Accurate Single-Molecule Kinetic Isotope Effects


ABSTRACT: An accurate single-molecule kinetic isotope effect (sm-KIE) was applied to circumvent the inherent limitation of conventional ensemble KIE by using graphene–molecule–graphene single-molecule junctions. In situ monitoring of the single-molecule reaction trajectories in real time with high temporal resolution has the capability to characterize the deeper information brought by KIE. The C–O bond cleavage and the C–C bond formation of the transition state (TS) were observed in the Claisen rearrangement through the secondary kinetic isotope effect, demonstrating the high detection sensitivity and accuracy of this method. More interestingly, this sm-KIE can be used to determine TS structures under different electric fields, revealing the multidimensional regulation of the TS. The detection and manipulation of the TS provide a broad perspective to understand and optimize chemical reactions and biomimetic progress.

INTRODUCTION

Insights into reaction mechanisms contribute to optimizing synthetic routes and realizing significant cost savings in laboratory preparation and industrial production. In particular, an understanding of the rate-determining step (RDS) is of crucial importance to high reaction yield and efficiency. As an effective method to demonstrate the transition state (TS) configuration and the RDS, KIE has been applied in many reactions for mechanism studies,^1−3 exemplified by studies into the aromatic nucleophilic substitution reaction (12C/13C KIE),^4 the Morita–Baylis–Hillman reaction (primary H/D KIE, involving the cleavage/formation of O–H/C–H bonds)^5 and the Claisen rearrangement (secondary H/D KIE, involving the cleavage/formation of O–C/C–C bonds).^6−15 In general, when the overall reaction rate changes as a consequence of switching isotopes, the isotope substitution site is likely to participate in the RDS to a greater extent (primary isotope effect) or to a lesser extent (secondary isotope effect), thereby helping to determine the main reaction trajectories and corresponding TS structures. However, it is hard to precisely measure KIEs because the KIE values are generally close to 1 (not much difference between the values).^1

As the reaction progresses, the reactants with a slower reaction rate are continuously accumulated. Therefore, when the reaction is nearing completion, the relative concentration of the reactant containing heavy-atom isotopes will increase. The relationship among the ratio of the isotope content in the reclaimed reactant to the isotope content in the reactant at the beginning (R/R_0), the extent of a reaction (F), and the KIE obeys eq 1:1

$$\frac{R}{R_0} = (1 - F)^{1/KIE - 1}$$  \hspace{1cm} (1)

In these experiments, nuclear magnetic resonance spectroscopy (NMR) is used to measure the content of 2H and 13C in the reactant. The KIE of each center can be measured by substituting the isotopic content ratio of each carbon atom or hydrogen atom into the above formula. However, this method is significantly limited by the naturally low abundance of isotopes and the extent of the reaction, resulting in low sensitivity and poor accuracy. In addition, the interpretation of experimental measurements of KIEs requires a known mechanism; otherwise, it may lead to inappropriate conclusions.^16 Therefore, taking advantage of KIEs to realize the visualization of intrinsic reaction pathways is an urgent and key challenge that needs to be solved.

Single-molecule detection (SMD) can shed light on the inherent laws of a reaction,^17 which is expected to be an ideal strategy to break through this bottleneck. SMD presents us with the opportunity to directly visualize the reaction process and possibly noteworthy KIEs, without being limited by abundance. To date, various kinds of SMDs based on molecular optical,^18,19 mechanical,^20,21 and electrical^22,23 properties have flourished for decades. Numerous single-
molecule effects, such as the field-induced effect, the Kondo effect, the Coulomb block, negative differential conductance, and quantum interference, have been discovered and studied. However, few studies of KIEs on single molecules have been reported, due to the limited inherent accuracy of monitoring chemical reactions. In contrast, the platform of graphene–molecule–graphene single-molecule junctions (GMG-SMJ) provides unprecedented accurate measurements of chemical reactions and intermolecular interactions, which lays the foundation to study the KIE effect on single-molecule reactions. In this work, we applied sm-KIE to probe the detailed properties of each intermediate in the Claisen rearrangement reaction and thus discover unexpected reaction behaviors (beyond the traditional textbook understanding) from their intrinsic dynamics.

