

Dynamic hyperconjugation induced bond-angle rotation isomerization mechanism and anionic conformational rigidity-dependent melting points of ionic liquids

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Abstract

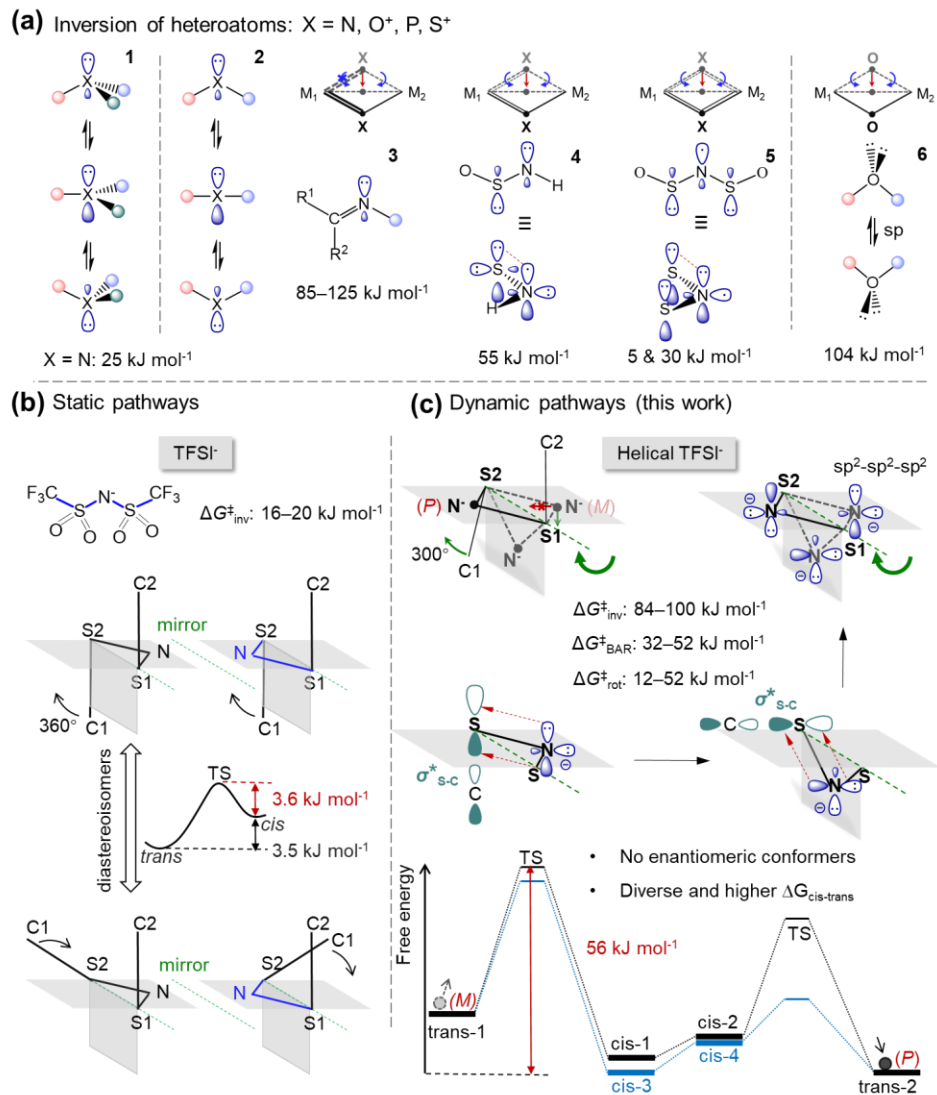
The pyramidal inversion of trisubstituted heteroatoms is well-recognized. However, the stereochemistry of bicoordinated heteroatoms is poorly developed and few examples assumedly isomerize by double-torsion motion or bond-angle inversion. Here, by using *ab initio* molecular dynamics combined with metadynamics simulation, we reveal an unexpected competing isomerization mechanism, namely dynamic $\pi \rightarrow \sigma^*$ negative hyperconjugation induced coupled rotation of bond and bond angle leads to helix inversion in bis(trifluoromethane)sulfonamide (TFSI⁻), making TFSI⁻ follow four distinct *trans-cis*

isomerization pathways with diverse energy barriers (12–52 kJ mol⁻¹) which are significantly higher than 3.6 kJ mol⁻¹ of sole pathway estimated by conventional static calculations. Our quantitative simulations and experiments results confirm the positive correlation between overall stability of *cis*-TFSI⁻ with polarity of counteranions. The melting points (T_m) of TFSI⁻-based ionic liquids (ILs) linearly rise with the conformational rigidity of TFSI⁻ in the ion pair state, offering a new fundamental perspective on the origin of low T_m of ILs.

Introduction

The pyramidal inversion **1**, a fundamental stereochemical process, is intensively studied and well-established in tricoordinated heteroatom compounds (X = N, O⁺, P, S⁺), which involves a planar transition state and thus sp³-sp²-sp³ re-hybridization at the heteroatoms X (Scheme 1a)¹⁻⁴. Many isolable phosphines¹ and sulfonium salts⁴ possess high inversion energy barriers of 120–185 kJ mol⁻¹ and 100–130 kJ mol⁻¹, whereas nitrogen pyramidal inversion is rapid and only takes 20–25 kJ mol⁻¹ in a tertiary amine due to quantum tunnelling⁵. By contrast, the bond-angle inversion (BAI) **2** at a singly bonded atom⁶, is much less exploited, though we have extended this isomerization mechanism to the bicoordinated heteroatoms in doubly bonded or conjugated systems. Nitrogen BAI (85–125 kJ mol⁻¹)^{7,8} and rotation around C=N bond (210–250 kJ mol⁻¹ barrier for photoinduced rotation^{9, 10}) provide alternative concerted mechanism for (E/Z) stereoisomerization in imine C(R₁R₂)=N(R₃) **3**. When nitrogen is merely conjugated with one of ligands, the BAI barrier drops to 55 kJ mol⁻¹ in thionylimide HNSO **4**¹¹. When nitrogen is conjugated with both ligands, the S-N-S inversion requires exceptionally low energy (5 & 30 kJ mol⁻¹) in

heterocumulene radical OSNSO **5**⁸, which is even lower than 40 kJ mol⁻¹ for two torsional motions. B-O-B inversion **6** overcomes an unusually high barrier of 104 kJ mol⁻¹, which is still lower than that of the double-torsion pathway involving only 90° motions, in singly bonded (BF)O(BF)-quinoxalinoporphyrin, owing to the geometrical restraints from B-O-B encapsulation⁶. Although all these examples involve a linear transition state, the oxygen BAI and nitrogen BAIs are distinct in the numbers of valence electron, bonding characters with ligands, and thus re-hybridization processes of central atoms (sp³-sp-sp³ (O) vs. sp²-sp-sp² (N)). To date, there are no progresses in identifying the isomerization mechanism for bicoordinated O or its valence isoelectronic atoms (e.g., S in disulfide^{12, 13}) or ions (e.g., N⁻) in the systems with conjugated M₁-X-M₂.



Scheme 1. Stereochemistry of heteroatoms. (a) Pyramidal inversion of tricoordinated heteroatoms **1**. Bond-angle inversion (BAI) (red arrows) of bicoordinated heteroatoms **2**, doubly bonded imine **3**, conjugated thionylimide HNSO **4** and heterocumulene radical OSNSO **5**, and singly bonded (BF)O(BF)-quinoxalinoporphyrin **6**. $\Delta G_{inv}^{\ddagger}$ for energy barrier of inversion, kJ mol^{-1} . Blue arrows for two torsional motions. (b) BAI and diastereomeric interconversion of TFSI⁻ anion with S–N–S moiety. (c) Dynamic $\pi \rightarrow \sigma^*$

negative hyperconjugation induced out-of-plane bond-angle rotation (BAR) leads to the helix inversion of TFSI⁻ conformers. $\Delta G_{\text{rot}}^{\ddagger}$ for lone S–C bond rotation.

