

Molecular dynamics simulation on glass transition temperature of isomeric polyimide

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Abstract. It is an abnormal phenomenon that glass transition temperature (T_g) of isomeric polyimide (PI) is higher than its corresponding symmetrical PI. To illustrate this phenomenon at the molecular scale, we applied molecular dynamics method to predict the T_g of PI, which were prepared based on 4,4'-oxydianiline (ODA) and 3,3',4,4'-biphenyltetracarboxylic dianhydride (3,3',4,4'-BPDA), and its isomeric system (2,2',3,3'-BPDA-ODA). Simulation result is consistent with experimental value. Non-bond energy plays an important role in glass transition process, for it has an abrupt change near T_g . The higher free volume fraction of isomeric PI can provide the polymer with more space to obtain segmental motion. However, from the torsion angle distribution calculations, it is shown that the torsion angle of its biphenyl group is constrained. Furthermore, from mean square displacement and vector autocorrelation functions calculations, this group is observed to rotate against other groups in the glassy state, and increases the chain rigidity to a great extent. So the isomeric PI needs much more relaxation time for the segment motion. Therefore, the higher T_g of isomeric PI is mainly attributed to the chain rigidity for the time scale, not the free volume for the space scale.

Keywords: modeling and simulation, isomeric polyimide, molecular dynamics simulation, glass transition temperature

1. Introduction

Glass transition temperature (T_g) is a key index to evaluate the thermal properties of heat-resistant polymers, such as polyimide (PI), polysulfone, polyether ether ketone. As for polyimide, isomeric PI, which incorporate geometrically asymmetrical unit in dianhydride or diamine structures, can improve the processability and solubility; at the same time, its thermal properties do not show obvious decrease for there is no composition change compared with corresponding symmetrical PI. Unexpectedly, the T_g of isomeric PI is relatively higher than that of symmetrical PI [1]. This interesting phenomenon is not observed with other polymers, such as polyamide [2] and poly(arylene ether ketone) [3, 4]. Several researchers have tried

to explain this phenomenon: Tong et al. [5] reported that the T_g of symmetrical PI, which was polymerized from 3,3',4,4'-biphenyltetracarboxylic dianhydride (3,3',4,4'-BPDA) and 4,4'oxydianiline (ODA), was 52 K lower than that of its isomeric system (2,2',3,3'-BPDA-ODA). Furthermore, they analyzed the relative energies of torsion angle for PI monomer, and suggested that the steric hindrance of the adjacent anhydride groups of 2,2',3,3'-BPDA inhibit the segmental motion, and increase the T_g , relative to that of the 3,3',4,4'-BPDA. Kochi et al. [6] reported similar results on these two systems, and analyzed the temperature of β relaxation using the dynamic mechanical analysis (DMA) method. Hasegawa et al. [7] studied on the isomeric phenomena of PIs, which were prepared

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based on BPDA and 1,4-phenylene diamine (PDA). Semiempirical method was used to study the conformation of a single PI chain in gas state. They inferred that the higher T_g for the isomeric polyimide is attributed to the asymmetric structure requiring a larger sweep volume for the crank shaft motion and accordingly more energy (higher temperature) to obtain this motion. The above studies facilitate us with the insight into the nature of the higher T_g of isomeric PI, however, there has been no systematic study investigating the effect of isomerism on T_g of polyimides. It is acknowledged that the T_g of polymeric materials is mainly determined by the segment motion of a polymer chain. If this motion can be observed, the polymer chain should have sufficient space (more free volume) and can complete this motion in a short time period (less relaxation time, it mainly depends on rigidity of a polymer chain). Nevertheless, these two factors can not be well discussed in the gas state [5, 7], so it is necessary to find an appropriate method to investigate them in the bulk state.

