

tutions in epoxy-amine networks and found a significant change (*para*- > *meta*- > *ortho*-) in the glass transition temperatures (T_g) of the networks due to minor structural changes in the monomer structure. These afore-mentioned studies are based on the use of structural isomers of multifunctional small molecule monomers.

Unlike small molecules, modulation of the spacing, regioselectivity, and sterics of a multifunctional unimolecular oligomer (*i.e.*, a macromer) is non-trivial and often requires extensive synthetic manipulations. Due to these synthetic challenges, studies into how macromer structure affects network connectivity and material properties are lacking. Addressing this question requires studying network formation with well-defined and easily accessible functional macromers. Sequence-defined macromers fill a void that exists between precise small molecule monomers and large homo- and copolymers. When assembled from functional monomers, these well-defined unimolecular macromers can serve as versatile building blocks for a wide range of polymeric materials and networks.^{22,23} However, simple gram-scale synthesis is required to understand how the structure and composition of a macromer can affect network topology and material properties.

Recently, our research group developed a robust platform for the synthesis of sequence-defined polyurethane macromers (SD-PUMs)²⁴ using a sustainable feedstock, vanillin, and high-yielding reactions. The SD-PUMs have modifiable pendant groups and a rigid aromatic vanillin-based backbone. The initial report by Hoff *et al.*²⁴ shows the importance of sequence on the properties of cross-linked networks. This work inspired further investigations, carried out in this study, where we investigate the effect of regioisomeric backbone and end-group substitutions on network topology and material properties. In this study, we show that the SD-PUM platform can be used to create and tune network properties by varying backbone structure and end-group composition. To study how changes in backbone and end-group structure affect network topology and properties, four macromers (two pairs of regioisomers) were

synthesized and cross-linked using multivalent thiols. The thermal and mechanical properties of the ensuing networks were measured and compared. The results reported herein illustrate how subtle changes in backbone and end-group structure, *via* the use of positional isomers, significantly impacts network performance.

Results and discussion

Synthesis of SD-PUMs

The SD-PUM synthetic platform involves the use of an inexpensive biorenewable compound, vanillin, to create SD-PUMs on the gram-scale (Fig. 1). Vanillin is converted to a divanillin carbonate monomer with two reactive sites: the aldehyde that undergoes a reductive amination reaction to generate a secondary amine and a carbonate that is reacted with the secondary amine to form a carbamate. The synthetic process involves sequential reductive amination and carbamate formation reactions, followed by a simple aqueous workup after each step (Fig. 1). Performing the assembly and purification steps in this manner enables the production of precision SD-PUMs at scale with only a round-bottom flask, a separatory funnel for stepwise purification, and a rotavap for solvent removal.

In this study, allylamine is used as the pendant group for all macromers. Macromers with 1,4-backbone substitutions were made using divanillin carbonate for the carbamate formation reaction with a secondary amine. However, for the macromers with a 1,3-backbone substitution, a 4-nitrophenyl activated carbonate of isovanillin was used. This was done due to the high pK_a of isovanillin (~ 8.8), which makes it a poor leaving group relative to vanillin ($pK_a \sim 7.4$) or nitrophenol ($pK_a \sim 7.2$). The final capped macromers were purified using column chromatography. The individual reaction steps had recovered yields of 70–85%. The recovered yields of the purified macromers were in the 16–22% range at the gram-scale. Each oligomer is named according to its monomeric units,