Centered with N⁻, the anion bis(trifluoromethyl-sulfonyl)imide (TFSI⁻) is assumed to be conjugated among –SO₂–N⁻–SO₂–. N⁻ BAI only required 16–20 kJ mol⁻¹ (Scheme 1b), estimated from the reported potential energy surface (PES)¹⁴⁻¹⁷. The two pairs of enantiomers of TFSI⁻ (*trans* or *cis* isomers) with the only energy difference ($\Delta E_{\text{cis-trans}}$) of 3.5 kJ mol⁻¹, underwent diastereomeric interconversion over the sole torsional barrier ($\Delta E_{\text{cis-trans}}^{\ddagger}$) of 3.6 kJ mol⁻¹.¹⁴⁻¹⁸ However, the barrier for rotation around single bond is about 12 kJ mol⁻¹ in ethane¹⁹, but 40 kJ mol⁻¹ in radical OSNSO⁸ and 85 kJ mol⁻¹ in an amide¹⁹ because of the conjugation effect. Moreover, previous theoretical studies on TFSI⁻ and even all the BAI examples are limited to the static electronic structure calculation, a nonergodic, chemical-intuition-dependent approach, which is insufficient to rationalize or predict the isomerization/reaction mechanism involving unforeseen processes (e.g., coupled motions) while ignoring dynamic effects.²⁰⁻²² TFSI⁻-based ionic liquids (ILs) exhibit multiple exotic thermal and mechanical properties, such as low melting points (a fingerprint of ILs)²³, peculiar sequence of phase transitions over temperature^{24, 25}, and pressure-induced liquid-liquid transition²⁶. The conformational flexibility of TFSI⁻ was considered to be one of possible driving factors. Nevertheless, the flexibility is inferred from the conventional theory for isolated TFSI⁻ and not includes the impact of co-existing counteranions in ILs.

Herein, we combine *ab initio* molecular dynamics (AIMD) simulation with metadynamics^{27, 28} to explore the isomerization mechanism of isolated TFSI⁻. Our simulations reveal that the conventional static conformation scanning approach significantly underestimated the restriction and complexity of TFSI⁻ isomerization. TFSI⁻ exhibits dynamic conformational rigidity with diverse energy barriers (12–52 kJ mol⁻¹) over four distinct *trans-cis* isomerization pathways. An unexpected isomerization mechanism, namely dynamic $\pi \rightarrow \sigma^*$ negative hyperconjugation enabled coupled rotation of bond and bond angle has been discovered for the first time, competing with other two fundamental intermolecular motions including bond rotation and bond-angle inversion (Scheme 1c). The concerted process provides a low-energy pathway for TFSI⁻ to invert its helicity and form six diastereomeric conformers, rather than three pairs of enantiomers. We uncover that the overall stability of *cis*-TFSI⁻ linearly increases with the polarity of counteranions in the isolated ion pair state, by using metadynamics, an approach enhancing sampling while maintaining dynamic coupling with environments. The fractional population of *cis*-TFSI⁻ follows the same trend in bulk ILs, which is confirmed by both experiments and simulations. In particular, we find that melting points of TFSI⁻-based ILs is linearly dependent on the conformational rigidity of TFSI⁻ in the ion pair state, rather than *cis:trans* ratio reported in molecular systems, or configurational entropy in BF₄⁻-based ILs. This indicates that the conformational rigidity can be used as a fundamental dynamic descriptor to quantify the disorder potential of ILs and provide a new perspective on the origin of low T_m of ILs and a guideline for the rational design of low- T_m ILs. The counterintuitive isomerization behavior of TFSI⁻ and its correlation with distinct environments can also offer molecular level basis for multiple other unusual

thermal/mechanical/electrical properties of ILs, ranging from phase transition²⁴, phase-dependent interfacial capacitance²⁹, direct piezoelectric effect³⁰ to long-lived electric fields³¹.

Results and Discussion

Conformational space of isolated TFSI⁻. Figure 1a shows the torsional free-energy landscape over two torsion/dihedral angles for presenting the conformational space of isolated TFSI⁻. The free-energy surface features that (1) the minima (stable/metastable states) and saddle points (transition states) are not symmetric with respect to the 180-180 diganol; (2) the saddle points are unexpectedly higher than the global minima, namely, TFSI⁻ undergoes restricted isomerization with diverse and high energy barriers (12–52 kJ mol⁻¹). To explore the relationship between free energy and conformational structure, we take the structural snapshots corresponding to the characteristic points from metadynamics simulation. The conformations are displayed in the Newman projection, looking along the imaginary S-S axle (Figure 1 b). We have identified two independent pathways (**I & II**) for the *trans-trans* process and four *cis-trans* isomerization subpathways (**1–4**). In pathway **I**, starting with *trans*-B1, the coupled clockwise motion comprising S2–C1 bond rotation and 180° S–N–S bond-angle rotation (BAR) around the S-S axle, leads to the formation of *cis*-C1 **1**. Subsequently, S2–C1 bond continuously rotates to *trans*-D, whereas S–N–S bond angle just oscillates slightly relative to the horizontal plane. In pathway **II**, these two steps are reversed. Interestingly, the BAR occurring in each pathway induces the M(left-handed)-to-P(right-handed) helix inversion between *trans*-B1 (16 kJ mol⁻¹) and energetically more favorable *trans*-D (0 kJ mol⁻¹), accompanied by a unidirectional 300°

rotation of C1–S2 bond around the S–S axle. The unprecedented rotation/oscillation of S–N–S bond angle (Figure 1c), and the different sequences of these two motions in each pathway seem to enable all the conformers to be diastereomeric with each other, but why do the motions occur in TFSI⁻ and are the *cis*–*trans* free-energy barriers unexpectedly higher and diverse?

