) and Transition State Identification (TS)

 $\Delta G = 5.06 \text{ kcal/mol}$

0.8

mol

Ba

En

0

0.5

- Ene 2.5 0.5 Tensile Force [nN]
- A step-by-step quantum chemistry simulation method was presented to analyze the mechanics of dynamic crosslinking structures.
- The analysis predicts that each ligand added to the crosslinking structure will significantly diminish both its mechanical response and energy barrier to breaking.
- The same analysis predicts more subtle changes in the mechanical response and energy barrier of the structure when changing the type of ligand.
- Further analyses may be necessary if non-bonded interactions of the complex with the polymer dominate and change appreciably between different ligands.

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