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Force-induced retro-click reaction of triazoles competes with adjacent single-bond rupture†

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The highly controversial force-induced cycloreversion of 1,2,3-triazole, its well-known retro-click reaction, is shown to be possible only for 1,5-substituted triazoles, but competes with rupture of an adjacent single-bond. We draw this conclusion from both static and dynamic calculations under external mechanical forces applied to unsubstituted and 1,4- and 1,5-substituted triazoles. The JEDI (Judgement of Energy DIstribution) analysis, a quantum chemical tool quantifying the distribution of strain energy in mechanically deformed molecules, is employed to identify the key factors facilitating the force-induced retro-click reaction in these systems. For 1,4-substituted triazoles it is shown to be impossible, but the parallel alignment of the scissile bond in 1,5-substituted triazoles with the acting force makes it generally feasible. However, the weakness of the carbon–nitrogen bond connecting the triazole ring to the linker prevents selective cycloreversion.

1 Introduction

The 1,3-dipolar cycloaddition of an azide and an alkyne falls into the category of click chemistry.^{1,2} The resulting 1,2,3-triazoles are versatile building blocks finding numerous applications in biology^{3–9} as well as in polymer and materials science.^{9–15} Considering the thermodynamic stability of the triazole moiety, it came as a surprise when sonochemical cycloreversion of 1,2,3-triazoles incorporated in poly(methyl acrylate) was presented in 2011 (Fig. 1).¹⁶ As polymers in an ultrasound bath are known to be subject to mechanical forces,^{17–22} this reaction was interpreted as a mechanically induced cycloreversion. Although the original publication¹⁶ has been retracted,²³ the mechanical cycloreversion of triazoles has attracted considerable interest within the past few years.²⁴

Using sonochemical experiments and extended Bell theory,²⁵ for example, the application of an external force to the triazole ring was found to indeed lower the activation energy barrier for cycloreversion.²⁶ In a purely theoretical work, on the contrary, the cycloreversion has been shown to be only favored for pulling forces below 1 nN and rupture at the α bond to be preferred at higher forces.²⁷ It was concluded that mechanical force alone is insufficient to explain cycloreversion of 1,2,3-triazoles. In yet another computational study of triazoles resulting from strained alkynes, the pulling geometry was identified as the key factor for successful cycloreversion.²⁸ The possibility of metal-assisted mechanical cycloreversion of triazoles has also been discussed recently.²⁹ Experimentally, cycloreversion was

observed for forces below 860 pN in an Atomic Force Microscope (AFM).³⁰ On the other hand, in another AFM study, it was not possible to determine reliably whether cycloreversion did indeed take place, but it was pointed out that forces in the nN regime would be needed.³¹ As can be seen from such contradictory experimental and theoretical findings, the feasibility of mechanical cycloreversion of 1,2,3-triazoles is widely debated.²⁴

Here we apply static quantum chemical calculations under a constant stretching force^{32–34} in combination with the JEDI (Judgement of Energy DIstribution) analysis.^{35–37} This allows us to quantify the distribution of mechanical strain energy in deformed molecules in general, and here to answer the specific question under which circumstances cycloreversion of 1,4- and 1,5-substituted triazoles is possible. Although the JEDI analysis is based on the harmonic approximation, it has been used successfully to gain detailed insights into the rupture mechanisms of knotted polymer strands³⁸ and mechanophores.³⁹ We first study several pulling coordinates within the isolated triazole to identify relevant coordinates triggering cycloreversion most efficiently. Then the mechanical properties and possible cycloreversion of triazoles substituted with aliphatic linkers are investigated. Therefore, the linkers are pulled in opposite directions to simulate the mechanical force acting in sonochemical experiments that is transmitted to the triazole *via* the linkers. In addition, Born–Oppenheimer Molecular Dynamics (BOMD) calculations under external forces are performed to simulate the experimental conditions more realistically. From an energetic point of view, the cycloreversion of 1,4-substituted triazoles is strongly disfavored. 1,5-Substituted triazoles, on the contrary, can in principle undergo cycloreversion, but this process competes with rupture of the carbon–nitrogen single-bond connecting the triazole ring to the aliphatic linker.

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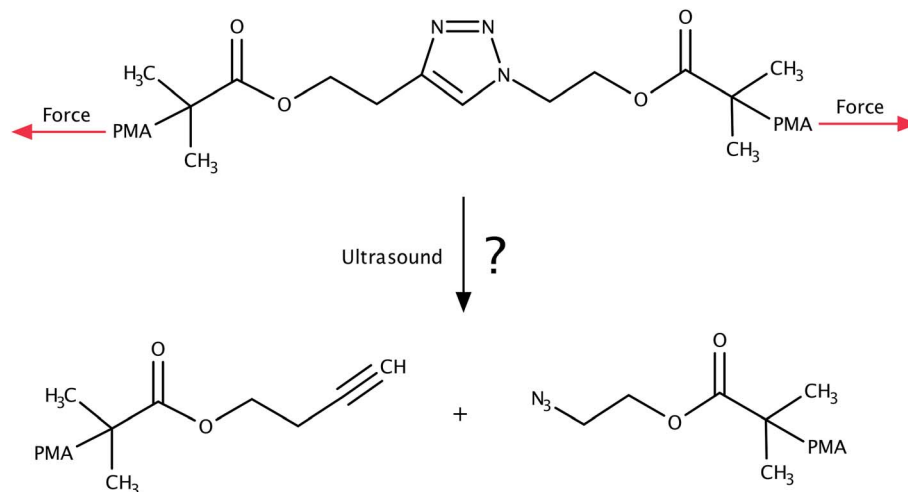