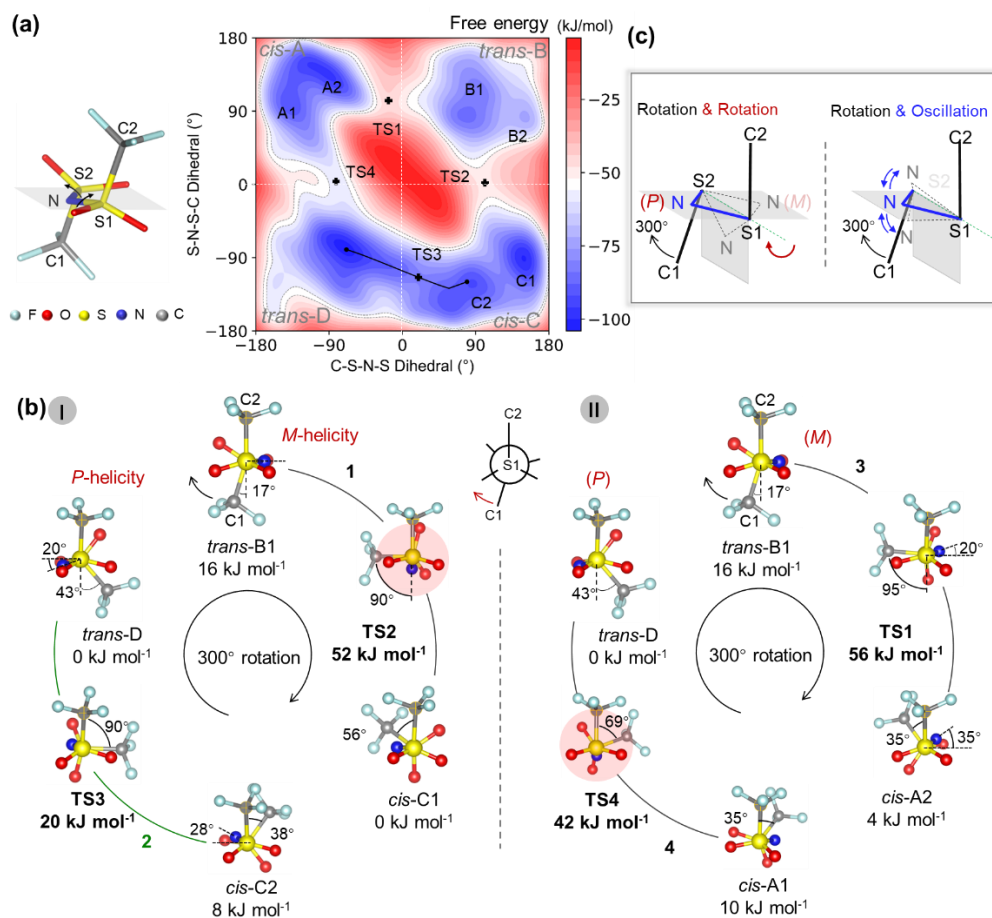


Figure 1. (a) Torsional free-energy landscape over two torsion angles/dihedrals (C1–S2–N–S1 and S2–N–S1–C2) of isolated TFSI⁻. The stable/metastable state A–D are separated by transition states (TSs). The white dashed lines for boundaries roughly separating the *cis* and *trans* conformer regions. (b) Newman projection of the structural snapshots corresponding to the characteristic points indicated in panel (a). Two distinct *trans*–*trans*

isomerization pathways (I: B–C–D, & II: B–A–D) and four subpathways (1–4) for *cis–trans* isomerization with relative energies. (c) Scheme of two types of motion adopted by S–N–S bond angle, accompanied by a unidirectional 300° rotation of S2–C1 bond: 180° out-of-plane bond-angle rotation (BAR) inducing M(left-handed)-to-P(right-handed) helix inversion (left) and oscillation on either side of the S1–C2 bond (right).

Negative hyperconjugation. Conjugation effect is known to play a critical role in stabilizing molecular structure and enlarging torsional energy barriers. The two lone pairs of electron from nitrogen, LP_A formed by deprotonation takes sp² hybridization, whereas LP_B is in a p orbital (Figure 2a, left up), distinct from the general sp³ hybridization in common sulfonamides³²⁻³⁴. Natural bond orbital (NBO) analysis shows that the π -type nitrogen lone pair (π_N) and oxygen lone-pair (π_O) NBOs overlap with $\sigma^*(S1-C2)$ antibond (left bottom). Their much larger interaction energies $E(2)$ (17.78 kcal mol⁻¹, 21.73 kcal mol⁻¹, and 23.70 kcal mol⁻¹), make π_N and π_O NBOs become the dominant contributors to the high NBO occupancy (0.34e) of $\sigma^*(S1-C2)$ (Table S2), in comparison with π_F (~9.0 kcal mol⁻¹). The negative $\pi_{O\&N} \rightarrow \sigma^*(S1-C2)$ hyperconjugation leads to $\pi_{O\&N} \rightarrow 3p(S)$ donation and further facilitates the formation of π_{SO} bonds and π_{SN} bond, and their conjugation among $-N^--SO_2-$ (middle up), reflected by the highest occupied molecular orbitals (HOMOs) of the extremely staggered and eclipsed conformations (middle bottom). The shorter length of S–N bond (~1.605 Å) (Table S2 shows bond lengths and angles for selective conformations of TFSI) also implies the presence of S–N π -bonding relative to the normal value of ~1.648 Å in sulfonamides^{33, 34}. Other electron donor-acceptor pairs may exist, but the $\sigma^*(S1-C2)$ acceptor orbital (0.14 a.u) is much more stable than others,

such as $\sigma^*(\text{S1-N})$ (0.21 a.u.) and $\sigma^*(\text{S1-O1})$ (0.40 a.u.) (Table S3), which renders the more favorable donation of π_{O} and π_{N} to $\sigma^*(\text{S1-C2})$.

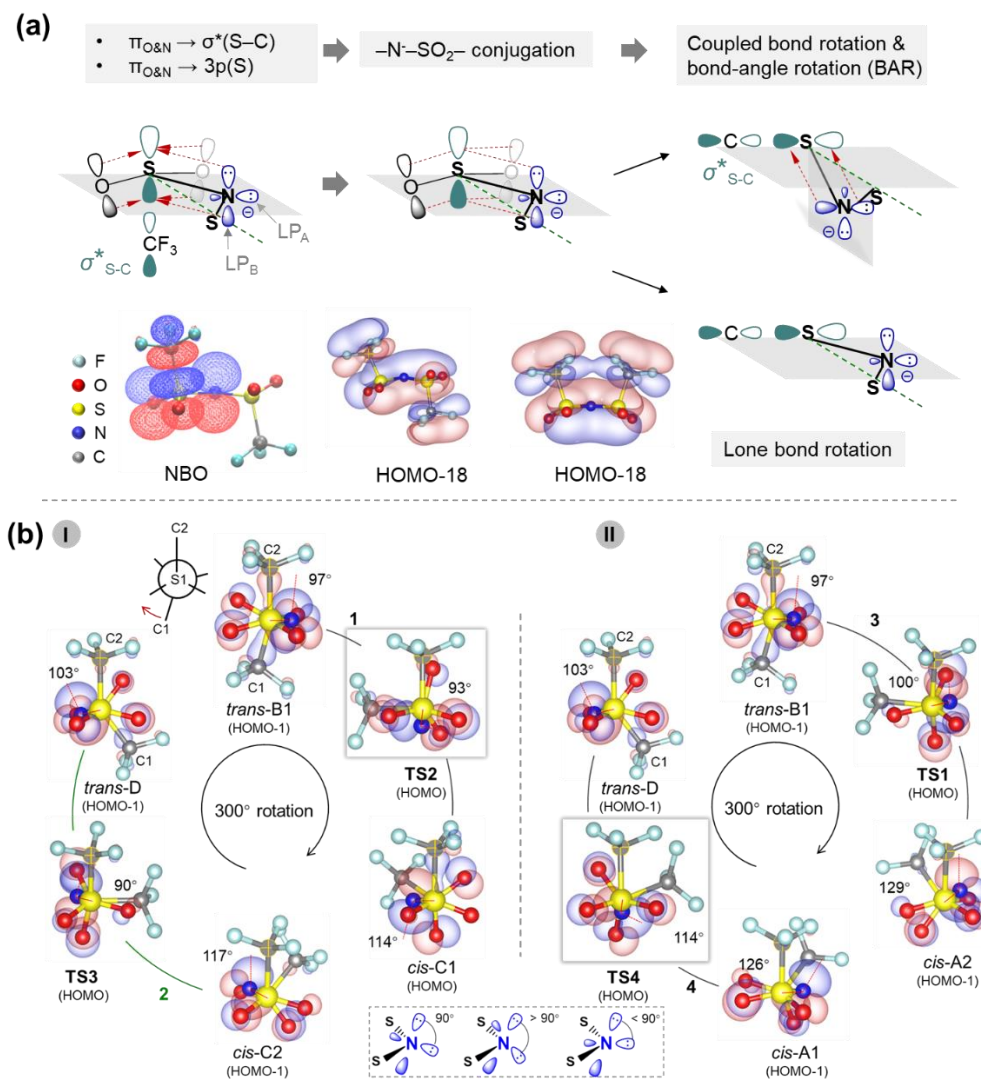


Figure 2. (a) Scheme for donation of nitrogen lone pair B (LP_B) and oxygen lone pairs which are in p orbitals, into the σ^* antibonding orbitals of S–C bond (left top). Overlapped natural bond orbitals (NBOs) on the extremely staggered conformation (isosurface value of 0.05) (left bottom). Π_{SO} bonds, π_{SN} bond, and their conjugation among $-\text{N}^--\text{SO}_2-$ (middle top), and highest occupied molecular orbitals (HOMOs) of the extremely staggered and eclipsed conformations (middle bottom) (isosurface value of 0.009). Alternative motions around the S–S axle: coupled rotation of S–C bond with S–N–S bond angle for