## RESULTS AND DISCUSSION

### Device Fabrication and Characterization

From the perspective of molecular engineering, phenol-based GMG-SMJ were prepared by covalently integrating a designed molecular bridge with a phenol functional center and amino terminals into nanogapped carboxyl-terminated graphene point electrodes (Figure 1A and Schemes S1 and S2). To confirm the successful formation of the single-molecule device, we compared the current–voltage (I–V) curves before and after the connection of the molecular bridge as shown in Figure S1. About 16 of 169 devices showed the successful connection (∼10% yield) and ∼95% probability of only one-molecule integration between two electrodes could be demonstrated by statistical analysis in the Supporting Information. Furthermore, under a constant bias voltage (300 mV), the bistable switching model between the low and high current levels in recorded I–t curves in an alkaline environment (Figure 1B) originates from the conversion of phenol and phenoxy anion. Those were demonstrated by their corresponding low and high electron transmission probabilities (Figure 1B and S2). This also elucidated the successful connection of the phenol functional molecular bridge. In addition, the bistable switching in alkaline concentration-dependent measurements (Figure S3) showed the preference for the phenoxy anion in an alkali-rich environment, which can be used as a nucleophile to react with allyl bromide to synthesize the substrate for a Claisen rearrangement at the single-molecule level. In advance, a fluorescein-substituted allyl compound, 5000 photos taken by a super-resolution fluorescence microscope with 50 ms exposure are reconstructed to obtain a single-molecule resolution photograph and enlarged image without the background. (D) Synchronous monitoring of current and fluorescent (the mean intensity of fluorescence in the region of the single-molecule functional center) signals.

### Claisen Rearrangement

To monitor a single-molecule Claisen rearrangement, we added allyl bromide to the device under alkaline conditions to produce allyl phenyl ether and monitored the current level at 300 mV bias voltage. An irreversible change in the current shows the S$_2$2 substitution of allyl bromide and successful preparation of allyl phenyl ether (Figure S5). Next, we replaced the solvent with a solution of trifluoroacetic acid (TFA) and further monitored the acid-catalyzed Claisen rearrangement. At the beginning, two conductance states were observed at room temperature, implying the reversible protonation of the allyl phenyl ether reaction substrate (RS) due to the relatively high energy barrier of the Claisen rearrangement (usually heating to 60–200 °C at macroscopic scales). The TFA concentration dependence of the binary conductance switching (Figure S6) provided a support for this assignment. There was a greater amount of the low-conductance state at higher TFA concentrations, illustrating that it should correspond to the protonated state (RS$^{+}$) and the high-conductance state should be assigned to the RS itself. Then, the reaction temperature was raised to 100 °C, and the I–t curves were recorded with six distinct conductance states (Figure 2A). Control experiments performed by replacing the allyl bromide solution with a 1-bromopropane solution are provided in