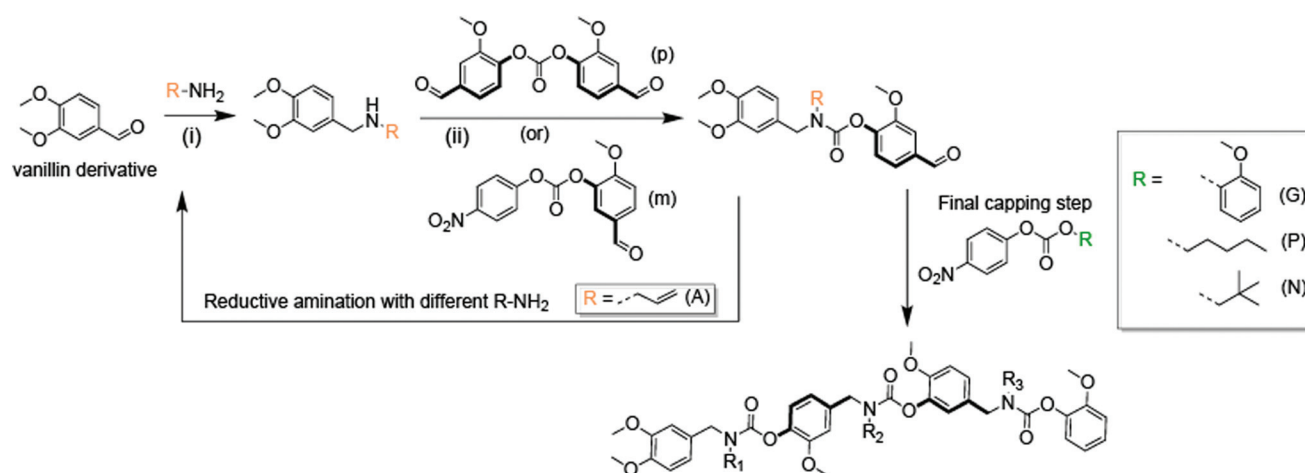


Fig. 1 General synthesis scheme for sequence-defined polyurethane macromers. (i) MeOH, NaBH₄, room temperature, (ii) triethylamine, CH₃CN, 50 °C. (A) – allyl, (G) – guaiacal, (P) – pentyl, (N) – neopentyl.

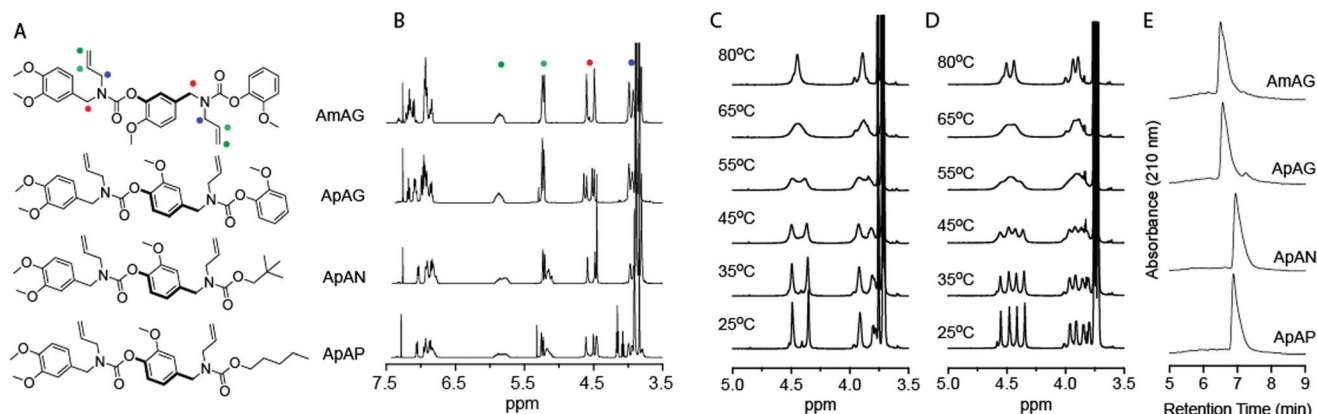


Fig. 2 (A) Chemical structure of macromers evaluated in this study; (B) ^1H NMR spectra of macromers; VT NMR of (C) AmAG and (D) ApAG; (E) retention time of the macromers on a C4 reverse-phase column.