Fig. 1 System investigated in ref. 16, in which the mechanochemical retro-click reaction of triazole that is substituted by poly(methyl acrylate) (PMA) chains in an ultrasound bath was reported. We investigate the feasibility of this reaction in this paper.

2 Computational methods

Static isotensional stretching^{32–34} was simulated using the EFEI (External Force is Explicitly Included)³³ method as implemented in the Q-Chem 4.3 program package.⁴⁰ Bond rupture forces were determined iteratively with a precision of 50 pN. For the Born–Oppenheimer Molecular Dynamics (BOMD) simulations, the Q-Chem 4.0.1 program package⁴¹ was used. In the BOMD simulations, forces were applied to the terminal carbon atoms of the linkers in a manner analogous to the EFEI approach. All calculations were carried out using Density Functional Theory (DFT)^{42,43} at the PBE⁴⁴/cc-pVDZ⁴⁵ level of theory, since this method has been shown to deliver satisfactory results in various mechanochemical pulling scenarios within reasonable computation times.^{46,47} We used custom Python routines interfaced with the Q-Chem program package to carry out the JEDI analysis. In particular, the strain energies in all internal coordinates *i*, *i.e.* all bonds, bendings and torsions of the molecule, were calculated *via*^{33–37}

$$\Delta E_i = \frac{1}{2} \sum_j^M \frac{\partial^2 V(\vec{q})}{\partial q_i \partial q_j} \bigg|_{\vec{q}=\vec{q}_0} \Delta q_i \Delta q_j. \quad (1)$$

Here, $\frac{\partial^2 V(\vec{q})}{\partial q_i \partial q_j} \bigg|_{\vec{q}=\vec{q}_0}$ is the Hessian matrix in redundant internal coordinates at the equilibrium geometry and Δq_i is the change in internal coordinate *i* upon mechanical deformation. *M* denotes the number of internal coordinates in the molecule. As the JEDI analysis is based on the harmonic approximation, it is useful to compare the harmonic strain energy ΔE_{harm} , *i.e.* the sum of the individual contributions ΔE_i , to the energy difference ΔE_{DFT} between the relaxed and the strained geometry, calculated with DFT. This allows an error estimation of the harmonic approximation.

3 Results and discussion

3.1 Static stretching

In a previous study,²⁸ a number of different pulling geometries in triazoles resulting from strained alkynes were investigated. Here, on the contrary, we first concentrate on several conceivable pulling geometries in isolated, unsubstituted 1,2,3-triazole to gain fundamental insight into the mechanical properties of triazole itself, and subsequently move to triazoles substituted by alkyl chains in order to model the original experimental setup¹⁶ more realistically. The coordinates tested (Fig. 2) include a symmetric expansion of the triazole ring (the “breathing” coordinate c_1), two distinct shearing coordinates (c_2 and c_3), two additional symmetric stretching coordinates, in which the azide is being removed from the alkyne (c_4 and c_5), and two unzipping coordinates (c_6 and c_7).

At the PBE/cc-pVDZ level of theory, only four of the seven investigated coordinates (c_3 and c_5 – c_7) lead to cycloreversion. Of these coordinates, cycloreversion *via* c_3 requires approximately twice the applied force (12.85 nN) and energy (2.05 eV) that is needed for cycloreversion *via* the other coordinates. This finding can be rationalized using the JEDI analysis (Fig. 2), which reveals that the entire triazole ring needs to be mechanically activated for cycloreversion *via* c_3 . Upon inspection of the results of the JEDI analysis for the coordinates c_5 – c_7 , by contrast, it becomes apparent that it is sufficient to activate one of the carbon–nitrogen bonds in the triazole ring to trigger cycloreversion. In the case of c_5 , both carbon–nitrogen bonds are activated and therefore the force needed to stretch each one of them (5.55 nN) is particularly low. However, since two bonds are elongated, the energy that is needed for cycloreversion (1.14 eV) is higher than in the case of c_6 (0.97 eV) and c_7 (0.81 eV). Moreover, the N3–C4 bond is especially weak, since it stores more energy in c_5 than the N1–C5 bond. In other words, stretching by the same force leads to a storage of more energy in the N3–C4 bond. This picture is supported by a comparison of