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**Figure 1.** Device structure and characterization of phenol-based GMG-SMJ. (A) Schematic diagram of a phenol-based single-molecule device. (B) I–t curves of the device in an alkaline environment (1 × 10$^{-5}$ mol/L NaOH solution), the corresponding histogram (right, showing bistable states), and the corresponding transmission spectra of phenol and phenoxy anion (Figure S2). In comparison with the phenol form, for the phenoxy anion form, the perturbed highest occupied molecular orbital (p-HOMO) has a transmission peak with an energy much closer to that of the unbiased electrode chemical potentials ($E_{0}$), leading to the higher conductance. (C) Characterization of a single-molecule connection. During the interaction between a single-molecule phenoxy anion and a fluorescein-substituted allyl compound, 5000 photos taken by a super-resolution fluorescence microscope with 50 ms exposure are reconstructed to obtain a single-molecule resolution photograph and enlarged image without the background. (D) Synchronous monitoring of current and fluorescent (the mean intensity of fluorescence in the region of the single-molecule functional center) signals.
two lower-conductance species, while these two conductance states cannot be transformed into each other (Figure 2B, left), implying that the highest conductance state was an inevitable intermediate during the reaction. The electron transmission spectra and potential energies of the RS, product state (PS), and possible intermediate state (IM) were also studied (Figure 2C, Figure S8, and Figure 2D). The simulated conductances from low to high were RS, PS, and IM (Figure 2C). In addition, the corresponding protonated species were found to have the same order but an overall lower order in comparison to the neutral species (Figure 2C), which is consistent with the previous TFA-dependent measurements (Figure S6) and the high-pass signals (Figure 2B). Similarly, the theoretical studies of the energy profiles showed that the sequence of the potential energy of the neutral species from low to high was PS, RS, and IM (Figure 2D), which matches their dwell times in the low-pass signals (Figure 2B). All of the corresponding protonated species have relatively high energies, demonstrating less probability of occurrence as shown in Figure 2A. On the basis of the filtering method, the different chemical reaction processes were distinguished conveniently.

In view of the slow rate of the Claisen rearrangement process, the number of events exhibited by 57600 Hz is few unless the acquisition time is greatly extended, which will create a large amount of data and make data processing difficult. Fortunately, the splitting of electrical signals can also be approached at the instrument and the Claisen rearrangement can be monitored via a reduced bandwidth (57.6 Hz). Therefore, the Claisen rearrangement can be observed more intuitively and the problem of excessive data volume caused by the high sampling rate can be avoided. The temperature-dependent measurements further prove the availability of the single-molecule platform and the accuracy of determining kinetic and thermodynamic properties, laying the foundation for sm-KIE measurements. The Claisen rearrangement at five different temperatures (353, 358, 363, 368, and 373 K) were recorded (Figure 2E). The ratios of the peak areas in the histograms and the lifetimes (derived from statistics of each time interval) of each state reflect the change in thermodynamics and kinetics with temperature (Figure 2F). The positive correlation between the ratio RS/PS and temperature reveals an overall exothermic reaction ($\Delta H < 0$). More quantitative analysis is provided in Figure S9. In addition, the shorter lifetime of each state at higher temperatures demonstrates the acceleration of each elementary step. Furthermore, the reaction rate constants $k = 1/\tau$ ($\tau$ is the lifetime) were found to have a linear relationship with $1000/T$, meeting the Arrhenius equation. Thus, the activation energy $E_a$ can be derived (Figure S10). A quantitative analysis shows that the single-molecule electrical detection platform has sufficient capacity to study the KIE.

**Single-Molecule KIE.** On the basis of the visualized trajectories and the detected intermediate (IM) discussed above, the 3,3-$\sigma$-migration process (RS to IM), which is slower than the proton shuttle (IM to PS), can be directly determined as an RDS (rather than via ensemble KIE). Further studies of the KIE of the elementary step (3,3-$\sigma$-migration) contribute to a deeper understanding of the mechanism. We synthesized two allyl compounds with different deuterated positions, which reacted with the photox anion through nucleophilic substitution, respectively, to generate Claisen rearrangement substrates: allyl phenyl ethers sp$^2$-D (deuterium connected to sp$^2$ C) and sp$^3$-D (deuterium connected to sp$^3$ C) (Figure 3E). The corresponding protonated species and corresponding dissociation. The conversion rates ($\sim\mu$s$^{-1}$) of the protonation process (Figure S6) also met at the cutoff frequency (1 kHz), which supports this assignment. Then, the other low-pass-filtered signal can be assigned to the Claisen rearrangement at 100 °C. The relatively low reaction rates in the macroscopic experiment also were in accordance with a frequency less than 1 kHz. The species with the highest conductance can be transformed into
3A–C). Under experimental conditions identical with those above, electrical measurements were performed on these