where “A” represents an allyl pendant group, “m” represents an isovanillin backbone unit and “p” represents a vanillin backbone unit. The end-groups used to cap the macromers were guaiacol (G), pentanol (P), and neopentanol (N). The four macromers that were studied were AmAG, ApAG, ApAN, and ApAP (Fig. 2A). The AmAG and ApAG macromers are regioisomers in their backbone, while ApAN and ApAP are regioisomers at their end-group. The progression of the SD-PUM synthesis was followed by ^1H and ^{13}C NMR spectroscopy and liquid chromatography-mass spectrometry (LC-MS) (ESI Fig. 1–4†). The final ^1H NMR spectra of the macromers after purification *via* column chromatography is shown in Fig. 2B. The full spectra of all macromers with complete integration can be found in ESI Fig. 5–8.†

Structural differences of regioisomeric SD-PUMs

The ^1H NMR spectra of the AmAG and ApAG regioisomers shared similar spectral features with notable differences in the splitting patterns of the benzylic protons centered around 4.6 ppm and the allylic protons at 4 ppm (Fig. 2B and ESI Fig. 5 and 6†). This hinted at differences in conformation or electronic environment. To probe the solution conformational dynamics of each macromer, a two-dimensional rotating frame Overhauser effect spectroscopy (^1H – ^1H ROESY) experiment was conducted for both macromers at 0 and 200 ms mixing times (ESI Fig. 9 and 10†). The presence of off-diagonal exchange peaks in the 200 ms spectra confirmed conformational exchange between the different positions due to rotation around the C(O)–N bond. The AmAG macromer only had two diagonal peaks at the benzylic protons, whereas ApAG had four diagonal peaks at the benzylic protons at 0 ms. At 200 ms, the AmAG macromer had a single set of exchange peaks while the ApAG macromer had two sets of exchange peaks (ESI Fig. 9B and 10B†). The rate constants for conformational exchange were calculated using EXSYcalc, and both macromers had similar rate constants ($k_1 = 8.23 \text{ s}^{-1}$ for AmAG and $k_1 = 8.56, 8.42 \text{ s}^{-1}$ for ApAG). Further studies *via* heteronuclear single quantum coherence spectroscopy (^1H – ^{13}C HSQC) (ESI Fig. 11 and 12†) confirmed the presence of eight

methylene protons (four benzylic and four allylic) for both macromers. The four benzylic protons in ApAG were separated both in the ^{13}C and ^1H spectra, indicating that the two benzyl groups in the macromer were in different environments. However, the benzylic protons in AmAG were separated only in the ^{13}C spectra suggesting that the two benzyl positions in the macromer were in identical chemical environments. This observation speaks to how a single positional change in the macromer backbone can alter the molecular electronic environment.

Variable temperature NMR (VT-NMR) was also used to confirm the structural dynamics of the macromers (Fig. 2C and D). As the temperature was increased from 25 to 80 °C, the two benzylic proton peaks at 4.6 ppm coalesced into a broad peak due to the rapid exchange between conformations brought about by rotation around the C(O)–N bond. The coalescence occurred at 65 °C for both macromers (Fig. 2C and D). The presence of two distinct peaks at 4.6 ppm at 80 °C in the ApAG spectra concurs with the conclusions from HSQC spectra, which suggests that the two benzyl groups in ApAG are in different environments. Similarly, the presence of a single peak at 4.6 ppm at 80 °C in the AmAG spectra confirms the identical environments for the two benzylic positions.

The retention times of the macromers on a C4 reverse-phase column is shown in Fig. 2E. The retention time is a measure of relative hydrophobicity, and the data shows that the macromers with the aliphatic end-groups have higher retention times (hence, more hydrophobic) than those with aromatic end-groups. The data also shows that the regioisomers, in both cases, have similar retention times.