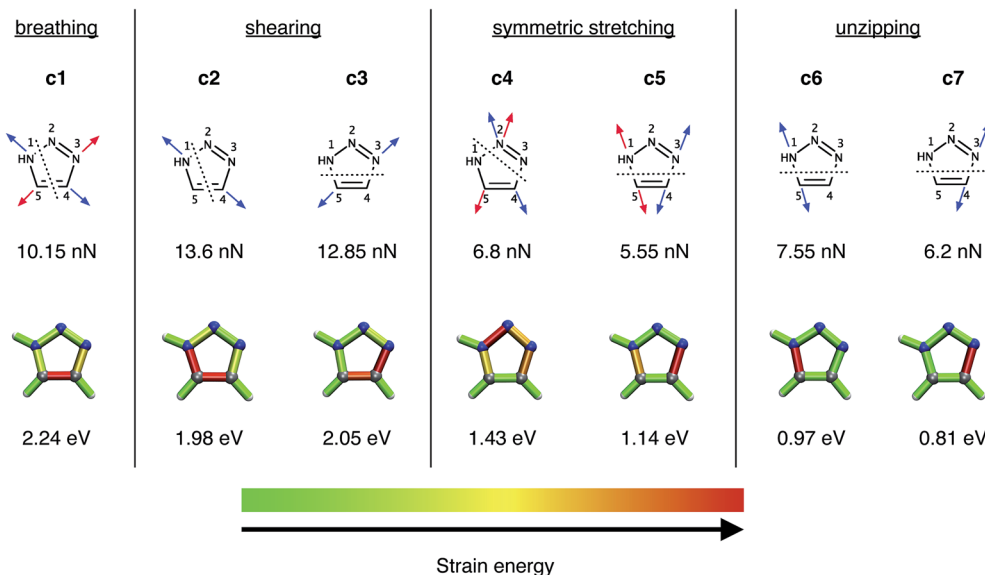


Fig. 2 The investigated breathing coordinate c_1 , the shearing coordinates c_2 and c_3 , the symmetric stretching coordinates c_4 and c_5 as well as the unzipping coordinates c_6 and c_7 . Forces were applied to the atoms indicated by the arrows, with arrows of the same color signifying the same EFEI stretching coordinate. Dotted lines indicate scissile bonds. The rupture forces are given below the coordinates. The results of the JEDI analysis are shown qualitatively by color-coded structures (gray : carbon, blue : nitrogen, white : hydrogen), the generation of which is described in detail in ref. 37. The rupture energies ΔE_{DFT} , calculated at the PBE/cc-pVDZ level of theory, are given below the color-coded structures.

the unzipping coordinates c_6 and c_7 , since more force and energy is needed for rupture of the N1–C5 bond *via* c_6 than the N3–C4 bond *via* c_7 . In the coordinates c_5 – c_7 , more than 90% of the strain energy is stored in the scissile bonds, while in the other stretching coordinates multiple bonds and bendings in the triazole ring store significant amounts of strain energy. Hence, cycloreversion proceeds only very inefficiently (c_3) or is not triggered at all (c_1 , c_2 and c_4).

Since the N3–C4 bond is ruptured most easily by an external force *via* c_7 and, as a result, the lowest amount of energy is needed for cycloreversion, the specific activation of this bond in experimental setups would lead to cycloreversion most efficiently. However, it is hard to imagine an experimental setup involving 1,4- or 1,5-substituted 1,2,3-triazoles, in which the N3–C4 bond is stretched in an isolated or at least in a preferred manner. Similar restrictions apply for the mechanically favored coordinate c_5 , *i.e.* simultaneous rupture of both carbon–nitrogen bonds. The bond that can arguably be activated most easily in an experiment is the N1–C5 bond by adding aliphatic linkers in a 1,5-substitution pattern. Upon stretching such molecules end-to-end, it is reasonable to expect a significant activation of the N1–C5 bond in analogy to the unzipping coordinate c_6 . End-to-end stretching in the case of a 1,4-substitution, however, resembles the shearing coordinate c_2 most closely, which may not lead to cycloreversion. Such scenarios are discussed in detail below.

In closing the discussion of the stretching coordinates within the isolated triazole ring it should be noted that the mere presence of a linker influences the forces and energies that are needed for rupture in a nontrivial manner (*cf.* the ESI, Fig. S1†). The linkers do not only fulfill the role of a transmitter of strain

energy from the environment to the triazole ring, but have an electronic influence as well. Similar effects have been documented previously in literature.^{48,49}

To simulate experimental conditions more realistically, alkyl linkers of different lengths (ethyl to hexyl) were introduced at positions 1 and 4 or 1 and 5 of the triazole ring, respectively (Fig. 3). The JEDI analysis reveals the triazole ring of the 1,4-isomer to be hardly strained when the molecule is stretched end-to-end (Fig. 3A, top). As a result, cycloreversion was not observed in any of the 1,4-substituted triazole systems under investigation. In the 1,5-isomer, on the contrary, the N1–C5 bond in the triazole ring is significantly strained (Fig. 3B, top), so that cycloreversion takes place in each case.