Figure 3. Characterization of sm-KIE. (A–C) I–t curves and corresponding enlarged images of single-molecule nondeuterated (A), sp²-D (B) and sp³-D (C) substrates. The corresponding lifetimes of each state and histograms are provided in the pictures on the left, respectively. The estimated reaction rate constants of each elementary step are provided on the top, respectively. (D) Box plots of the forward and reverse rate constants of 3 kinds of devices based on the statistics of 10 devices.

GMG-SMJIs with three functional centers of nondeuterated (H), sp²-D and sp³-D. In comparison with the nondeuterated reaction rates, the changes in the reaction rate of sp²-D and sp³-D are not noticeable in the inset histograms of the lifetime, \( \tau \). Similarly, the statistics of the ratios of each conductance state vary very little among the three kinds of devices, revealing a slight effect on the kinetics and thermodynamics by isotopic substitution, which is consistent with the ensemble experiments. However, the subtle difference can be shown with the quantitative analysis by a maximum interval likelihood rate estimate. At 373 K, we tested 10 additional devices for each reaction center to obtain the conversion rate constant \( k \) of each elementary step at 300 mV, including the rate constant of the 3,3-\( \sigma \)-migration process in the Claisen rearrangement (RS to IM, abbreviated as \( k_\text{C} \)), as shown in Figure 3A–C and Figures S11–S39. The box plots of the rate constants \( k_{\text{C}} \) of 3,3-\( \sigma \)-migration in the rearrangement process (RS to IM) are provided in Figure 3D. All three kinds of devices revealed obvious differences in both the forward and reverse 3,3-\( \sigma \)-migrations, while there is only a slight difference in the proton shuttle (Figure S40). This also proves the correctness of the conductance state assignment. In addition, obvious solvent KIEs were found in the proton shuttle progress (Figure S41), providing more opportunities for our platform to further detect solvent effects and proton transformation at the single-molecule level (which are not the focus of the current study). According to the definition of the kinetic isotope effect, the KIE values of the forward and reverse reactions of sp²-D and sp³-D are obtained, as shown in Table S1. For the forward reaction, the KIE value of sp³-D is less than 1, while the KIE value of sp²-D is greater than 1, which is consistent with the KIE theory. The resistance to the bending vibration decreases as the carbon atom undergoes a change from sp³ to sp² hybridization, which has a greater influence on the C–H bond than on the C–D bond, leading to the difference in the activation energy of the reaction. Therefore, the reaction rate decreases when the C–H bond is replaced by the C–D bond \( (k_{\text{fi}}/k_{\text{fi-D}} > 1) \). Conversely, when a carbon atom undergoes a change from sp³ to sp² hybridization, the reaction rate increases \( (k_{\text{fi}}/k_{\text{fi-D}} < 1) \). Specifically, the sm-KIE values at a general bias voltage (300 mV) and 373 K are \( k_{\text{fi}}/k_{\text{fi-D}} = 0.798 \pm 0.004 \) and \( k_{\text{fi}}/k_{\text{fi-D}} = 1.290 \pm 0.060 \), which are more obvious than the results \( (k_{\text{fi}}/k_{\text{fi-D}} = 0.950 \pm 0.020 \) and \( k_{\text{fi}}/k_{\text{fi-D}} = 1.180 \pm 0.020 \) in previous work on the aromatic Claisen rearrangement at 443–468 K.\(^{13}\) The more obvious KIE at the single-molecule level can be attributed to three main reasons. (1) Due to the catalysis by an acid, we conducted the single-molecule Claisen rearrangement at a relatively low temperature (373 K). According to \( \frac{k_{\text{fi}}}{k_{\text{fi-D}}} = \frac{H}{D} \) showing an inverse proportion to some extent,\(^{42}\) the lower temperature leads to a larger KIE. (2) Without the influence of the natural abundance and the difference in the reaction extent between H- and D-substituted substrates, the pure single (isotope substituted) molecule shows a more obvious KIE. (3) The focused single-molecule phenol moiety was necessarily substituted at the 2- and 5-positions to span the gap. The KIE of the Claisen rearrangement was reported to have a large substitution effect,\(^{4} \) which results in the difference between ensemble and single-molecule measurements.