Yet, molecular dynamics (MD) simulation method provides a proper means to study bulk polymer at molecular scale. Some researchers have applied it to study the glass transition temperature of polymers, such as polyethylene [8], polypropylene [9], polystyrene [9], poly(oxymethylene) [8], polysiloxane [10], polylactic acid [11], epoxy resin [12]. The results indicate a good agreement with experimental value. In the filed of polyimides, Liang et al. [13] successfully predict the glass transition temperature of PI with different chromophores using MD method, and they deduced that T_g of PIs are correlated to the height of energy barrier of bridging bonds between aromatic rings. Recently, Yani and Lamm [14] studied on the glass transition temperature of PI/polyhedral oligomeric silsesquioxane (POSS) blends, showing that the simulation results are comparable to experimental observation and incorporation of the appropriate content POSS can improve glass transition temperature. Thus, MD method is an effective method to predict the glass transition temperature of polymers at the molecular scale.

In this study, we firstly applied MD simulation method to study the effect of isomerism on the glass transition temperature of bulk polyimides, using 3,3',4,4'-BPDA-ODA and 2,2',3,3'-BPDA-ODA systems (Figure 1) as model molecules. The

Figure 1. Structures of two polyimides: 3,3',4,4'-BPDA-ODA and 2,2',3,3'-BPDA-ODA

rest of the paper is organized as follows: In section 2, simulation details of MD method are described. Section 3 is result and discussion. We obtained comparative value of T_g in PIs, and analyze the role of energy component in glass transition process. Furthermore, the reason for higher T_g of 2,2',3,3'-BPDA-ODA systems is given by the analysis at time scale (chain rigidity) and space scale (free volume). Finally, we summarize our major results and conclusions in section 4.

2. Simulation details

Simulations were performed using the Materials Studio 4.0 software purchased from Accelrys Inc [15]. The Dreiding force field was used in molecular mechanics (MM) and molecular dynamics simulations [16]. The total potential energy (E_{total}) can be expressed as:

$$\begin{split} E_{total} &= E_{bond} + E_{non-bond} = \\ &E_b + E_{\theta} + E_{\phi} + E_{inv} + E_{vdW} + E_{elec} \end{split} \tag{1}$$

Equation (1) is divided into two parts: valence terms (E_{bond}) and non-bond interaction terms $(E_{non-bond})$. The valence terms include E_b , E_{θ} , E_{ϕ} , E_{inv} for bond stretching, angle bending, dihedral torsion and inversion (out-of-plane) terms. The non-bond interaction terms include E_{vdW} and E_{elec} for calculating van der Waals force and electrostatic interactions respectively. The non-bond interactions were calculated in this study with cut-off being set to 0.95 nm, which is less than half of the cell length (during the glass transition process, cell length was kept around 2.2-2.5 nm). The amorphous polymer models were built with the Theodorou and Suter's method [17, 18] using periodic boundary conditions. Each cell consisted of a parent PI chain with 20 repeat units. Though the system size of 20 repeat units is not enough to represent the conformation of a real polymer chain, previous researchers have reported reasonable result when they used 10–15 repeat units [13, 19]. In addition, previous studies demonstrate that simulated T_g values are comparable with experimental one by simulating a single polymer chain [8-10, 13]. So we can save a lot of computational resource in this way. The cell was first optimized in a low initial density of 0.5 g/cm³ by smart minimization, which incorporated steepest decent with conjugate gradient methods. Then, relaxation of the cell was performed by MD simulation at 1000 K for 2 ns using an integration step of 1 fs under constantpressure constant-temperature (NPT) ensemble. The conformation of low potential energy structure was selected and minimized again. In this procedure, density of each PI system was above 0.8 g/cm³. To make sure a system reached equilibrium, a number of thermodynamic quantities were checked. As a result, they remained stable during the last 500 ps, and it was concluded that the system was equilibrated.

The glass transition temperature of polymeric materials can be determined by plotting specific volume versus temperature with constant-NPT MD simulation and noting where the slope changes [14]. The system was cooled from 750 K stepwise to 400 K by decrements of 50 K. At each temperature, constant-NPT MD simulation was conducted for 1 ns. In the last 100 ps, the cell density was sampled at every picosecond, and the average value was taken as the result.

In addition to the glass transition temperatures, we have also calculated torsion angle distribution, mean square displacement and vector autocorrelation functions of polyimides. The equilibrated configuration in the constant-NPT MD simulation was used as the initial structure, and then was calculated by constant-volume constant-temperature (NVT) MD simulation for 2 ns. Data samples were collected at every picoseconds.