To study these effects more closely, the internal coordinates of the molecules, *i.e.* their bond lengths, bond angles and dihedral angles, were partitioned into three groups. The first group consists of internal coordinates in which the involved atoms are located solely in the ring. Analogously, the second group comprises internal coordinates in which the atoms are located solely in the linker. The third group consists of internal coordinates that involve atoms from both the ring and the linker (*e.g.* the bond between the ring and the linker). The application of this partitioning scheme in the JEDI analysis reveals that the strain energy in the triazole ring of the 1,4-isomer is rather low during the entire end-to-end stretching process and becomes even lower at higher forces (Fig. 3A). This demonstrates that triazoles are mechanically robust in general, and cycloreversion cannot be achieved by a purely mechanical activation of the entire ring *via* a shearing geometry. If the forces pulling at the terminal carbon atoms are low, most strain energy is stored in the interface between ring and linker, but



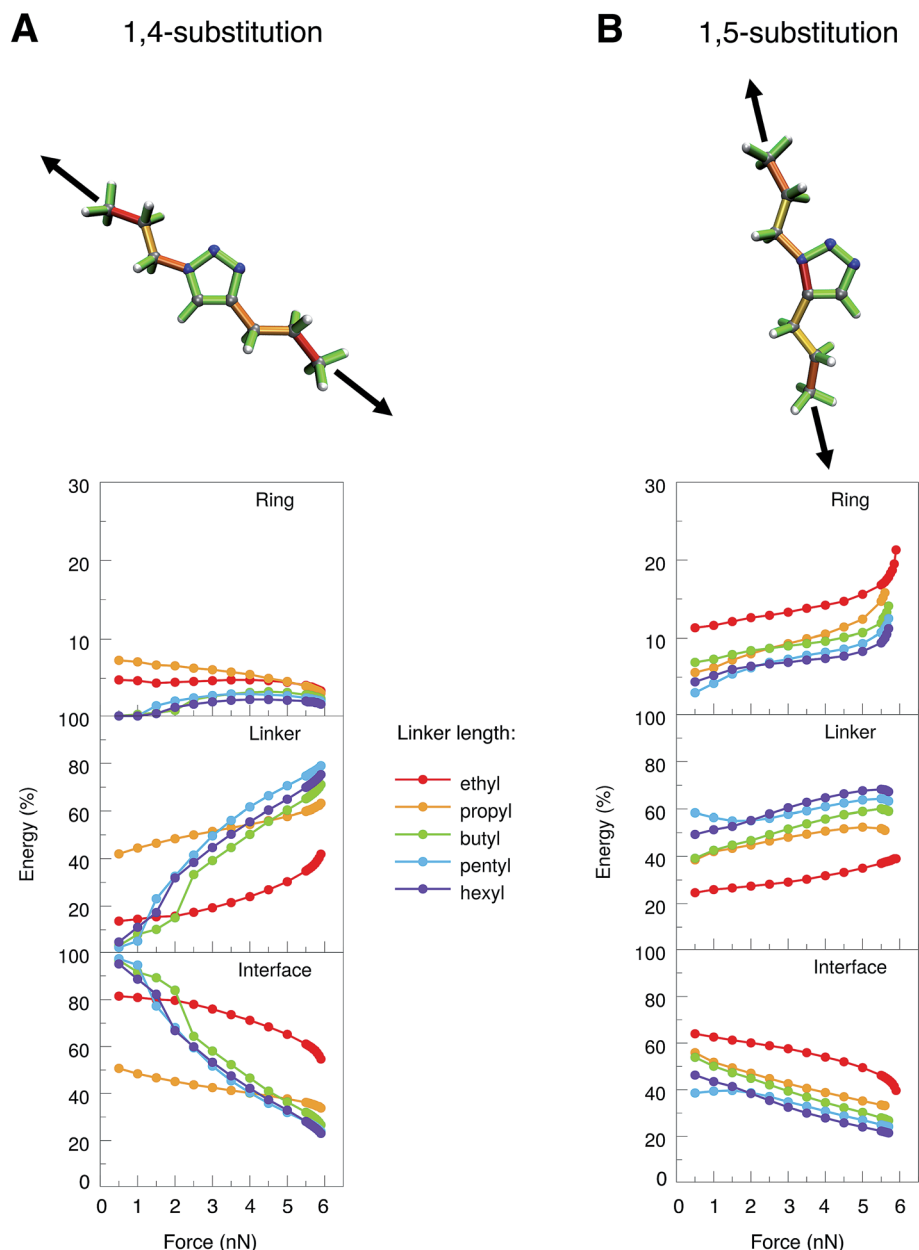


Fig. 3 JEDI analysis of 1,4- (A) and 1,5-substituted (B) 1,2,3-triazoles with linkers of different sizes (ethyl to hexyl) that were stretched end-to-end. Color-coded structures show the results of the JEDI analysis qualitatively for propyl-substituted triazole rings under a constant stretching force of 5 nN. The internal coordinates of the molecules were partitioned into coordinates that are located solely on the ring (top panel of the plots) or solely on the linker (middle panel). The bottom panel shows interface coordinates that include atoms from both the ring and the linker.