**Regulation of the Transition State.** According to KIE = EIE' (EIE is the equilibrium isotope effect, \( E = \frac{k_{\text{fi}}}{k_{\text{fi-D}}} \) and \( i \) is the breakage or formation degree on the corresponding deuterated carbon),\(^{43} \) \( i \) can be derived from the simple formulas

\[
EIE = \frac{k_{\text{fi}}}{k_{\text{fi-D}}} = \frac{k_{\text{fi}}}{k_{\text{fi-D}}} = \frac{H}{D} = KIE_{\text{EIE}} \quad \text{KIE}
\]

\[
i = \frac{\ln KIE_{\text{E}}}{\ln KIE_{\text{E}} - \ln KIE_{\text{E}}}
\]

For sp²-D, \( i \approx 27\% \), which means that the C–C bond formation degree is \( \sim 27\% \). For sp³-D, the degree of C–O bond cleavage is \( \sim 57\% \) (the calculation process of \( i \) is provided in the Supporting Information). These reveal the structure of the TSs: the cleavage degree of the C–O bond is greater than the formation of the C–C bond, which proves the concerted but asynchronous mechanism of 3,3-\( \sigma \)-migration. Note that the derived \( i \) represents not only the C–C and C–O bond stretching but also the variations of the vibrational frequencies in the reaction coordinate. Specifically, the C–H vibrational frequencies (such as the transition of the out-of-plane bending vibration between sp² and sp³ carbons) resulting from the change in the type of carbon hybridization during the reaction can be reflected in the observed KIE. The derived \( i \) implies the change degree of the labeled C–H bond vibration frequency from the substrate to the TS. A higher value was found on the sp³ carbon, showing more change in this C–H vibrational frequency and the type of the hybridization, which also implies a greater degree of the C–O bond breaking in comparison to the C–C bond formation. Moreover, the sum of bond breakage and bond formation is less than 1, indicating that the transition state is more like the substrate than the product and the reaction experiences an early transition state. To better describe the multidimensional view of the Claisen rearrange-
ment, a More O’Ferrall–Jencks diagram is provided in Figure S42, also revealing the concerted but asynchronous mechanism.

Due to the relatively high polarity of the TS in the 3,3-σ-migration, water was found to accelerate the reaction. In the current case, the natural external electric field (EEF) in the single-molecule junction is considered to regulate the apparent reaction rates and TS structures. To identify the TS structure, \(I-t\) curves of three kinds of devices were recorded under different bias voltages (Figures S11–S39) and the corresponding rate constants and KIEs are provided in Tables S1 and S2. A set of \(I-t\) curves of the nondeuterated device is provided in Figure 4A. All of the forward reaction rate estimates show a nonmonotonic relationship between \(k_{r,S}\) and the bias voltage (Figure 4B), implying a regulatory effect on the Claisen rearrangement. However, only one conductance (RS) was recorded when we switched the bias voltage direction, revealing no reaction of the substrate (Figure S43). The benzene ring was fixed and optimized under the electric field parallel to the molecular bridge (Z axis in Figure 4C). The spatial orientation of the allyl group connected by the C–O–C bond varies under different electric fields. The entire allyl group shifts away from the reaction site (the distance is marked as an orange dashed line) and is unable to form the six-membered ring TS required for rearrangement under a negative bias (forming an electric field antiparallel to the Z axis), thus resulting in a no-reaction state. On the other hand, the structural change of the allyl group was restricted to the neighboring reaction site under the electric field parallel to the Z axis, which makes it easier to meet the requirement of rearrangement. In addition, the EEF plays a role in stabilizing the polar TS and further accelerates the reaction, as demonstrated by the positive correlation between \(k_{r,S}\) and voltages at a low bias range (100–300 mV) in Figure 4B. However, when the EEF was further increased, the allyl groups were more strictly aligned in the direction of the electric field, and the structural change became difficult. Under these conditions, the orientation of the allyl group has changed to be in plane with a larger distance from the reaction site (Figure 4C, left), causing a certain deviation between its position and the TS as well as a relatively higher energy barrier. This is consistent with the negative correlation between \(k_{r,S}\) and voltages at a high bias range (400–600 mV). The bias-voltage-dependent measurements reveal the regulation of the apparent reaction rates and both the forward and reverse catalysis by EEFs.

