

Ultrafast formation dynamics of D_3^+ from bimolecular reaction of the D_2 - D_2 dimer

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Bimolecular reactions involving small diatomic molecules constitute a fundamental category of chemical processes. Recent experimental breakthroughs have enabled precise timing of the ultrafast formation dynamics of D_3^+ originating from the D_2 - D_2 dimer. In this paper, we systematically survey diverse theoretical approaches for numerically simulating bimolecular reaction dynamics, considering various initial geometric configurations of the dimer. Our findings demonstrate the consistency and reliability of both classical and quantum methods in predicting the reaction timescale for D_3^+ formation. Additionally, we illustrate the intricate microscopic details of bimolecular reactions, laying the foundation for comprehending intermolecular interactions.

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I. INTRODUCTION

Bimolecular reactions represent a class of fundamental processes in chemistry, encapsulating intricate molecular dynamics phenomena. Among these, the formation of the trihydrogen cation H_3^+/D_3^+ from the hydrogen dimer H_2 - H_2/D_2 - D_2 represents a basic while pivotal instance. The significance of this reaction extends beyond its unique bonding properties and versatility. As a proton donor, the resulting H_3^+ serves as the initiator for the majority of chemical reactions within interstellar clouds, a phenomenon that has attracted substantial attention in the past two decades [1–12]. Previous investigations have revealed the origin of trihydrogen cation formation within interstellar molecular clouds, attributing it to the cosmic-ray ionization of hydrogen molecules [13]. The subsequent reaction ($H_2 + H_2^+ \rightarrow H_3^+ + H$) is widely regarded as the principal pathway leading to H_3^+ formation, with extensive experimental research elucidating kinetic energy and angular distribution of products, reaction rate coefficients, and state-selective cross sections [14–16].

Recent experimental advances have allowed for precise temporal characterization of the formation dynamics of trihydrogen cations, pinpointing a reaction timescale on the order of hundreds of femtoseconds [17,18]. This achievement was facilitated by starting from the initial configuration of a dimer state, providing a known initial separation and, consequently, a well-defined spatial starting point. The temporal starting point, essential for timing the bimolecular reaction, was enabled by employing a femtosecond pump-probe setup. Notably, these studies employed classical trajectories to

describe nuclear motion, similar to the approach used in most prior molecular dynamics calculations. A crucial question thus remains unanswered: does the timescale for ultrafast formation dynamics differ when accounting for the quantum nature of the bimolecular reaction?

Chemical reactions inherently involve the cleavage and formation of chemical bonds. In systems with a high degree of freedom, the reaction dynamics can become intricate, often obscuring the underlying physical processes. In such scenarios, one would typically focus on the principal coordinates relevant to the reaction, often termed the reaction coordinates, which clarify and simplify the understanding and tracking of reaction dynamics. After defining these reaction coordinates, potential energy surfaces can be constructed through first-principle quantum chemical calculations. Subsequent nuclear dynamical evolution occurs on these potential energy surfaces and can be examined using various theoretical methods. In addition to employing classical trajectories, the nuclear wave packet can be propagated on the potential energy surface (PES) by numerically solving the time-dependent Schrödinger equation (TDSE), offering a quantum perspective on reaction dynamics. Effective information contained within the time-dependent nuclear wave packet can be extracted using approaches such as the virtual detector method [19–22], Bohmian mechanics (BM) [23–30], and backpropagation (BP) [31–34].

In this article, we focus on the elementary ultrafast reaction dynamics of D_3^+ formation from the D_2 - D_2 dimer, aiming to explore nuclear motion from both quantum and classical perspectives. We systematically compare different theoretical methods for achieving time resolution in ultrafast bimolecular reaction dynamics. Our results reveal that, when properly implemented, these methods yield similar timescales for the chemical reaction, thus affirming their reliability. It also demonstrates the limited influence of quantum effects on nuclear motion. In addition, we systematically explore the

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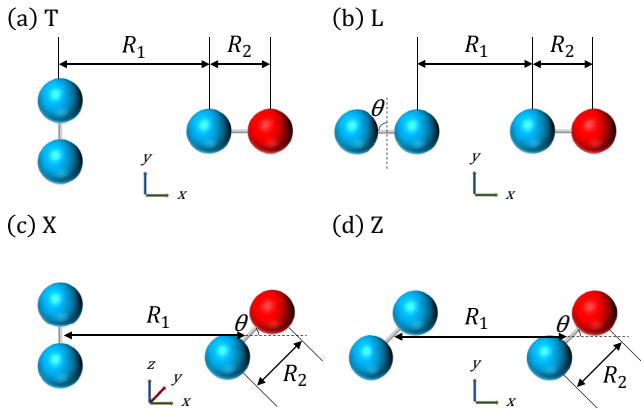


FIG. 1. Different initial configurations of the D_2 - D_2 dimer.

ultrafast reaction dynamics starting from different initial geometric configurations of the neutral D_2 - D_2 dimer. The intricate microscopic details of bimolecular reactions are important for further understanding intermolecular interactions.