In this study, the pressure of system was set to $1.0 \cdot 10^5$ Pa. Nose thermostat [20, 21] and Berendsen

barostat [22] were applied to control temperature and pressure in constant-NPT MD simulation.

3. Results and discussion

3.1. Glass transition temperature

The temperature dependence of the specific volume of two PI systems is shown in Figure 2. With increasing temperature, the increment of specific volume is less pronounced below T_g yielding a kink in the curve to determine T_g . Detailed method has been well described in ref. [11]. According to this method, simulated glass transition temperatures of these two PIs were obtained. The T_g of isomeric PI (2,2',3,3'-BPDA-ODA) is 593 K which is much higher than symmetrical PI's (3,3',4,4'-BPDA-ODA, 548 K). Simulation result is consistent with experimental values shown in Table 1. For comparison, we also applied group contribution method to predict T_g of PIs [13]. This method is based on the contribution of each functional group to thermodynamic properties of polymeric materials, and it has been widely used in alkyl polymers [23, 24]. However, in this study, the T_g values of two PIs are the same using this method for it can not resolve the position effect of the imide group. Therefore, MD simulation is a relatively effective method to predict T_g of isomeric polyimides.

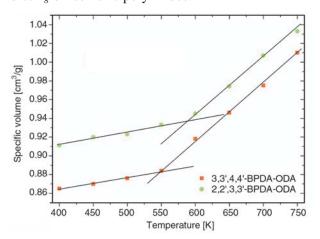


Figure 2. Plots of specific volume of MD simulation versus temperature for two polyimides: 3,3',4,4'-BPDA-ODA (■); 2,2',3,3'-BPDA-ODA (●)

 Table 1. Glass transition temperatures of PIs with MD simulation, correlation prediction, and experiment methods

PI systems	Glass transition temperature [K]		
	MD simulation	Correlation predictiona	Experiment ^b
3,3',4,4'-BPDA-ODA	548	576	539
2,2',3,3'-BPDA-ODA	593	576	591

^aPredicted by Synthia module in Materials Studio software package

^bExperimental value in ref. [5]

Also, we checked the density of PIs at 298 K. The density for 3,3',4,4'-BPDA-ODA system is 1.20 g/cm^3 while the density of 2,2',3,3'-BPDA-ODA system is 1.13 g/cm^3 . And the density of 3,3',4,4'-BPDA-ODA system is about 17% smaller than experiment value [25] (density of 2,2',3,3'-BPDA-ODA system has not been reported in existing reference). However, when we want to get the comparable T_g from the MD simulation, the dynamic change of density (or specific volume) near T_g should be emphasized, not the density itself.

3.2. Free volume

According to the Fox and Flory's theory of glass transition, the free volume of the polymer experiences an abrupt change during the cooling down at a point called the glass-transition temperature. In this study, we applied a probe sphere, whose radius is 0.1 nm (distance between neighboring atoms) or 0.3 nm (distance between neighboring molecules), to probe the equilibrated polymer cells to calculate the free volume [8]. Furthermore, the free volume fraction (FVF) can be obtained by Equation (2):

$$FVF = \frac{V_f}{V_f + V_0} \tag{2}$$

where V_f is free volume, and V_0 is occupied volume of the polymer chains.

The plot of FVF of two PIs with a probe sphere of 0.1 nm radius versus temperature is shown in Figure 3, indicating similar T_g with the plot of specific