this energy percentage decreases with increasing force, whereas the energy percentage in the linkers increases. Bond rupture in the 1,4-substituted triazoles was observed exclusively at the chain ends, which can be rationalized by the observation that the terminal bond angles of an aliphatic chain are typically softer than in its central part, thus leading to better alignment of the terminal bonds with the external force.⁵⁰

Apparently, the end-to-end stretching coordinate in 1,4-substituted triazole is similar to the shearing coordinate c_2 within the triazole ring. Rupture upon stretching the 1,4-isomer end-to-end occurs in the linker and not, as in the case of the shearing coordinate c_2 , within the ring, because very large forces

and energies are needed to initiate rupture in the triazole ring *via* the c_2 shearing coordinate ($F_{\text{max}} = 13.6$ nN). Rupture in the linker, on the other hand, requires much less force and energy ($F_{\text{max}} \approx 6$ nN), this process is thus triggered in the end-to-end stretching coordinate.

In 1,5-substituted triazoles (Fig. 3B), on the other hand, the N1–C5 bond is activated directly. The increase in strain energy in the triazole ring at higher stretching forces can be traced back to the error of the harmonic approximation resulting from rupture of the N1–C5 bond, which leads to cycloreversion. Initially, however, a large amount of strain energy is also stored in the linker and the interface, but this proportion generally



decreases towards the end of the stretching coordinate. Of course, the longer the linker the more energy is stored in it, since there are more internal coordinates that can act as reservoirs of strain energy.

We conclude from these considerations that cycloreversion of 1,5-substituted triazoles is much more feasible than cycloreversion of the 1,4-isomers, since, in the former case, the activation of the entire mechanically robust triazole ring can be circumvented due to the optimal coupling between the end-to-end stretching coordinate and a scissile bond in the triazole ring. Thus, the results of the JEDI analysis provide the quantitative basis for straightforward geometric arguments.²⁶ If cycloreversion of the 1,4-isomers indeed occurs in the experiment, mechanical forces resulting from longitudinal waves in an ultrasound bath cannot play the key role. An earlier study suggested rapid heating due to the collapse of cavitation bubbles, electronically excited states, solvent-assisted cycloreversion or processes including radicals as additional factors influencing scission of 1,2,3-triazole in an ultrasound bath,²⁷ all of which were not considered here. Since this study focuses on the mechanochemical aspect of the cycloreversion of triazoles, we concentrate on the 1,5-substitution pattern in the following.

However, even in 1,5-substituted triazoles it is not possible to activate the N1–C5 bond exclusively by stretching the aliphatic linkers. We found that the bonds connecting the linker to the ring are significantly strained as well. Therefore, two additional stretching coordinates, in which these bonds were pulled apart in an isolated manner (*cf.* the blue and green arrows in Fig. 4A), were investigated. In all cases, linkers of different sizes were considered and the bond rupture forces and energies were

compared. We refrained from an isolated stretching of the terminal carbon–carbon bonds, since the rupture probability of polymer chains in an ultrasound bath is highest within the central 15% of the chain.^{21,22}

In general, the rupture forces change only slightly when the linker length is altered (Fig. 4B, top panel). Interestingly, the scission of one of the interface bonds (N1–C6) requires less force when stretched in an isolated manner than the scission of the N1–C5 bond when stretched end-to-end. Nevertheless, it is the N1–C5 bond in the triazole ring that breaks when the molecule is stretched end-to-end, thereby initiating cycloreversion, since the N1–C5 bond is aligned optimally with the force vector in this stretching scenario. As a result, only rupture events of the N1–C5 bond in the end-to-end stretching coordinate have been observed, regardless of the linker length. Breaking the other interface bond (C5–C7), however, requires significantly more force, which is in agreement with the previous observations that the carbon–nitrogen bonds in the molecule are weaker than the carbon–carbon bonds.