maintaining conjugation (right top), and lone S–C bond rotation for interrupting conjugation (right bottom). (b) HOMOs of characteristic conformations in two isomerization pathways (I & II) with isosurface value of 0.059. Inset, distortion of trigonal planar nitrogen.

The peculiar π -conjugation enables TFSI to carry out two possible types of internal motion around the S–S axle (Figure 2a): the unprecedented coupled bond rotation with bond-angle rotation (BAR) for maintaining conjugation (right top) and the lone bond rotation for interrupting conjugation (right bottom). The spatial evolution of molecular orbitals (MOs) (Figure 2b) elucidates that, in pathway **I**, LP_B is initially in a balanced position aligned with both S–C bonds in *trans*-B1. Subsequently, it only keeps aligned with S2–C1 bond when TFSI converts into TS2 by the coupled S–N–S BAR with S2–C1 bond rotation. Then, it tunes its aligned objective into S1–C2 bond during the TS2-to-*cis*-C1 process, leading to the decoupling of the rotary motions. By nitrogen oscillating, it tunes its alignment with S1–C2 bond in the following *cis*-C1-to-*trans*-B1 process, and finally settles in another position in line with both S–C bonds. In pathway **II**, the order of its alignment with one of S–C bonds is reversed. In both pathways, the trigonal planar structure of nitrogen undergoes geometric deformations to different extents (Figure 2b insets & Figure S1), but the energetically favourable π -electron delocalization is preserved.

The geometric structure of a molecule is also influenced by intramolecular steric repulsions. The steric clash between two aligned S–C bonds, S–C bond with S–O bond, and two S–O bonds are well-known, but the repulsion from nitrogen LP_A with S–C bonds and S–O bonds is easily overlooked (Figure S2). Furthermore, our Bader charge analysis (Table S4 & Figure S3) exhibits that each *cis-trans* isomerization process experiences

~0.3e intramolecular charge transfer mainly between carbon atom and fluorine atom, which suggests the dynamic repulsion between S–CF₃ moieties and S–O bonds. The net effect from the dynamic negative hyperconjugation and the five distinct steric repulsions drives TFSI⁻ to isomerize into six diastereomeric conformers with diverse and unusually high energy barriers, rather than three pairs of enantiomers reported by previous studies.¹⁴⁻¹⁸

Counteraction effect. Two typical counteranions, pyrrolidinium-based PYR₁₃⁺ and imidazolium-based BMIM⁺, are chosen to examine their impact on the isomerization of TFSI⁻ in the bulky ionic liquid state by performing AIMD simulations. Figure 3a–3f show the combined probability distribution of characteristic dihedrals of TFSI⁻ evolving with time in the bulk PYR₁₃TFSI and BMIMTFSI. The conformational distribution of TFSI⁻ indicates that, no *trans*-to-*cis* isomerization occurs in bulk PYR₁₃TFSI during 30 ps (Figure 3a–3c). In fact, some transition states between *cis*-C and *trans*-B even turn back to *trans* conformer. By contrast, in bulk BMIMTFSI (Figure 3d–3f), some TFSI⁻ anions initially adopt *trans* isomer or transition states in B region, but after 30 ps, they completely convert into *cis*-C. These observations point out the preferential stabilization of *cis*-TFSI⁻ in bulk BMIMTFSI than PYR₁₃TFSI, which conflicts the widely accepted *cis*: *trans* ratio of 1:1. In order to further quantify the conformational propensity of TFSI⁻ in the two bulk ILs, we compare their Raman spectra computed from AIMD simulation (Figure S4 & S5), which are not accessible by static calculations. The conformer population distribution (Figure 3g) derived from the simulated Raman spectra exhibits that 60% of TFSI⁻ adopt *trans* conformer in bulk PYR₁₃TFSI, whereas *cis*-TFSI⁻ occupies 60% of population in bulk BMIMTFSI. This prediction is confirmed by the experimental Raman spectra analysis

(Figure 3h & Figure S6). Specifically, *cis*-TFSI (67.11%) prevails in the bulk BMIMTFSI whereas it's less populated (41.96%) in bulk $\text{PYR}_{13}\text{TFSI}$, which are derived by deconvoluting characteristic peaks centered at 397 and 405 cm^{-1} for *trans*-TFSI and *cis*-TFSI (Figure 3i-3k).³⁵⁻³⁷