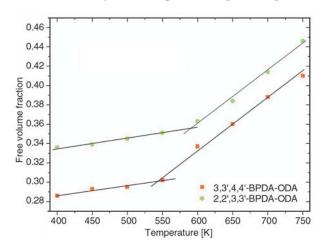


Figure 3. Plots of free volume fraction versus temperature for two polyimides: 3,3',4,4'-BPDA-ODA (■); 2,2',3,3'-BPDA-ODA (●)

volume versus temperature. However, increasing the radius to 0.3 nm, we can not get the clear T_g by the same method, for it is difficult to determine van der Waals surface with the probe sphere of big radius in this measurement. During the glass transition process, free volume determines how much space can be provided for the segment motion of a polymer chain. For polymers with similar compositions, the lower FVF can bring with higher T_g of that polymeric materials, for it can not provide sufficient space for segment motion at the same temperature. It is clear that free volume fraction of 2,2',3,3'-BPDA-ODA system is higher than 3,3',4,4'-BPDA-ODA system at the same temperature, as the same result for specific volume showing in Figure 2. In 2,2',3,3'-BPDA-ODA system, incorporation of asymmetry group decreases the regularity of polymer chain, inhibits the close packing of inter polymer chain, and increases free volume at molecular scale, namely facilitates the segment motion to some extent. This result may go against the result of higher T_g of isomeric PIs. However, free volume is not the single variable to evaluate the T_g of PIs, other factors that may affect the T_g will be discussed in the following sections.

3.3. Role of the energy components in glass transition process

To understand the role of each energy component playing in glass transition process, we examined the plot of the energy component versus temperature for two PI systems, and the result shows that the PIs show the same trend in the plot. Take isomeric 2,2',3,3'-BPDA-ODA system for example, the simulated results of bond energy, angle energy, torsion energy, inversion energy and non-bond energy versus the temperature were plotted in Figure 4. There is a kink only in the plot of non-bond energy versus temperature, indicating the occurrence of glass transition. At both below and above T_g , non-bond energy increases almost linearly with increasing temperature with a break at T_g . But for the other energy components, they consistently increase linearly with increasing temperature in the whole temperature range. These results show that the non-bond energy plays an important role in the glass transition process of the polymer system. The above results are consistent with the same analysis on epoxy resin by Wu and Xu [12]. However, Li

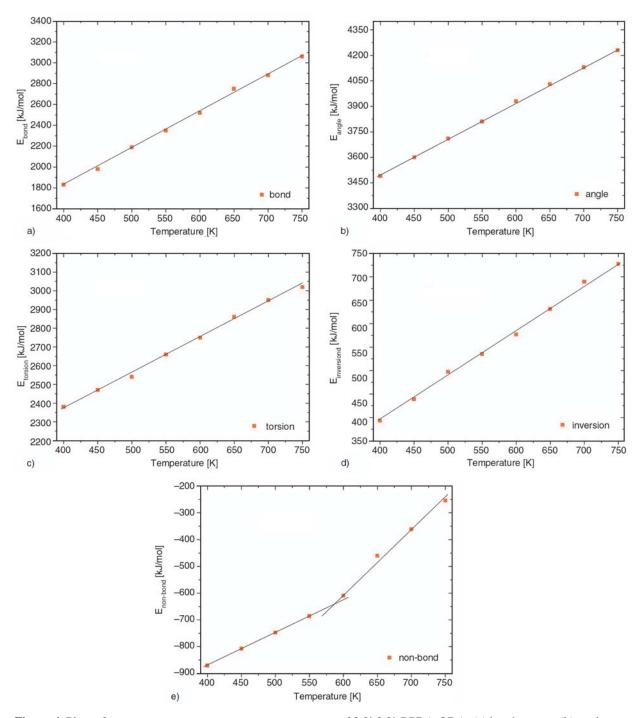


Figure 4. Plots of some energy components versus temperature of 2,2',3,3'-BPDA-ODA: (a) bond energy; (b) angle energy; (c) torsion energy; (d) inversion energy; (e) non-bond energy

and coworkers [8, 26] also performed the same analysis on polyoxymethylene (POM) system [8] and poly(3-hydroxybutyrate) (PHB)/poly(ethylene oxide) (PEO) blend system [26] and drew a different conclusion from ours: both torsion and nonbond energy played important roles in glass transition. In fact, the freedom of bond stretching and angle bending could be equilibrated fast in the simulation, therefore the linear increasing trend are

shown. Unlike the common alkyl polymers, aromatic PI has much higher T_g for it has plenty of phenyl and imide rings, so the dihedral torsion and out-of-plane motion of aromatic PIs are constrained, and no kink has been observed in this study. However, the segment motion ability of aromatic PIs are mainly attributed to the limited torsion of linking of phenyl and imide rings [13]. Influenced by the thermal fluctuation, limited link-

ing groups show distinct mobility, yielding the conformation transition, and furthermore, resulting in the sudden change of non-bond energy near T_g .