The article is organized as follows. In Sec. II, we summarize the initial stable geometric configurations of the neutral D_2 - D_2 dimer, elucidate the ultrafast process of the bimolecular reaction, and detail the theoretical methods for the timing of the reaction dynamics. In Sec. III, we comparatively investigate the role of quantum effects of the nuclei using a variety of theoretical methods starting from the most stable and populated initial configuration. In Sec. IV, we explore the influence of the initial nuclear momentum on the subsequent molecular dynamics. In Sec. V, we systematically study the details of ultrafast reaction dynamics starting from different initial geometric configurations. Conclusions are given in Sec. VI. Atomic units are used throughout unless stated otherwise.

II. THEORETICAL FRAMEWORK

A. Initial configurations

The D_2 - D_2 dimer complex, comprising four atoms, can be fully characterized by a six-dimensional potential energy surface. This surface exhibits a number of local minima, each supporting a stable local geometric configuration. These stable points correspond to the initial configurations of the neutral dimer, a crucial factor enabling precise time resolution of the ensuing reaction dynamics by providing a well-defined spatial starting point.

Previous investigations have revealed several stable initial structural configurations of the neutral dimer, which encompass the T, L, X, and Z shapes [35], as visually depicted in Fig. 1. For each of these geometric configurations, we establish an approaching coordinate denoted as R_1 and a departing coordinate represented by R_2 . As the reaction progresses, the approaching distance R_1 gradually decreases, culminating in the formation of trihydrogen, while the departing distance R_2 concurrently increases, leading to the departure of the remaining hydrogen atom, indicated in red. In the case of the most stable T configuration, the definition of R_1 and R_2 as the reaction coordinates is sufficient, as they are the principal coordinates directly relevant to the reaction. However,

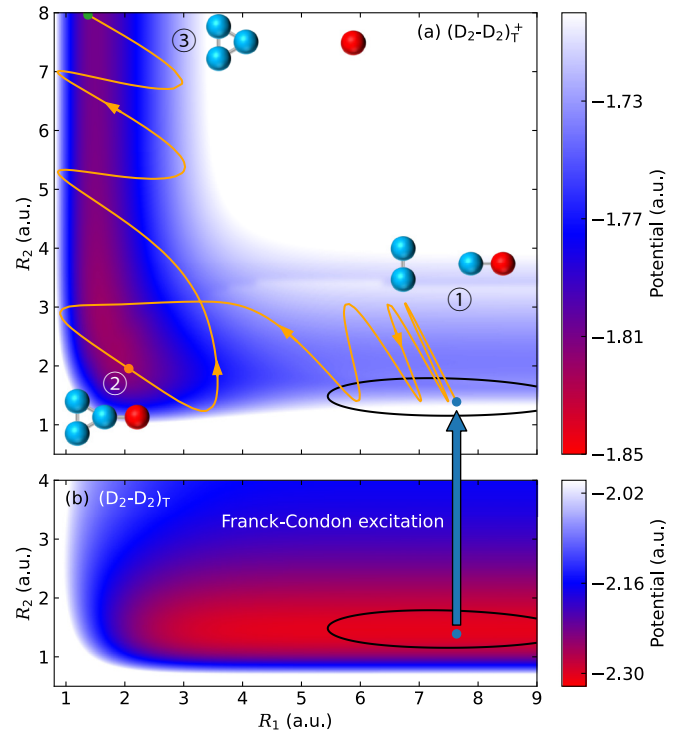


FIG. 2. Potential energy surfaces of the T-type D_2 - D_2 dimer complex in both its (a) cationic and (b) neutral states, with the ground-state probability distribution highlighted by black contour lines. In panel (a), a representative reaction trajectory leading to the formation of D_3^+ is depicted by the orange curve. The dimer configurations at various points along this trajectory are displayed as insets.

for other initial configurations, an additional rotation angle, θ , must be introduced to enable the formation of the stable, triangle-shaped D_3^+ .