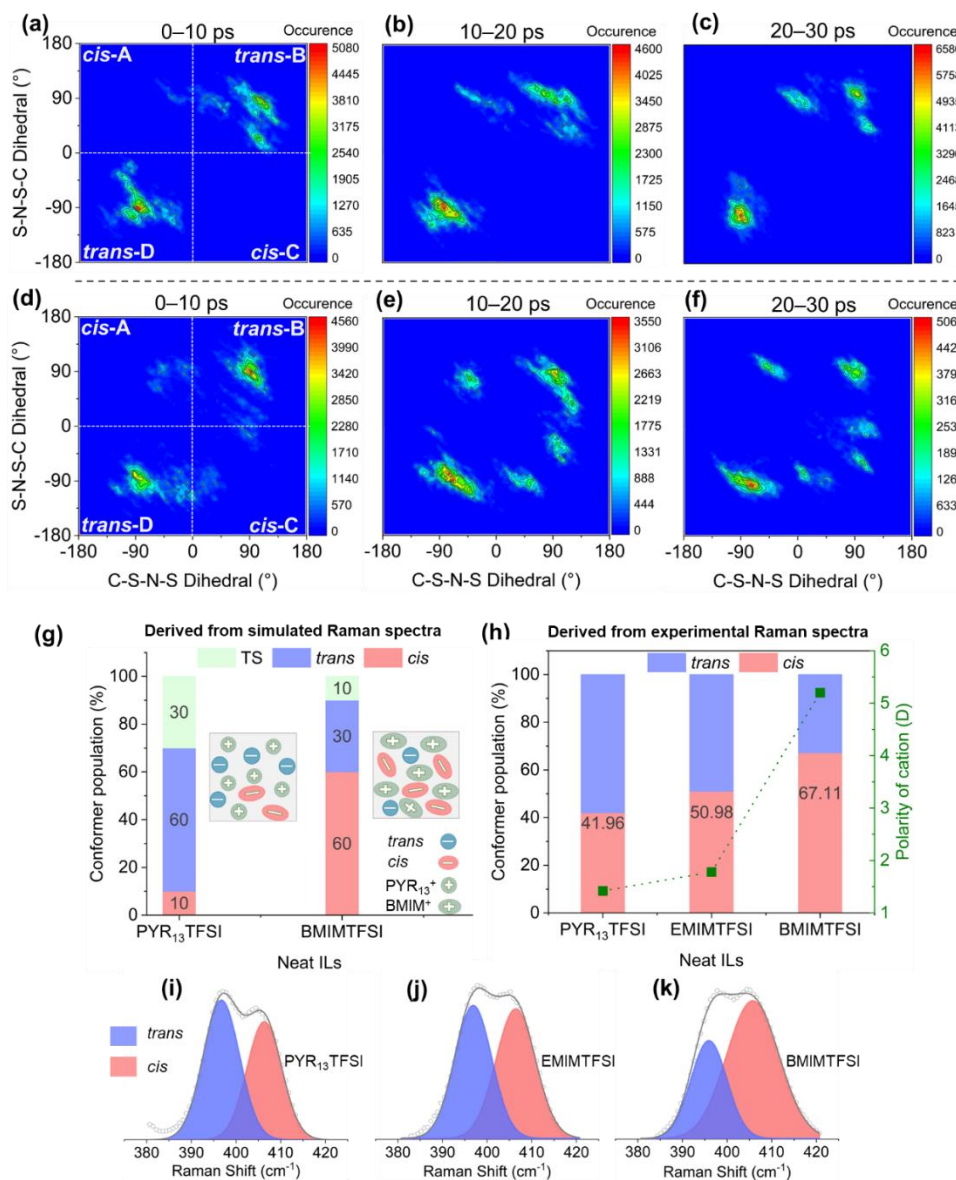


Figure 3. Combined dihedrals (C1-S1-N-S2 and S1-N-S2-C2) distribution of TFSI⁻ evolving over time (a) 0–10 ps, (b) 10–20 ps, (c) 20–30 ps in bulk PYR₁₃TFSI and (d) 0–10 ps, (e) 10–20 ps, (f) 20–30 ps in bulk BMIMTFSI. Fractional population distribution of TFSI⁻ conformers over cations with various polarities in the bulk ILs, derived from (g) simulated Raman spectra and (f) experimental Raman spectra. (Insets) Scheme for *trans/cis*-TFSI⁻ population distribution in bulk ILs.

To probe the nature of the countercation-dependent conformational preference of TFSI⁻, we analyzed the differences in structures and physicochemical properties between *cis*-TFSI⁻ and *trans*-TFSI⁻, PYR₁₃⁺ and BMIM⁺, respectively. *cis*-TFSI⁻ and *trans*-TFSI⁻ have almost identical polarizability (Table S5), but *cis*-TFSI⁻ has significantly larger dipole moment (4.44 D for *cis*-C1) than *trans*-TFSI⁻ (1.61 D for *trans*-D) (Table 1). The AIMD sampling trajectories show that both PYR₁₃⁺ and BMIM⁺ also adopt two main conformers in the bulky systems, which is induced by flexible side alkyl chains. *cis*-PYR₁₃⁺ (1.12 D) with a N-C-C-C dihedral of 65° is slightly less polar than *trans*-PYR₁₃⁺ (dihedral of 171°, 1.42 D) (Figure S7a). The 0.32:1 ratio of *cis*-PYR₁₃⁺ to *trans*-PYR₁₃⁺ in the accumulated frequency of occurrence indicates that *cis*-PYR₁₃⁺ is the minor conformer of PYR₁₃⁺ in bulk. By contrast, *cis*-BMIM⁺ (N-C-C-C dihedral of 60°, 4.44 D) has the probability of 61.24% whereas *trans*-BMIM⁺ (N-C-C-C dihedral of 180°, 5.96 D) possesses the probability of 38.76% in all possible conformations of BMIM⁺ (Figure S7b). The conformational geometry and distribution analysis of the countercations reveals that PYR₁₃⁺ is intrinsically weakly polar, independent of its isomers, whereas BMIM⁺ is strongly polar. Therefore, we infer that the polarity of counteraction contributes to the stereoselective isomerization of TFSI⁻ in TFSI⁻-based bulk ILs. The relationship is also

found in bulk EMIMTFSI (Figure 3h & 3g), where EMIM⁺ cation (1.13–1.78 D) has a shorter alkyl chain than BMIM⁺ (Figure S7d). To further support the inference, we also investigated the conformational distributions of BMIM⁺ and TFSI⁻ in bulk BMIMTFSI initiating with *cis*-TFSI⁻, compensating the narrow scope of sampling by AIMD simulation in the limited time. The majority of *trans*-BMIM⁺ (62.83%, 5.96 D) corresponds to the dominant population distribution of *cis*-TFSI⁻ in the latter bulk (Figure S8), whereas the minority of *trans*-BMIM⁺ (38.76%) matches the less population of *cis*-TFSI⁻ in the first bulk BMIMTFSI (Figure 3d–3f). This comparison uncovers the TFSI⁻-countercation mutual conformational selectivity and their coupled isomerization behavior. Altogether, we propose that the proportion of *cis*-TFSI⁻ in TFSI⁻-based bulk ILs is positively correlated with the polarity of countercations.

Table 1. Dipole moments for cationic/anionic components of ILs, and solvents in this work.

	Dihedral	Dipole moment (D)		
Isolated TFSI ⁻	5.06 (eclipsed)	4.44 (<i>cis</i> -C1)	1.61 (<i>trans</i> -D)	0.84 (staggered)
Interfacial TFSI ⁻	C-S-S-C	5.70 (<i>cis</i> , 15°)	5.03 (<i>cis</i> , 77°)	1.95 (<i>trans</i> , 153°)
EMIM ⁺	C-N-C-C	1.13 (8.6°)	1.78 (124°)	1.76 (159°)
BMIM ⁺	N-C-C-C	-	4.44 (<i>cis</i> , 60°)	5.96 (<i>trans</i> , 176°)
PYR ₁₃ ⁺	N-C-C-C	-	1.12 (<i>cis</i> , 65°)	1.42 (<i>trans</i> , 171°)
PYR ₁₄ ⁺	N-C-C-C	-	2.87 (<i>cis</i> , 65°)	3.27 (<i>trans</i> , 180°)
		Gas phase	Liquid phase	
NMP	-	4.03	4.06 (CFF) ³⁸	4.09 (Exp.) ³⁹
DMSO	-	3.99	4.96 (CFF) ⁴⁰	4.06 (Exp.) ⁴¹

CFF: Classical Force Field. Exp.: Experimental data. Interfacial TFSI⁻: BMIMTFSI-water interface starting with *cis*-TFSI⁻