3.4. Torsion angle distribution

As discussed in section 3.3, mobility of linking groups exerts an important influence during the glass transition process. In order to elucidate the reason for the higher T_g of isomeric PI, torsion angle distributions of linking in two PIs were plot-

ted (Figure 5 and Figure 6). Torsion angles Φ 1, Φ 2 and Φ 3 were defined in Figure 1. For facilitating the discussion, we defined C_p for carbon atom in phenyl ring and C_i for carbon atoms in imide ring linking the ether group. Therefore, we can define $C_p-C_p-C_p-C_i$ (Φ 4) as the torsion angle of C_p-C_p between phenyl and imide rings, and $H-C_p-C_p-C_p$ (Φ 5) as torsion angle of C_p-C_p in the phenyl ring which is between imide ring and ether group. Furthermore, the Φ 4 and Φ 5 can be used for study of the co-plane properties of these aromatic rings. As

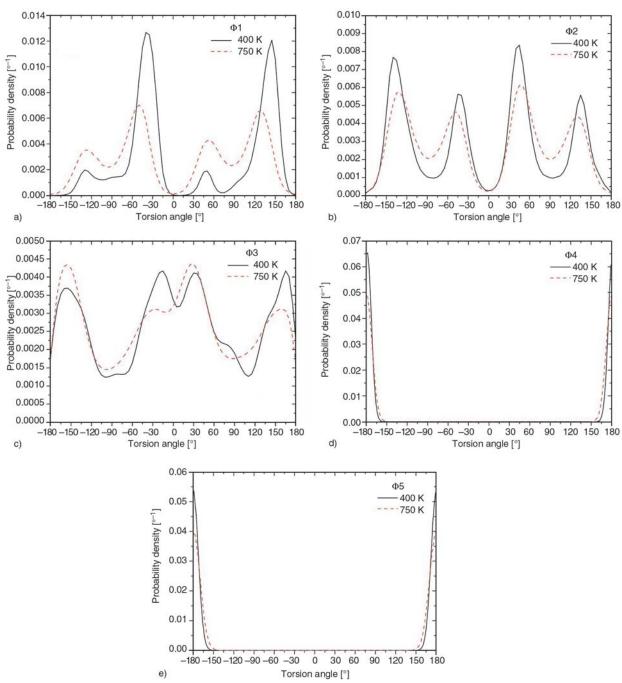


Figure 5. Torsion angle distributions of 3,3',4,4'-BPDA-ODA: (a) Φ 1, (b) Φ 2, (c) Φ 3, (d) Φ 4 and (e) Φ 5. The results have been obtained with constant-NVT MD simulation at T = 400 K (——) and 750 K (——).