Effect of end-group composition and regioisomerism on macromer physical properties

To study the effect of regioisomerism on the physical properties of the macromers, viscosity measurements were performed using a parallel plate rheometer at 45 °C with a constant shear rate of 0.1 s^{-1} over 60 seconds. The measured viscosities were compared to motor oil and honey in order to provide a physical frame of reference using common house-

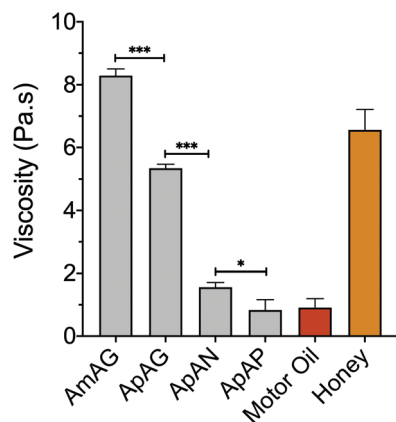


Fig. 3 Viscosity measurements made with a parallel plate rheometer at a temperature of 45 °C and a constant shear rate of 0.1 s⁻¹. ***, *p*-value < 0.0001, *, *p*-value < 0.05.

hold items. The viscosities of the SD-PUMs with rigid aromatic guaiacol end groups are comparable to honey, with AmAG and ApAG having viscosities of 8.29 and 5.35 Pa.s, respectively (Fig. 3). The viscosities of SD-PUMs with aliphatic end groups are comparable to motor oil, with ApAN and ApAP macromers having viscosities of 1.56 and 0.84 Pa.s, respectively. The viscosity clearly decreases with increasing end-group rotational flexibility, *i.e.*, going from a rigid guaiacol cap in AmAG and ApAG to aliphatic end caps in ApAN and ApAP. Flexible chains in the molecular structure increase the available free volume in the fluid, which results in enhanced mobility and diminished resistance to flow. ApAN, due to its bulky and rigid neopentyl end-group, had a higher viscosity (almost 2-fold) than ApAP as expected. Interestingly, the subtle change in the backbone structure from a 1,3 substitution in AmAG to a 1,4 substitution in ApAG led to a ~40% decrease in the viscosity (Fig. 3).

Preparation and characterization of SD-PUM networks

To understand the effect of composition and regioisomerism on the thermal and mechanical properties of thermosetting networks, the allyl groups in the SD-PUMs were cross-linked with a tetravalent and trivalent thiol *via* thiol-ene reactions. The two cross-linkers were selected to study the effect of cross-linker valency on material properties. The size of the cross-linkers were kept small, such that their chain flexibility would not mask the effect of SD-PUM structure. Networks were prepared *via* a thiol-ene photopolymerization reaction between the allyl groups of the SD-PUM and the thiol groups on the cross-linker. The four SD-PUMs and two cross-linkers were used to prepare eight unique thioether linked networks.

The networks were prepared by solvent casting a mixture of the SD-PUM, multivalent cross-linker, and a photoinitiator in acetonitrile into a rectangular PTFE mold (25 mm × 20 mm × 1 mm deep). Cross-linking was initiated with UV light (365 nm, 20 mW cm⁻²), and the networks were cured for 60 minutes. Following irradiation and gelation, each film was removed from the mold and swelled in acetonitrile for 48 h to remove unreacted macromers, cross-linker, and photoinitiator from the solid network. The residual acetonitrile solvent was removed, and the film was cut into ~5 mm wide rectangular strips, as seen in Fig. 4B, and dried in a vacuum oven for 12 h at 100 °C.

To examine the extent of cross-linking across all networks, the gel fractions of the networks were determined gravimetrically by subtracting the mass of the residual mixture from the total mass of the starting materials (not including the initiator). The gel fraction results in Fig. 4C indicate that the extent of cross-linking is similar (>90%) for all eight networks. Using the fluorescence signal of the SD-PUMs (ESI Fig. 13†), fluorescence measurements were acquired to measure the amount of residual SD-PUM in the recovered mixture by generating calibration curves for each macromer. The residual