Furthermore, the bond length of the TS under different EEFs obtained from the theoretical studies provided evidence of the regulation of the TS structure via EEFs. With an increase in the applied electric field, the C–C bond’s distance in the transition state gradually decreases, and the C–O bond’s distance increases (Figure 4D). The computational results (the bond lengths and the atomic coordinates of TS) are provided in Tables S3–S8. The theoretical simulation has accuracy limitations, but the ultrasensitive single-molecule experiments provide a more accurate description. The breakage (C–O bond) or formation (C–C bond) degrees of the TS under different bias voltages were derived from the KIE of two kinds of deuterated devices (Figure 4E) As the bias voltage increases, the degree of the C–O bond breakage and the degree of the C–C bond formation gradually increase, indicating that the TS shifts to the right (the structure of the early TS is more similar to that of the product). At the same time, the sum of the degrees of both bond breakage and bond formation is less than 1, showing that the transition state is still more like a reactant than the product, and the reaction experiences an early transition state. The TS structure shows its position change along the reaction coordinate under different EEFs. In combination with the influence of the EEF on the height of the TS, we found the diversification of the EEF regulation: the molecular configuration, the energy barrier (orthogonal direction), and the TS position (horizontal direction). From another perspective, this sm-KIE approach offers a multidimensional description of the TS, the Holy Grail of chemical research, which forms the foundation of direct detection of the TS.

**CONCLUSION**

In summary, we demonstrate sm-KIE as a distinctive method to reveal the intrinsic mechanism of chemical reactions. The filtering technology has been creatively used in distinguishing electrical signals, which allows us to analyze different reaction processes in one reaction data set and simultaneously determine the KIE of forward and reverse reactions. In addition to quantitative analyses of thermodynamics and kinetics of sm-KIE, different EEFs are also found to not only control the occurrence of the reaction but also realize the precise regulation of the TS structures. We firmly believe that,
with the advantages of high detection sensitivity and accuracy, this sm-KIE approach is ready to be applied immediately to a variety of single-molecule dynamics studies, which is crucial to the development of fundamental chemical reactions and an understanding of basic life sciences.

ASSOCIATED CONTENT

Supporting Information
The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/jacs.1c12490.

Molecular synthesis, fabrication procedure of the graphene FET array and single-molecule device, single-molecule connection analysis, $I−V$ curves of single-molecule devices, transmission spectra of phenol and phenoxy anion, pretreatment of graphene--phenol--graphene junctions, photoelectrical integration, preparation of the substrate, protonation of the substrate, control experiment for single-molecule devices, the transmission spectra of six species, plots of thermodynamic and kinetic properties, additional $I−t$ curves of three kinds of devices, the summary of the rate constants and sm-KIEs, rate constants of the proton shuttle, the calculation process of $i$, a More O’Ferrall–Jencks diagram, $I−t$ curves under negative bias voltages, the bond lengths and the atomic coordinates of the TS, and the theoretical simulation methods (PDF)

Fluorescent intensity and current signals (MP4)

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Notes
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REFERENCES