In addition to the comparison of bond rupture forces, it is insightful to compare the energies needed for bond rupture in each of the three coordinates using the JEDI analysis. Since the JEDI analysis is based on the harmonic approximation, the bond rupture energies yielded by the JEDI analysis for the coordinates shown in Fig. 4A overestimate ΔE_{DFT} due to pronounced anharmonic effects. However, a satisfactory correction of the energies is possible.³⁷ In the case of end-to-end stretching, for example, we are primarily interested in the energy stored in the scissile N1–C5 bond. As a step towards correcting this value, the bond length immediately prior to

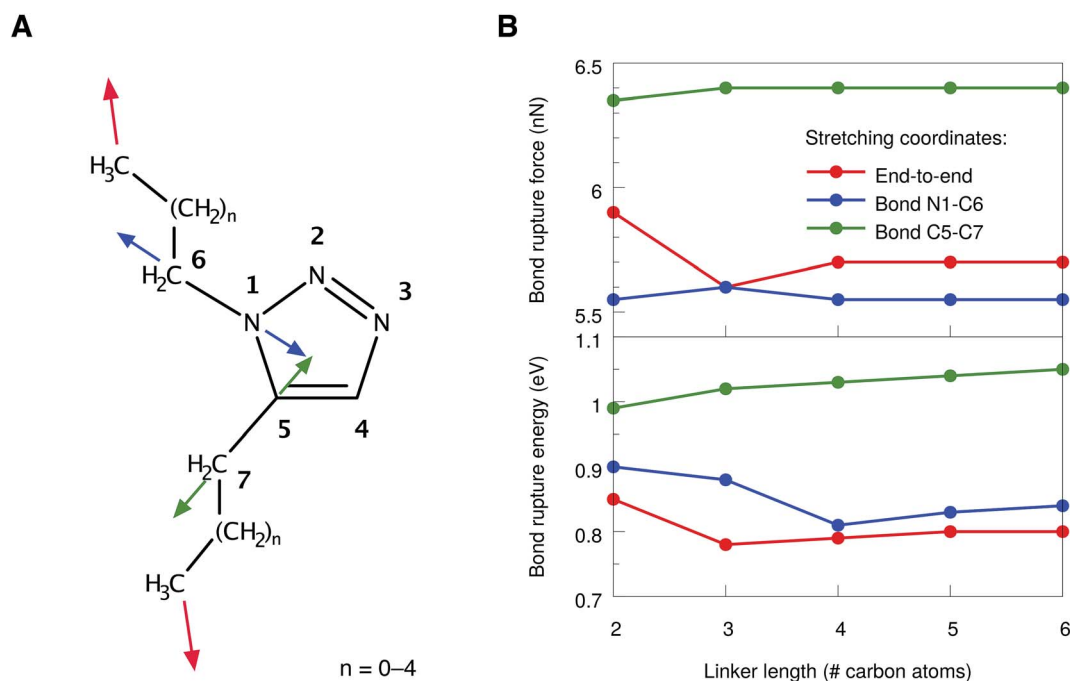
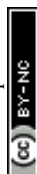


Fig. 4 (A) Numbering scheme and stretching coordinates in the 1,5-substituted triazole ring. Arrows of the same color belong to the same stretching coordinate. (B) Forces (top panel) and energies (bottom panel) required to achieve rupture in the stretching coordinates depicted in (A) for linkers of different sizes. Lines are included to guide the eye.



rupture was adjusted in an isolated bond stretching scenario, which corresponds to the coordinate c_6 . Subsequently, a COGEF (CONstrained Geometries simulate External Force)^{51,52} calculation was carried out. The harmonic error within this isolated stretching setup was used to compensate the error of the energy stored in the N1–C5 bond in the end-to-end stretching scenario. In the case of the interface bonds N1–C6 and C5–C7, the energy correction is much more straightforward, since only one bond is stretched and the overestimation of ΔE_{DFT} by ΔE_{harm} can be used directly to correct the energy.

The resulting corrected energies from the JEDI analysis are shown in Fig. 4B (bottom panel). As expected, similar energies are needed to rupture the bonds N1–C5 and N1–C6. Although the N1–C6 interface bond breaks at slightly lower forces, the energy needed to rupture the N1–C5 bond in the triazole ring and to achieve cycloreversion is lower. A straightforward explanation of this effect is that the energy is proportional to the bond elongation and the N1–C5 bond does not need to be elongated as much as the N1–C6 bond in order to achieve rupture. The C5–C7 bond is again found to be significantly more stable. Interestingly, the rupture forces and energies correlate only weakly with the length of the linker, although the total amount of strain in the molecule increases linearly with the chain length (Fig. S2†).

Hence, from a static point of view, rupture of the bonds N1–C5 and N1–C6 require similar amounts of force and energy. From these considerations we predict that, if cycloreversion takes place in the 1,5-substituted triazole, it is a statistical process that competes with rupture of the adjacent single-bonds connecting the linker to the ring. However, these conclusions were drawn entirely from static considerations and thus we now turn to a discussion of the end-to-end stretching process based on dynamics calculations.