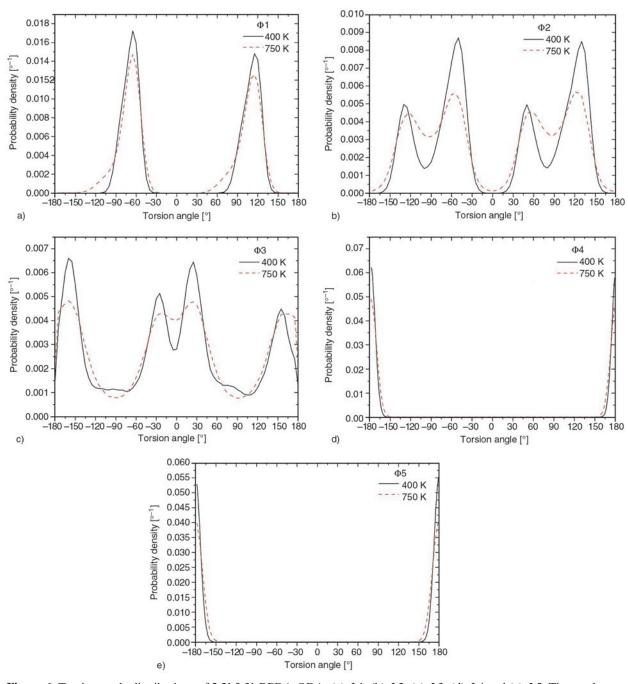


Figure 6. Torsion angle distributions of 2,2',3,3'-BPDA-ODA: (a) Φ 1, (b) Φ 2, (c) Φ 3, (d) Φ 4 and (e) Φ 5. The results have been obtained with constant-NVT MD simulation at T = 400 K (——) and 750 K (——).

for $\Phi 1$ (torsion angle of C_p – C_p between phenyl ring), 3,3',4,4'-BPDA-ODA system has peaks in $\pm 35^{\circ}$ and $\pm 135^{\circ}$, but in isomeric 2,2',3,3'-BPDA-ODA system, there are only two peaks located in the $\pm 60^{\circ}$. This result is similar to the plot of relative energy of torsion angles for PI monomers in the same systems obtained by molecular mechanics in the previous study [5]. Higher rotational barrier corresponding to $\Phi 1$ in 2,2',3,3'-BPDA-ODA system is responsible for the two symmetrical two peaks in the torsion angle distribution, and eventu-

ally results in its higher T_g . Two PIs have the same peak position but different relative proportion for $\Phi 2$ (torsion angle of N–C_p between imide and phenyl rings) and $\Phi 3$ (torsion angle of C_p–O in diamine structure). As for $\Phi 4$ and $\Phi 5$, two PIs have only two half peaks in $\pm 180^\circ$, illustrating that phenyl and imide ring can keep in a plane well under the premise of phenyl ring's keeping in a plane. To explore the effect of temperature on torsion angle distribution, two PIs' torsion angle distributions in 400 and 750 K were given. These

distribution functions exhibit peak at the same position at two different temperatures, only a wide distribution at the higher temperature, especially $\Phi 2$ and $\Phi 3$. No matter at glassy or rubbery state, the torsion angles of $\Phi 4$ and $\Phi 5$ are kept around $\pm 180^\circ$. It proves that atoms in phenyl and imide ring of PIs can be kept in a plane respectively during the glass transition process. Furthermore, it also confirms that the torsion energy does not have sudden change in the glass transition process due to confined torsion, which has been discussed in section 3.3.

3.5. Mean square displacement

The value of T_g is correlated well with the polymer rigidity, namely the segmental mobility in polymer chain. To observe the mobility of PI chains during glass transition process, the mean square displacements (MSD) were calculated [14]. With the increasing temperature, the slope of MSD of both two PI systems becomes steeper. Take 2,2',3,3'-BPDA-ODA system as an example (Figure 7), the slope of MSD above T_g (700 and 750 K) are much higher than that below T_g (400 and 500 K), indicating higher mobility of PI chains above T_g . Furthermore, we have also compared the mobility of 3,3',4,4'-BPDA-ODA and 2,2',3,3'-BPDA-ODA systems. Figure 8 shows the mean square displacement plot for the two PI systems at 400 and 750 K. It is shown that the MSD for 2,2',3,3'-BPDA-ODA system is lower than that for 3,3',4,4'-BPDA-ODA system at all times. This means 2,2',3,3'-BPDA-ODA system has lower mobility than 3,3',4,4'-BPDA-ODA system, and hence it needs more time

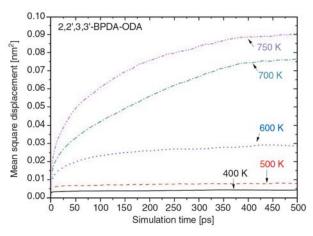


Figure 7. Mean square displacements of 2,2',3,3'-BPDA-ODA system as a function of temperature

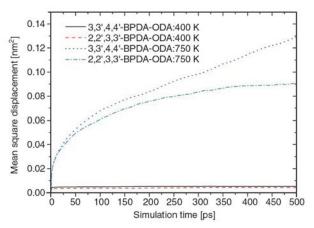


Figure 8. Mean square displacements of 3,3',4,4'-BPDA-ODA and 2,2',3,3'-BPDA-ODA systems at T = 400 K and 750 K

(or higher temperature) to obtain segment motion. The above observations support the result of higher T_g of 2,2',3,3'-BPDA-ODA systems.