To explore the intrinsic origin of the polarity-dependent conformational selectivity, the torsional free energy landscapes were sampled for TFSI⁻ in gaseous cation-anion pairs. For PYR₁₃TFSI (Figure 4a), *trans*-TFSI⁻ is the most stable conformer with the global minimum energy of -142 kJ mol⁻¹ in the D region, much lower than the local minimum of -116 kJ mol⁻¹ (A1) for *cis*-TFSI⁻. Other local minima corresponding to *trans*-TFSI⁻ are also deeper than the local minima for *cis*-TFSI⁻ (Table S6). Furthermore, the lowest $\Delta G^{\ddagger}_{\text{trans-cis}}$ increases to 54 kJ mol⁻¹ (B3-TS2-C1) from 20 kJ mol⁻¹ for isolated TFSI⁻ (D-TS3-C in Figure 1a), whereas the lowest $\Delta G^{\ddagger}_{\text{cis-trans}}$ holds the same value (12 kJ mol⁻¹, C2-TS3-D). Thus, PYR₁₃⁺ not only favors *trans*-TFSI⁻ thermodynamically, but also hinders the *trans*-to-*cis* conversion of TFSI⁻ kinetically. For BMIMTFSI (Figure 4d), *cis*-C (-84 kJ mol⁻¹) demonstrates the global minimum in the free energy profile whereas *trans*-TFSI⁻ is slightly less stable at the local minima of B (-80 kJ mol⁻¹) and D (-76 kJ mol⁻¹), which implies that BMIM⁺ supports *cis*-TFSI⁻ thermodynamically. Moreover, the lowest $\Delta G^{\ddagger}_{\text{trans-cis}}$ keeps the same value (20 kJ mol⁻¹, D-TS4-A) with that of isolated TFSI⁻, whereas the lowest $\Delta G^{\ddagger}_{\text{cis-trans}}$ increases to 24 kJ mol⁻¹ (A-TS4-D). That is, BMIM⁺ exerts completely opposite control on the TFSI⁻ isomerization in both thermodynamics and kinetics in comparison with PYR₁₃⁺. The conformational energy surfaces of TFSI⁻ are also dramatically modulated by another two moderately polar cations, EMIM⁺ and PYR₁₄⁺ (2.87–3.27 D) (Details in Table S6 and Figure S7). To more clearly quantify the impact of counteranions with various polarities on the stabilization of TFSI⁻ conformers, we extract all energy minima from the complex landscapes and calculate the ratio of *cis*-TFSI⁻ to *trans*-TFSI⁻ in overall energies ($r = E_{\text{cis}}/E_{\text{trans}}$). Figure 4e displays that the ratios are in a linear relationship with the polarities of cations from the four ion pairs. This result is excellently consistent

with the correlation between fractional population of *cis*-TFSI⁻ and polarities of cations, which is found in the bulky systems (Figure 3h).

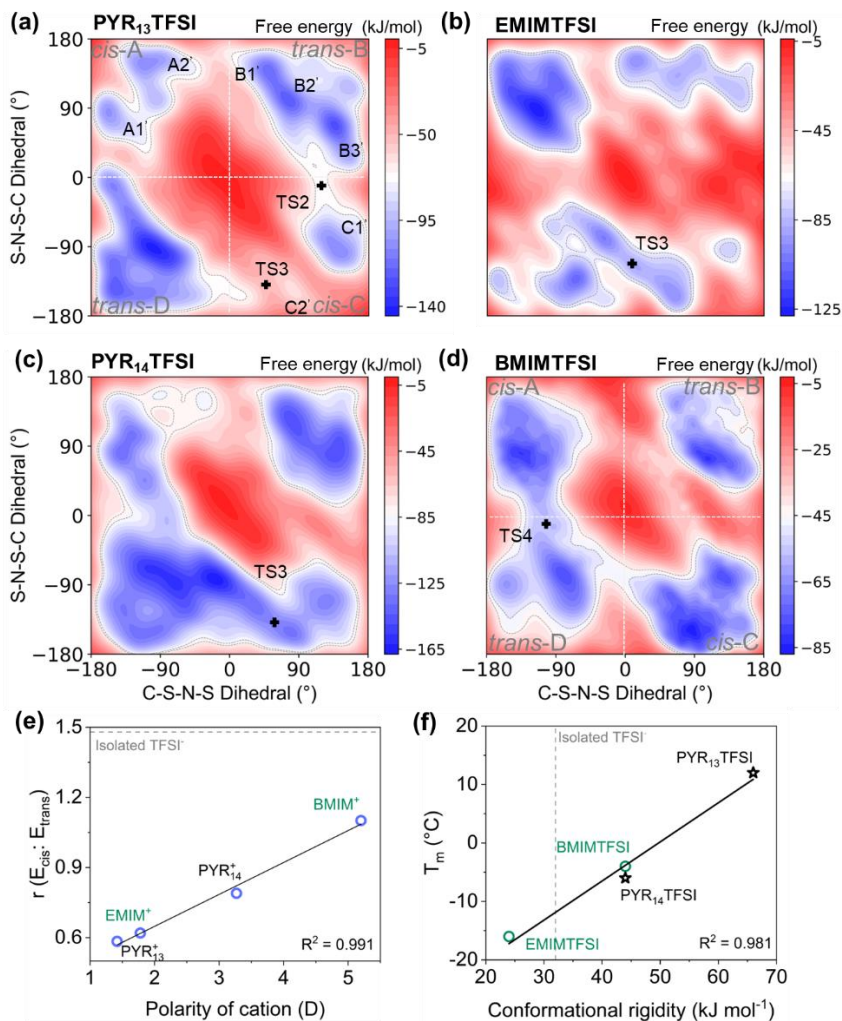


Figure 4. Torsional free-energy landscapes of TFSI⁻ in gaseous ion pair of (a) PYR₁₃TFSI, (b) EMIMTFSI, (c) PYR₁₄TFSI, and (d) BMIMTFSI. (e) Ratio of *cis*-TFSI⁻ to *trans*-TFSI⁻ in overall energies ($r = E_{cis}/E_{trans}$) as a function of the polarity of counteranions. The overall free energy is a sum of all energy minima. (f) Correlation of ILs melting point (T_m) with the conformational rigidity of TFSI⁻ in ion pairs.

Conformational rigidity of TFSI⁻. *cis* alkenes, the molecular systems, tend to have lower melting points (T_m) than *trans* alkenes due to their higher polarity and lower symmetry.^{42, 43} Although we have confirmed that the proportion of *cis*-TFSI increases with the polarity of counteranions, the melting points don't drop down as the general trend. In fact, PYR₁₄TFSI/EMIMTFSI is the minimum turning point of the curve displaying melting points of pyrrolidinium-based (green line) or imidazolium-based (blue line) ILs over the alkyl-chain length of counteranion (Figure S9). By quantitatively comparing ILs with alkali halides, a latest study⁴⁴ proposes that the characteristic low T_m of ILs, relative to other salts, mainly origin from the structural entropy (configurational and conformational entropy), particularly configurational/interionic entropy, rather than the widely believed enthalpy (Coulombic interaction and lattice energy). However, only conformational entropy from the cations (EMIM⁺ & BMIM⁺) is considered in that study, since anion PF₆⁻ is rigid and not deformed. By contrast, TFSI-based ILs are much more complex in additional conformational diversity of TFSI⁻, interionic conformational selectivity and coupled isomerization, and configurational diversity. Accordingly, it's challenging to decompose enthalpy/entropy into individual terms, and consequently identify their role in the low T_m of ILs. Here, we find that the conformational rigidity (a sum of the lowest $\Delta G^{\ddagger}_{\text{trans-cis}}$ and $\Delta G^{\ddagger}_{\text{cis-trans}}$) of TFSI⁻ in the ion pair state, is a fundamental descriptor to quantify the impact of counteranions on the melting points of TFSI-based ILs (Figure 4f). The free-energy-based dynamic descriptor comprehensively including the enthalpic and entropic contribution from intraionic and interionic interactions, delineates the disorder potential of TFSI-based ILs under thermal stimuli.