3.2 Dynamic stretching

As dynamics simulations have proven to be valuable tools in the description of mechanochemical bond rupture processes,^{53–55} Born–Oppenheimer Molecular Dynamics (BOMD) simulations have been carried out, in which a stretching force was applied to the terminal carbon atoms of 1,5-hexyl triazole. To mimic experimental conditions, the stretching force was increased by 250 pN each 500 time steps (1 time step = 0.484 fs), until a maximum force of 5 nN was reached. Thus, the total simulation time amounted to 4.84 ns. With this setup, a loading rate of approximately 10^9 nN s^{−1} was simulated, which lies in the range estimated for ultrasound baths (10^8 – 10^{10} nN s^{−1}).²⁸ Since elevated temperatures play a major role in sonochemistry^{56,57} and thermal oscillations have a significant impact on bond rupture in general,^{58–61} we adjusted the temperature in the BOMD simulations to 500 K and sampled the initial velocities *via* a Boltzmann distribution. Due to the size of the system and the immense computational cost, only ten BOMD simulations could be conducted, although a larger number of trajectories would be beneficial for a more robust statistical evaluation. We are aware that our setup does not model the experimental conditions in an ultrasound bath fully realistically, since solvent effects, radicals

and the influence of pressure are not simulated. However, we concentrate on the mechanical component of ultrasound baths and note that a computational protocol for modeling sonochemical experiments realistically is not available.²⁴

The end-to-end distances and the lengths of the bonds N1–C5, N1–C6 and C5–C7 during the BOMD simulations are shown in Fig. 5. In the ten trajectories, eight cycloreversions, initiated by rupture of the N1–C5 bond, and two ruptures of the N1–C6 bond have occurred. Scission of the C5–C7 bond was not observed. The forces needed to achieve bond rupture lie between 3.75 and 4.75 nN. In general, the gradual application of a stretching force leads to a gradual increase in bond lengths. After rupture of a bond, the other bond lengths relax. An example of this effect is provided by the two cases in which rupture of the N1–C6 bond occurs and the N1–C5 bond length relaxes as a consequence (*cf.* the orange and purple lines in Fig. 5B). Although the inclusion of a wider variety of experimental parameters and the calculation of a larger number of trajectories would be beneficial for a quantitative discussion of the behavior of triazoles in ultrasound baths, the results are in qualitative agreement with the static EFEI calculations (Fig. 4B), where it was found that the forces needed for rupture of the bonds N1–C5 and N1–C6 are very similar.

To draw more quantitative conclusions, we used the JEDI analysis to investigate the distribution of potential energy among the internal coordinates of the molecule in the course of the BOMD simulations. For this, we averaged the geometries over 500 consecutive time steps in order to minimize the influence of thermal oscillations and used these mean geometries as the mechanically strained structures within the JEDI analysis.³⁷ The full set of JEDI analyses for all BOMD simulations can be found in the ESI (Fig. S3†). The results of a typical trajectory, in which cycloreversion occurs, are shown in Fig. 6A. The energies in the bonds N1–C5, N1–C6 and C5–C7 are similar throughout a large part of the simulation. As soon as cycloreversion occurs after approximately 4 ns, however, anharmonic effects result in a sharp increase in energy of the N1–C5 bond and the results are no longer quantitative. Up to this point, however, the energies in the bonds N1–C5 and N1–C6 are almost the same, thus supporting our conclusion that rupture of one of these bonds is a statistical phenomenon.

Similar effects can be observed in the JEDI analysis of one of the trajectories in which rupture of the N1–C6 bond occurs (Fig. 6B): the energies in the bonds N1–C5 and N1–C6 are very similar throughout the simulation, but the sharp increase in the N1–C6 bond energy indicates rupture at approximately 4 ns. The observation of both cycloreversion and interface rupture events are in good agreement with the results from the static JEDI analyses based on EFEI calculations (Fig. 4B), where it was found that the energies needed for the rupture of the bonds N1–C5 and N1–C6 are almost the same.

Finally, we note that the forces needed to induce rupture are larger by almost a factor of 2 when they are applied gradually in comparison to a BOMD simulation in which the force is applied suddenly (Fig. S4†). If the force is applied gradually, the molecule has time for equilibration, *i.e.* for a redistribution of strain energy among all internal coordinates. If the force is applied