3.6. Vector autocorrelation function

To further characterize the mobility of PI chains, the mobility of linking groups were examined. It can be quantified by vector autocorrelation functions (VACFs) of these groups along the polymer chain (Equation (3)):

$$M_2(t) = \frac{3 < (u(t) \bullet u(0))^2 > -1}{2} \tag{3}$$

where u(t) and u(0) represent unit vectors of the segment at time t and time 0, respectively. The <> indicates an ensemble average over such vectors [12]. At the same simulation time, the higher value of $M_2(t)$ indicates the higher rigidity of the segment

The segmental vectors corresponding to the torsion angle Φ 1, Φ 2, Φ 3, Φ 4 and Φ 5 are shown in Table 2. They are defined as the vectors from the first atom to the fourth atom of the torsion angles. The plots of VACFs of these segments in two PIs are given at two temperatures: 400 K (Figure 9) and 750 K (Figure 10) [12]. At 400 K, the value

Table 2. Definition of segmental vectors corresponding to torsion angle Φ 1, Φ 2, Φ 3, Φ 4 and Φ 5

Torsion angle	Atoms	Segmental vector
Ф1	C_{p1} - C_{p2} - C_{p3} - C_{p4}	$C_{p1} \rightarrow C_{p4}$
Ф2	C_{i} - N - C_{p1} - C_{p2}	$C_i \rightarrow C_{p2}$
Ф3	C_{p1} - C_{p2} - O - C_{p3}	$C_{p1} \rightarrow C_{p3}$
Ф4	C_{p1} – C_{p2} – C_{p3} – C_{i}	$C_{p1} \rightarrow C_i$
Ф5	$H-C_{p1}-C_{p2}-C_{p3}$	$H \rightarrow C_{p3}$

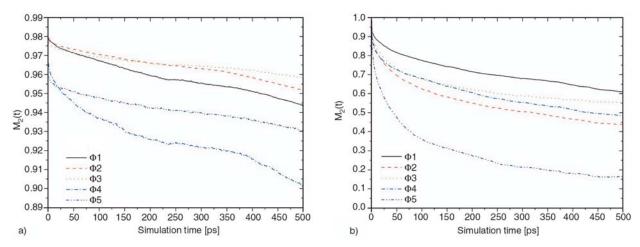


Figure 9. Vector autocorrelation functions of vector bonds corresponding to Φ 1, Φ 2, Φ 3, Φ 4 and Φ 5 in 3,3',4,4'-BPDA-ODA with constant-NVT MD simulation at (a) T = 400 K and (b) 750 K

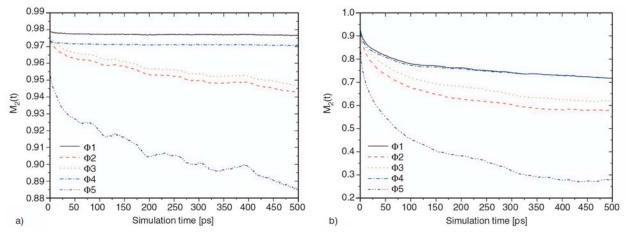


Figure 10. Vector autocorrelation functions of vector bonds corresponding to Φ 1, Φ 2, Φ 3, Φ 4 and Φ 5 in 2,2',3,3'-BPDA-ODA with constant-NVT MD simulation at (a) T = 400 K and (b) 750 K