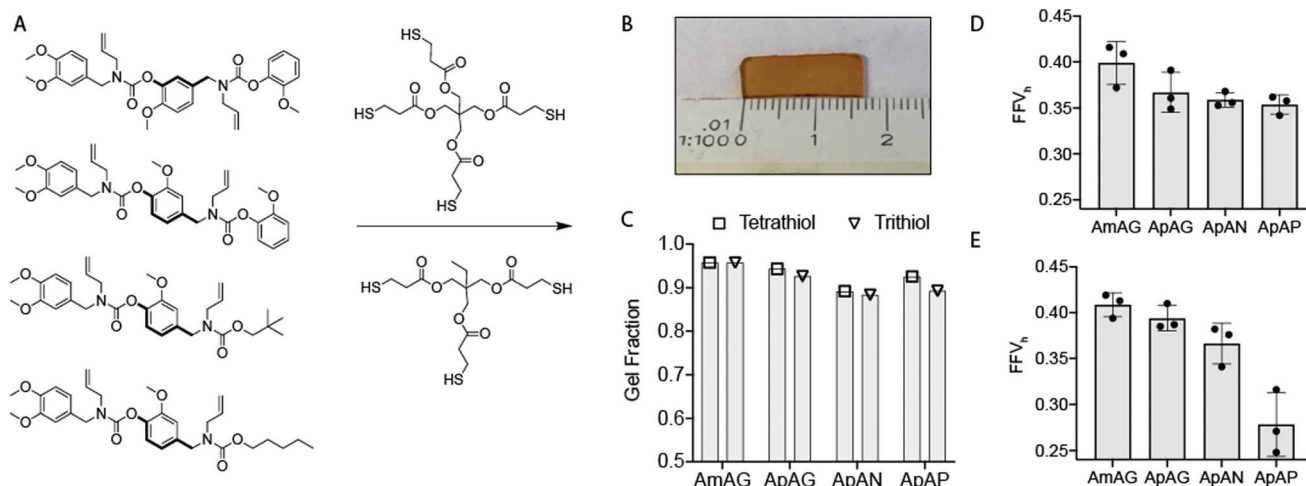


Fig. 4 (A) Macromer cross-linking *via* a photo-initiated thiol-ene reaction with a tetrathiol and trithiol. (B) Image of crosslinked film (C) Gel fraction for each of the crosslinked networks after 60 minutes of irradiation. Fractional free volume of static holes (FFV_n) of macromers crosslinked with the (D) tetrathiol, and (E) trithiol cross-linker.

mixture gave no fluorescence emission, which suggests that all macromers were incorporated in their respective networks.

We also examined the initial rate of cross-linking across all networks using a model kinetics experiment with a monothiol such that –ene conversion could be measured *via* ^1H NMR. Thiol–ene reaction kinetics were monitored by reacting the SD-PUMs with 3-mercaptopropionic acid (3-MPA) under the same reaction conditions as the network cross-linking reaction. The reaction conversion was observed at various time points by monitoring the disappearance of the allyl olefin peaks at 5.9 and 5.2 ppm *via* ^1H NMR spectroscopy (ESI Fig. 14†), and the conversion was plotted as a function of time (ESI Fig. 15†). The kinetics data shows similar reaction kinetics across all SD-PUMs. This, along with the gel fraction data in Fig. 4C, indicates that the rates and extent of cross-linking are not affected by the SD-PUM structure. It follows then that any differences in the network properties can be attributed to monomer structure and composition.

The fractional free volume,^{25–30} an empirical parameter that characterizes static hole volume in polymers in the glassy state, was measured for all eight cross-linked networks at -10°C . This static hole volume is the unoccupied space, in excess of the occupied volume (V_0), that is created due to the orientation and arrangement of the molecules in the frozen network. Beyond T_g , molecular vibrations are large enough to create enough free volume, in excess of the static hole volume and V_0 , to allow for large-scale segmental motions of polymer chains, which allows the material to flow. In the glassy state, higher static hole volumes are associated with polymer networks that are unable to pack efficiently due to rigid or bulky groups with constrained centers or shapes. Here, the fractional free volume of static holes (FFV_h), is the ratio of the estimated specific static hole volume (SSHV), also commonly referred to as the specific free volume (SFV), and the experimentally measured specific volume (V) (eqn (3)).