Solvent effect. Although the applications of the electrolytes comprising TFSI⁻-based ILs⁴⁵ or Mⁿ⁺(TFSI⁻)_n (M = Li, Mg, Al ...) salts⁴⁶⁻⁴⁸ with solvents are ubiquitous in electrochemical energy storage systems, there are no reports on quantifying the conformational proportion of TFSI⁻ in the solutions, which potentially influences the thermal/electrochemical properties of electrolytes, such as stability, interfacial capacitance and ionic conductivity.⁴⁹⁻⁵¹ We collected the Raman spectra of the solutions comprising TFSI⁻-based ILs with polar, aprotic solvents, and quantified the impact of solvents on the population distribution of *cis/trans* conformers of TFSI⁻ at various concentrations. Figure 5a shows that the ratio of *cis*-TFSI⁻ in the three solutions containing PYR₁₃TFSI, EMIMTFSI and BMIMTFSI are all raised by 1-Methyl-2-pyrrolidone (NMP) (4.09 D and Figure S10a for the Raman spectrum of neat NMP), evidenced by the higher peak intensity at 405 cm⁻¹ relative to 397 cm⁻¹ (left three), and the enlarged fractional population of *cis*-TFSI⁻ corresponding to the decreased concentration of ILs (right one). Even though for PYR₁₃TFSI with weakly polar cation and high $\Delta G_{\text{trans-cis}}^{\ddagger}$ (≥ 54 kJ mol⁻¹), the increased fractional population of *cis*-TFSI⁻ reaches 27% at the concentration of 1: 5 (molar ratio). Another solvent, namely dimethyl sulfoxide (DMSO) (4.06 D and Figure S10b for the Raman spectrum of neat DMSO) exhibits its better promotion performance in the larger raise of *cis*-TFSI⁻ fractional population (35%) in PYR₁₃TFSI-DMSO solution (molar ratio of 1: 5) relative to neat PYR₁₃TFSI, making *cis*-TFSI⁻ fully dominate in the solution (Figure 5b). We also examined the torsional free energy surface of TFSI⁻ in the ion pair of PYR₁₃TFSI dissolved in bulk DMSO. Figure 5c–5d indicate *cis*-TFSI⁻ is more stable in the solution with two lower local energy minima of -134 kJ mol⁻¹ (A2' & C'), in contrast to -116 kJ mol⁻¹ (A1) and -108 kJ mol⁻¹ (C1) in the ion pair of PYR₁₃TFSI. As a result, the

minimum $\Delta G_{\text{trans-cis}}$ drops to 8 kJ mol⁻¹, versus 26 kJ mol⁻¹ in the neat PYR₁₃TFSI. Moreover, the *trans*-to-*cis* isomerization is catalyzed by DMSO through the energy pathway of B1'–TS1'–A2' (minimum $\Delta G_{\text{trans-cis}}^\ddagger = 12$ kJ mol⁻¹), saving 78% of the energy and even lower than that in isolated TFSI (≥ 20 kJ mol⁻¹). The $\Delta G_{\text{trans-cis}}^\ddagger$ in other pathways also drop to different degrees. On the other hand, the minimum $\Delta G_{\text{cis-trans}}^\ddagger$ jumps from 12 kJ mol⁻¹ (C2–D, gaseous PYR₁₃TFSI) to 42 kJ mol⁻¹ (A1'–TS4'–D2'). The energy analysis reveals that strongly polar solvents stabilize *cis*-TFSI, catalyze the *trans*-to-*cis* process, and hinder the reverse transition, which breaks most of the constraints exerted on TFSI by PYR₁₃⁺.

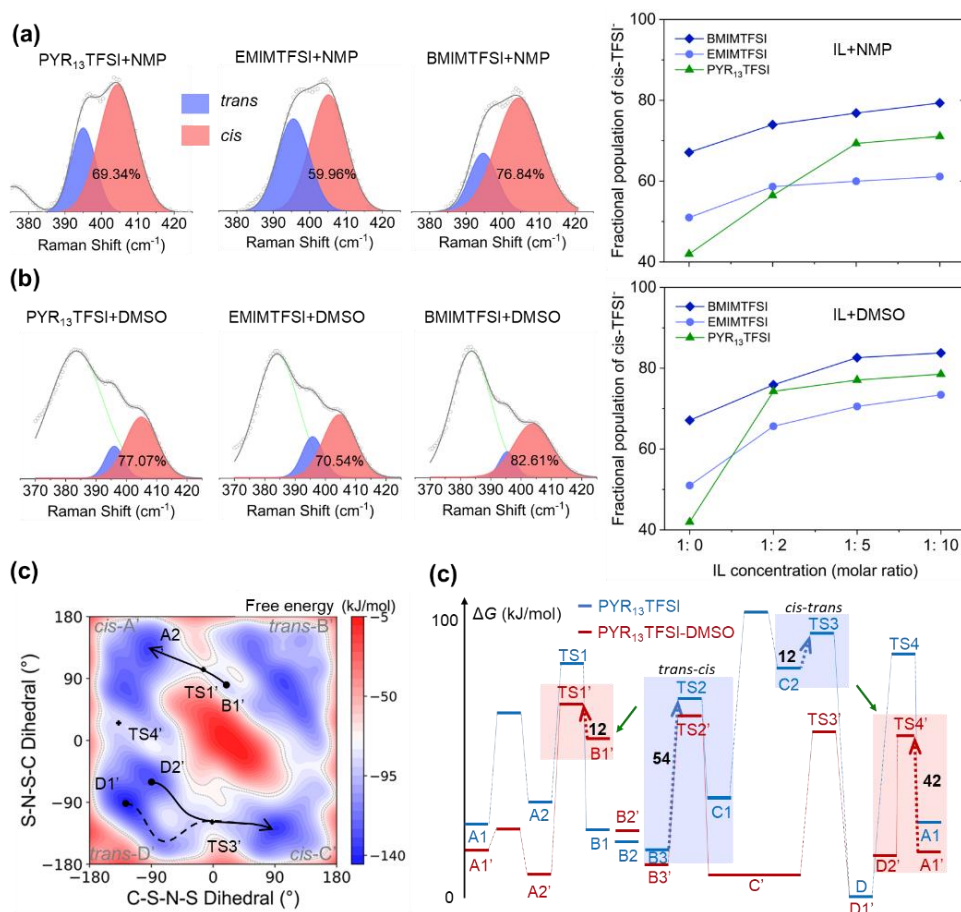


Figure 5. (a & b) Characteristic Raman spectra of IL-solvent solutions at molar ratio of 1: 5 and IL concentration-dependent fractional population of *cis*-TFSI⁻, derived from peak deconvolution. The ILs include PYR₁₃TFSI, EMIMTFSI and BMIMTFSI. NMP, 1-Methyl-2-pyrrolidone. DMSO, dimethyl sulfoxide. (c) Torsional free-energy landscape of TFSI⁻ in ion pair of PYR₁₃TFSI dissolved in bulk DMSO (explicit solvent model) and its comparison with TFSI⁻ isomerization in neat ion pair of PYR₁₃TFSI (d). Red and blue regions display the minimum energy pathways.