 $M_2(t)$ in two PIs are above 0.9 during the simulation time, indicating high chain rigidity of PIs. In 3,3',4,4'-BPDA-ODA system, the mobility of Φ 1 is higher than Φ 2 and Φ 3, but the opposite result is obtained in 2,2',3,3'-BPDA-ODA system. The reason is that higher steric hindrance between biphenyl group can lead to lower torsion mobility of 2,2',3,3'-BPDA-ODA system. Because the aromatic rings can keep in a plane respectively during the simulation, we can define $C_{p1} \rightarrow C_i (\Phi 4)$ to characterize the torsion mobility of phenyl and neighboring imide rings. Also, we can define $H\rightarrow C_{p3}$ $(\Phi 5)$ to characterize that of phenyl ring in diamine structure. In this way, the cooperative motion among Φ 1, Φ 2 and Φ 3 can be characterized through Φ 4 and Φ 5. In 3,3',4,4'-BPDA-ODA system, the mobility of $\Phi 4$ is much higher than $\Phi 1$ and $\Phi 2$, for it exists a positive cooperative motion between Φ 1 and Φ 2. Yet, in 2,2',3,3'-BPDA-ODA system, mobility of $\Phi 4$ lies between $\Phi 1$ and $\Phi 2$, which

indicates that Φ 1 has reverse torsion motion with Φ 2, thereby it weaken the mobility of Φ 4. As for Φ 5, its mobility is much higher than Φ 2 and Φ 3 in two PI systems, showing the positive cooperative motion between $\Phi 2$ and $\Phi 3$. Possible torsion model is shown in Figure 1, and the result confirms the inference of torsion analysis via DMA method in the previous study [6]. In two PI systems, the mobility of five torsion angles shows an obvious increment at 750 K, and the same arrangement order in mobility, that is $\Phi 5 > \Phi 2 > \Phi 3 \approx \Phi 4 > \Phi 1$. This result indicates that the $\Phi 2$ and $\Phi 3$ are easier than Φ 1 to be influenced by thermal fluctuation. The mobility of $\Phi 2$ and $\Phi 3$ increase more quickly at higher temperature, and then the conformation transitions are easier to take place. So the corresponding torsion angles are distributed wider (see Figure 5 and Figure 6). The mobility of $\Phi 1$ is the lowest among the five torsion angles, and hence it can be used to compare the mobility of the polymer chains in two PI systems. At 750 K, the value of $M_2(t)$ in 2,2',3,3'-BPDA-ODA system is higher than that of 3,3',4,4'-BPDA-ODA system. This result indicates that 2,2',3,3'-BPDA-ODA system has lower mobility than 3,3',4,4'-BPDA-ODA system. And it also confirms the claim that the higher T_g in isomeric 2,2',3,3'-BPDA-ODA system is mainly attributed to the limited torsion motion between biphenyl groups. Yet, in two PI systems, the mobility of Φ 4 lays between Φ 1 and Φ 2, while that of $\Phi 5$ is much higher than $\Phi 2$ and $\Phi 3$. In other words, the positive comparative motion of $\Phi 1$ and Φ2 in 3,3',4,4'-BPDA-ODA system disappears in the rubbery state. It may be resulted from partial reverse motion of diamine structure above glass transition temperature.

4. Conclusions

MD simulation was performed to study the glass transition temperature of isomeric PIs. Simulated T_g is consistent with experimental value: T_g of isomeric 2,2',3,3'-BPDA-ODA system is higher than 3,3',4,4'-BPDA-ODA system.

The glass transition temperature of polymeric materials is mainly determined by the segment motion of a polymer chain. It mainly involves the factors at space scale (free volume) and time scale (chain rigidity). Higher rigidity and less free volume lead to the higher T_g of polymers, however, these two factors are in opposition in isomeric PI (2,2',3,3'-BPDA-ODA). The higher free volume fraction of isomeric PI can provide the polymer with more space to obtain the segment motion. However, from the torsion angle distribution calculations, it is shown that the torsion angle of its biphenyl group is constrained. Furthermore, from mean square displacement and vector autocorrelation functions calculations, this group is observed to rotate against other groups in the glassy state, and increases the chain rigidity to a great extent. So the isomeric PI needs much more relaxation time for the segment motion. Therefore, the higher T_g of isomeric PI is mainly attributed to the chain rigidity for the time scale, not the free volume for the space scale.

Acknowledgements

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