$$\text{SSHV} = V - V_0 \quad (1)$$

$$V_0 = 1.3 \times V_{\text{vdw}} \quad (2)$$

$$\text{FFV}_h = \text{SSHV}/V = 1 - (1.3 \times V_{\text{vdw}})/V \quad (3)$$

The SSHV was calculated using the Bondi group contribution method²⁷ in eqn (1) and (2), where V_0 is the occupied volume of the constituent moieties that comprise the network and V_{vdw} is the van der Waals volume, *i.e.*, the volume occupied by the molecule that is impenetrable by other molecules. The coefficient of 1.3 represents the average packing density of polymers at 0 K and is an approximation adopted from Van Krevelen²⁶ and Privalko³¹ due to the lack of data for these exact networks. The experimentally measured specific volume is the inverse of the density, which was obtained by measuring the mass and volume of the rectangular strips of films (3 replicates measured at -10°C). The van der Waals volume of the network was estimated numerically using values from Park *et al.*³⁰ as well as MarvinSketch (ESI Fig. 16 and 17†). The two estimates were similar, and thus volume estimates using

MarvinSketch were used to calculate FFV_h (Fig. 4D and E). One limitation of the FFV_h parameter as calculated is that the packing density embedded within it does not take into account different spatial arrangements due to the regioisomeric SD-PUMs. The positron annihilation lifetime spectroscopy measurement (PALS)^{32–34} is a unique probe for characterizing the local hole volume in networks and could also be used to measure FFV_h, though it has been reported that densely cross-linked networks such as epoxy networks tend to have small hole volumes that push the limit of the PALS method.³⁵

The data Fig. 4D and E show the AmAG SD-PUM with the highest FFV_h in both tetrathiol and trithiol cross-linked networks. In the tetrathiol networks, ApAG, ApAN, and ApAP gave similar FFV_h values while the trithiol network showed ApAP with a significantly lower FFV_h than its regioisomer, ApAN. As indicated earlier, high FFV_h values are associated with polymer networks that are unable to pack efficiently due to rigid or bulky groups with constrained centers or shapes. This suggests that materials with high FFV_h values have a high barrier to bond rotation, which is required for polymer unwinding and segmental motion of chains. Thus, our results suggest that the AmAG network is geometrically constrained in the glassy state relative to the other SD-PUM networks, and this should translate to higher T_g . Furthermore, our results imply that the ApAP trithiol linked network, given its much lower FFV_h, should have a low T_g relative to the other SD-PUM networks.

Effect of end-group composition and regioisomerism on thermal and mechanical properties

The T_g of the cross-linked networks were evaluated by differential scanning calorimetry (DSC) (Fig. 5A and B). Overall, the data showed higher T_g 's for networks made with the tetrathiol cross-linker (Fig. 5A) relative to those made with the trithiol cross-linker (Fig. 5B). We also see a general trend of decreasing T_g with increasing end-group flexibility as we go from AmAG and ApAG with rigid aromatic end-groups to ApAN and ApAP with flexible aliphatic end-groups. A predictable regioisomeric



Fig. 5 Differential scanning calorimetry of the (A) tetrathiol and, (B) trithiol cross-linked networks.