Local environment effect. Based on the understanding of the strong TFSI⁻-cation (two-body) interaction, we have extended our exploration to the TFSI⁻-cation-solvent (many-body) system. Although the dipolar interactions are inherently anisotropic⁵² in the IL-solvent solutions, the environment of the ion pair is homogeneous, since each solvent molecule is identical in shape and polarity. Nevertheless, bulk TFSI⁻-based ILs consisting of two strongly interacting components with deformable structures and diverse polarities, exhibit additional anisotropy and heterogeneity, which further complicates the analysis of TFSI⁻ isomerization in the ion pair dissolved in the “TFSI⁻-based IL solution”. The trajectories of two bulky BMIMTFSI systems initiating with *trans*-TFSI⁻/*cis*-TFSI⁻, show that both stay as *trans* conformer during 30 ps (Figure S11a–11c & Figure S12a–12c), but *trans*-TFSI#1 is surrounded by one *cis*-TFSI⁻, one TS and four *trans*-TFSI⁻ (Figure 6a, left), whereas *trans*-TFSI#1* is envired by five *cis*-TFSI⁻ and one TS (Figure 6b, left). By metadynamics approach again, we can obtain the whole conformational space of *trans*-TFSI#1 and *trans*-TFSI#1* while maintaining their coupling with environments and characterize the impact of surrounding ions on their isomerization. The conformational free-energy surface of *trans*-TFSI#1 (*trans*-B) seems to resemble the landscape of gaseous

BMIMTFSI ion pair (Figure 4d), but the energy difference between *cis*-C and *trans*-B increases from 4 kJ mol⁻¹ to 12 kJ mol⁻¹ while the lowest $\Delta G^\ddagger_{\text{trans-cis}}$ or $\Delta G^\ddagger_{\text{cis-trans}}$ jumps by 20 kJ mol⁻¹ (Figure 6a, right). In a sharp contrast, *trans*-TFSI#1* (*trans*-D) overmatches its competing *cis* conformers with the environment in a large energy difference of 28–42 kJ mol⁻¹ and high $\Delta G^\ddagger_{\text{trans-cis}}$ of 46–70 kJ mol⁻¹ (Figure 6b, right). These simulation results signify that the local environment in the bulky TFSI-based IL systems also influence the conformational stability and rigidity of TFSI⁻ by net electrostatic interactions, in addition to the counteranions, suggesting the complex many-body correlation^{52, 53}.

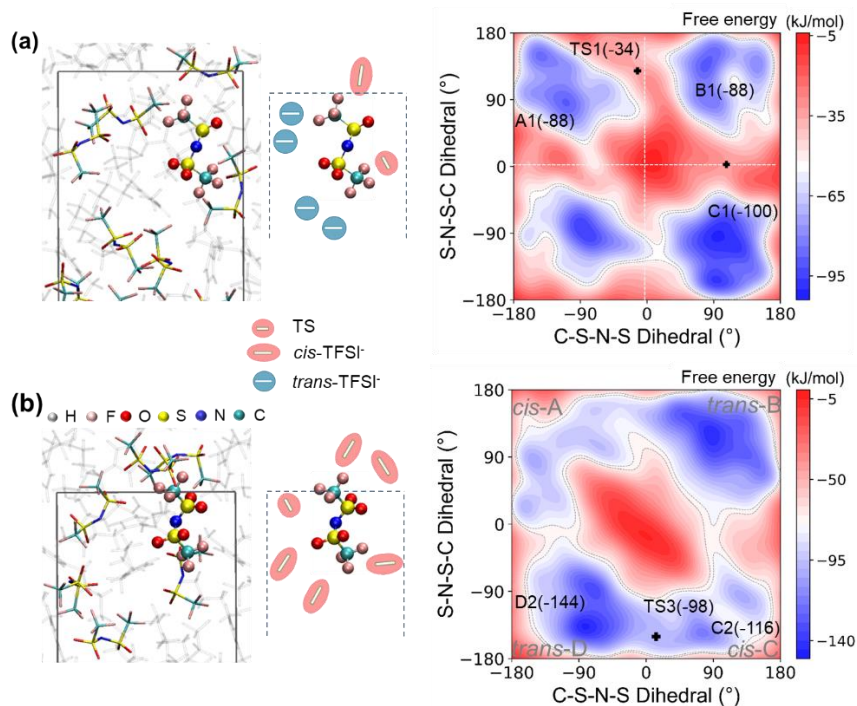


Figure 6. Snapshots (left) at 30-ps AIMD trajectories and torsional free-energy landscapes (left) of TFSI#1 (a) in the bulk BMIMTFSI initiating with *trans*-TFSI, and TFSI#1* (b) in the bulk BMIMTFSI initiating with *cis*-TFSI. Black solid line and blue dashed line (scheme) illustrating the unit cell. Cations are displayed in grey stick model.

Conclusions

The conventional static electronic structure calculation of isolated TFSI⁻ showed that the sole and extremely low energy barrier between *cis* and *trans* conformer endowed the conformational flexibility of TFSI⁻, which might contribute to the unusual thermal and mechanical properties of ILs. By using AIMD combined with metadynamics, we revealed that gaseous TFSI⁻ experiences restricted *cis-trans* isomerization following four distinct pathways with strikingly higher and diverse free-energy barriers. We have identified an unprecedented coupled motion comprising bond rotation and bond-angle rotation at the central N⁻, which competes with other two isomerization mechanisms in TFSI⁻, a system with bicoordinated heteroatoms. Discovering theoretical evidence for the 180° bond-angle rotation also brings our knowledge one step further towards an alternative strategy for controlling the molecular helical inversion, dynamic chirality and unidirectional rotation of molecular motor. The negative $\pi_{\text{O\&N}} \rightarrow \sigma^*(\text{S-C})$ hyperconjugation leads π electrons to delocalize among $-\text{SO}_2-\text{N}^--\text{SO}_2-$ and allows TFSI⁻ to carry out the coupled motion for maintaining conjugation. The net dynamic effect from the negative hyperconjugation and five distinct steric repulsions drives TFSI⁻ to rigidly and diversely isomerize into six diastereomeric conformers, rather than three pairs of enantiomers.

Counteranions can significantly control the conformational stability and rigidity of TFSI⁻. The overall stability of *cis*-TFSI⁻ linearly increases with the polarity of counteranions in the isolated ion pair state. The fractional population of *cis*-TFSI⁻ follows the same trend in bulk ILs, which is confirmed by both experiments and simulations. The

conformational rigidity of TFSI⁻ in the ion pair, a free-energy-based dynamic descriptor, can excellently quantify the impact of countercations on the T_m of TFSI⁻-based ILs, owing to the inclusion of cation-anion coupling. This relationship provides new fundamental insights into the origin of low T_m of ILs. The unified descriptor can serve as a guideline towards the rational design of low- T_m ILs. The similar effects including *cis*-TFSI⁻ stabilization, *trans*-to-*cis* catalysis, and reverse transition hinderance are also found in strongly polar solvents which overcomes most of the constraints imposed on TFSI⁻ by countercations. The distinct local environments in the highly anisotropic and heterogeneous TFSI⁻-based IL bulks can exert different additional influence on the isomerization behavior of TFSI⁻ by net electrostatic interactions. This feature in association with interionic mutual conformational selectivity and coupled isomerization imply that TFSI⁻-based ILs are the complex many-body strongly correlated systems.

The counterintuitive isomerization behavior of TFSI⁻ and its correlation with distinct environments can also provide molecular level basis for multiple other exotic thermal/mechanical/electrical properties of ILs. (1) The diversity in the relative stability of conformers, isomerization pathway and energy barrier of TFSI⁻ leads to the peculiar sequence of phase transitions over temperature, phase-dependent transport characteristics and interfacial capacitance. (2) Pressure forces *trans*-to-*cis* conversion (helical folding) of TFSI⁻, cation-anion reconfiguration, and re-orientation of those dipoles, thus induces liquid-liquid transition and direct piezoelectric effect, a first report in a neat liquid. (3) The alignment of ion dipoles also forms long-lived electric fields of IL bilayers under external electric field pulse.

ASSOCIATED CONTENT

Supporting Information

The Supplemental Information (including Raman spectroscopy characterization, computational details, Tables S1–S6, and Figures S1-S12) is available free of charge at <https://pubs.acs.org/doi/>

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L.Y.: conceptualization, simulation and experimental characterization; **A.R.:** experiments design and writing editing; **D.K.:** supervision, writing editing; **H.B.:** supervision; **T.Z.:** conceptualization, supervision, writing editing, funding acquisition, resources.

Notes

The authors declare no competing financial interests.

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