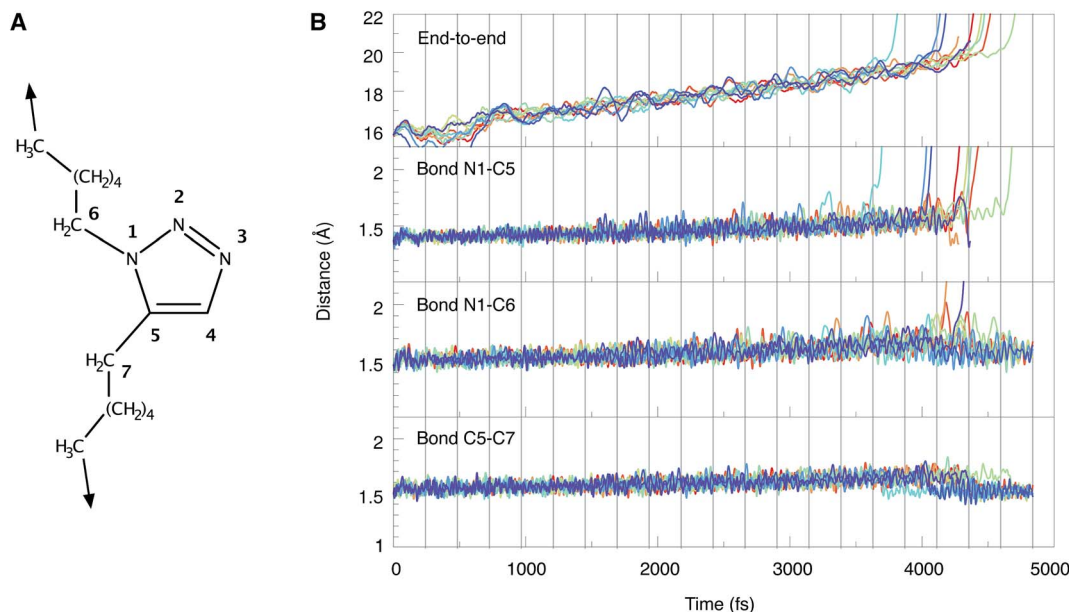


Fig. 5 (A) Numbering scheme and stretching coordinate used in the BOMD trajectories. (B) Temporal progression of the end-to-end distance as well as the lengths of the bonds N1–C5, N1–C6 and C5–C7 during the BOMD simulations. Each line represents one trajectory. Vertical gray lines indicate the points in time when the force was increased by 250 pN.

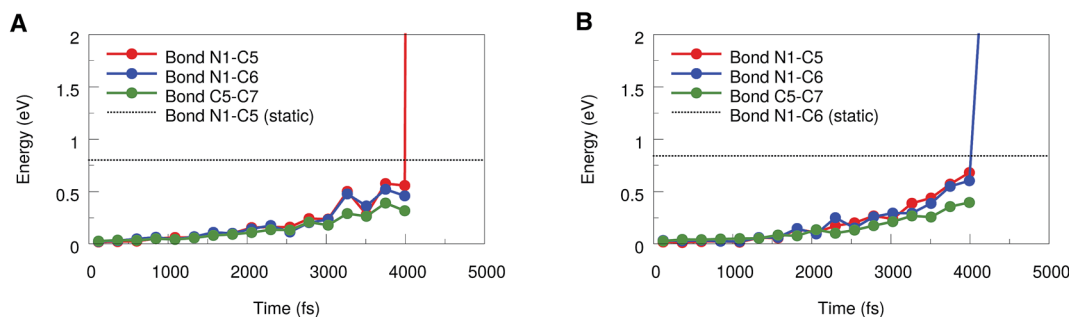


Fig. 6 JEDI analysis of a representative BOMD trajectory in which cycloreversion (A) or rupture of the N1–C6 bond (B) occurs, respectively. The potential energies in the bonds N1–C5, N1–C6 and C5–C7 were calculated based on averaged geometries. The bond rupture energies from static EFEI calculations, which were corrected for anharmonicities, are included as dotted lines.

suddenly, by contrast, the strain propagates rapidly from the linkers to the central triazole moiety, where constructive interference results in efficient bond rupture. Hence, one way of reducing the forces required for rupture in triazoles would be to increase the loading rate. However, the close proximity of the bonds N1–C5 and N1–C6 impedes selective cycloreversion (see ESI†).

4 Conclusions and outlook

In this paper, we have applied static EFEI and dynamic BOMD calculations in combination with the JEDI analysis to answer the question whether mechanically induced cycloreversion of 1,4- and 1,5-substituted 1,2,3-triazoles is generally possible. We have found that cycloreversion of 1,4-substituted 1,2,3-triazoles is impossible if stretching forces are applied to the ends of the aliphatic linkers, since the entire mechanically robust triazole ring would need to be activated. This process is energetically

much less favorable than rupture somewhere in the aliphatic chain. In contrast, mechanical cycloreversion of 1,5-substituted 1,2,3-triazoles is possible, since the N1–C5 bond in the triazole ring, which needs to be ruptured for cycloreversion, can be activated efficiently by stretching forces due to the optimal alignment of the scissile bond with the force vector.

Cycloreversion in 1,5-substituted triazoles, however, competes with rupture of the carbon–nitrogen bond connecting the triazole ring to the aliphatic linker. The forces and energies needed to rupture these bonds are very similar, so that a statistical mixture of cycloreversions and rupture events of the interface bond in stretching experiments is expected. Since only a few BOMD trajectories could be calculated and several important experimental parameters (*e.g.* temperature, pressure, solvent effects, loading rate, length and nature of the linkers) have been neglected or sampled only insufficiently, we cannot predict the distribution of cycloreversion and interface rupture events in experiments quantitatively. In the future, experimental



conditions need to be modeled more realistically and a larger number of *ab initio* molecular dynamics simulations need to be carried out to predict the probability of mechanical triazole cycloreversion in a given experimental setup.

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