Fig. 6 Thermal and mechanical properties of sequence defined networks (60 min. UV-cure). (A–D) Dynamic mechanical analysis of cross-linked networks. The $\tan \delta$ curves of networks cross-linked with (A) tetrathiol and (B) trithiol. Plots of storage modulus (log scale) as a function of temperature for (C) tetrathiol cross-linked networks and (D) trithiol cross-linked networks.

effect was observed; ApAN networks with the bulky neopentyl group had a higher T_g relative to the ApAP network composed of a linear and flexible pentyl end-group. This observation is consistent with the conclusions from the FFV_h data. Furthermore, the data showed that the AmAG networks, made with both tetrathiol and trithiol, had the highest T_g values (Fig. 5A and B). Again, this observation is consistent with the conclusions from the FFV_h data. In addition to DSC measurements, T_g was also determined from DMA studies *via* $\tan \delta$ plots (Fig. 6). Similar trends seen in the DSC data were also recorded in the $\tan \delta$ plots. For the tetrathiol cross-linked networks, the T_g 's for AmAG, ApAG, ApAN, and ApAP were 35.4 °C, 24.1 °C, 22.6 °C, and 6.1 °C, respectively. Similarly, the T_g 's for the trithiol cross-linked networks were 22.1 °C, 21.3 °C, 11.2 °C, and –4.1 °C in the same order. Again, T_g decreases with increasing end-group flexibility. The difference in the T_g of the networks made with AmAG relative to ApAG could be due to the higher molecular rigidity or constrained shape of the AmAG macromer within the network. This effect is more pronounced in the tetrathiol cross-linked networks than in the trithiol networks. We posit that the additional connectivity in the tetrathiol amplifies the geometric constraints of the macromers in the networks.

The storage modulus, determined *via* DMA measurements as a function of temperature, is shown in Fig. 6C and D. Above T_g , all networks had similar storage moduli values ranging between 2–4 MPa in the rubbery regime. The rubbery modulus for the ApAG tetrathiol cross-linked network was ~45% lower than that of its backbone regioisomer, AmAG. However, the rubbery modulus of the ApAN network was ~40% lower than its end group regioisomer, ApAP. The latter is observed with both the tetrathiol and trithiol cross-linked networks (Fig. 6C and D). At room temperature, the storage modulus largely follows similar trends to the T_g and viscosity measurements.

The FFV_h values of the networks are correlated with the measured T_g . Networks with higher T_g values comprise chains that are composed of structurally “constrained” moieties, as observed in the viscosity data (Fig. 3). This molecular constraint results in the accumulation of larger static hole volumes in the glassy state, hence high FFV_h. Polymeric networks that are unable to pack efficiently due to these molecular constraints, for the same reason, have a high barrier to

bond rotation, which is required for polymer unwinding and segmental motion of chains. Thus, networks with poor packing efficiencies due to geometric constraints have a high FFV_h and high T_g relative to others. Overall, our results, when paired with the gel fraction and kinetics data, support the conclusion that the observed differences in thermomechanical properties are related to the impact of SD-PUM structure (*i.e.* regioisomerism).

Conclusions

This study sought to investigate the effect of regioisomerism and composition on material properties *via* a recently developed scalable SD-PUM platform. This was achieved by designing two sets of regioisomeric SD-PUMs with different backbone and end-groups. Analysis of the synthesized SD-PUMs *via* NMR and viscosity measurements showed changes in physical properties owing to their structural differences. Following cross-linking with multivalent thiols, the thermomechanical properties of the networks were obtained experimentally. Here, we observed that the change in backbone connectivity influenced the T_g to a greater extent when cross-linking was performed with a tetrathiol relative to a trithiol. The difference in the thermomechanical properties was determined to be solely due to composition and structure by ruling out other factors such as the extent and rate of cross-linking. With the use of regioisomers, this study clearly illustrates how subtle changes in backbone and end-group structure significantly impacts network performance. We discovered that a subtle change from 1,3 to 1,4 connectivity leads to a decrease in viscosity as well as attenuated thermomechanical properties of the ensuing networks. This response in material properties to the subtle change in backbone connectivity could be used to alter the properties of commercial resins, *e.g.*, tuning the mechanical, self-healing, and adhesive properties of phenylene diamine-based epoxy resins. Armed with the versatility of the SD-PUM platform, the next step beyond backbone conformation and terminal group composition is to combine these elements along with sequence to generate multifunctional macromers with multiple handles that can be used to further tune material properties.

Conflicts of interest

There are no conflicts to declare.

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