B. Reaction mechanism

To explore the quantum effects on reaction dynamics and gain insights into the application of different quantum methods in molecular dynamics simulations, we conduct a comparative analysis between classical and quantum methods to calculate the timescale of trihydrogen cation formation. Considering the various structural configurations of this reaction, we initiate our investigation with the most stable T-type configuration. As illustrated in Fig. 1(a), the T-type configuration features two molecular axes perpendicular to each other; this configuration has the lowest equilibrium energy and represents the most probable configuration. As shown in Fig. 2, by defining two reaction coordinates, R_1 and R_2 , the PES of the cationic $(D_2-D_2)^+$ dimer and the neutral D_2 - D_2 dimer are calculated using the complete-active-space self-consistent field (CASSCF) method at the CAS(3,4) and CAS(4,4) levels with the aug-cc-pVTZ basis set. The active space includes two occupied and two unoccupied orbitals and all electrons are active, which is sufficient for the calculations of the D_2 - D_2 dimer. The initial equilibrium geometry of the neutral dimer is determined at $R_1 = 7.635$ a.u. and $R_2 = 1.390$ a.u.

The formation of D_3^+ can proceed via three steps, as illustrated in Fig. 2. Initially, upon Franck-Condon excitation, the cationic state populates around the equilibrium geometry of the neutral state (referred to as step ①). Under the force field of the cationic state, the nuclear dynamics follow a typical trajectory indicated by the orange curve in Fig. 2(a). Notably, as the hydrogen molecule at the right end undergoes single ionization, the D_2^+ ion begins to vibrate around the new equilibrium internuclear distance of $R_2 = 2.0$ a.u. while simultaneously approaching the neutral D_2 molecule at the left end. This results in an intermediate state where all four particles become adjacent (step ②). Finally, the departure of the neutral D atom at a longer time delay signifies the formation of the D_3^+ cation (step ③).

C. Simulation methods

In this section, we provide an overview of the main computational methods employed in our study, including both classical and quantum approaches. Our benchmark begins with classical-trajectory Monte Carlo (CTMC) simulations, which are then compared to the outcomes of TDSE, BM, and BP.

1. Classical-trajectory Monte Carlo

The Born-Oppenheimer approximation separates the motion of atomic nuclei and the calculation of electron structure, streamlining the analysis of nuclear motion. CTMC, as a powerful implementation of continuum propagation, neglects quantum effects but fully accounts for molecular potential energy during nuclear motion. Setting the initial conditions of trajectories, involving both initial position and velocity, constitutes the initial crucial step in CTMC. In our simulations, the initial positions are sampled from the ground-state nuclear wave function on the neutral dimer state (depicted by black contour lines in Fig. 2). This approach naturally incorporates anharmonic effects compared to traditional methods. The ground-state nuclear wave function is derived from imaginary-time propagation by solving the TDSE, as elaborated below.

Once the initial conditions are set, the trajectories are evolved according to the Newtonian equation of motion:

$$m\ddot{\mathbf{r}} = -\nabla V(\mathbf{r}), \quad (1)$$

where $V(\mathbf{r})$ represents the PES of the cationic $(D_2-D_2)^+$ dimer and m is the generalized mass, which depends on the specific configuration and coordinate system where it is defined. Within the R_1 - R_2 coordinate system of the T-type configuration as in Fig. 1(a), m can be written as

$$m = \begin{bmatrix} m_D & m_D/2 \\ m_D/2 & 3m_D/4 \end{bmatrix}, \quad (2)$$

where m_D is the mass of the D atom. To ensure sufficient statistical accuracy, a swarm of 10 000 nuclear trajectories is evolved on the PES with zero initial velocity. Once all trajectories have completed their propagation, we extract time-related information. We consider D_3^+ formed when the approaching distance R_1 is smaller than 4 a.u. and the departing distance R_2 is larger than 5 a.u. Subsequently, trajectories leading to the formation of D_3^+ are collected, allowing us to

construct a histogram of the formation time of these trajectories for detailed analysis of the reaction timescale.

2. Time-dependent Schrödinger equation

As a well-developed quantum theory, TDSE offers a comprehensive quantum perspective for studying molecular dynamics. In the current work, the TDSE is solved within the framework of nonrelativistic quantum mechanics to describe the motion of the nuclear wave packet governed by

$$i\frac{\partial}{\partial t}\psi(\mathbf{r}, t) = \left[\frac{\mathbf{p}^2}{2m} + V(\mathbf{r}) \right] \psi(\mathbf{r}, t). \quad (3)$$

The TDSE is propagated using the split-operator Fourier method. The initial nuclear wave function is obtained as the ground state of the neutral dimer, which is calculated through imaginary-time propagation by setting $V(\mathbf{r})$ as the PES of the neutral state.

Following Franck-Condon excitation, the nuclear wave function undergoes quantum dynamical evolution on the PES of the cationic state. To this end, we populate the initial nuclear wave function on the PES of the cationic $(D_2-D_2)^+$ dimer and allow the nuclear wave function to evolve freely.

The PES of the T-type D_2 - D_2 dimer has been calculated within the scope of $0.5 < R_1 < 13.8$ and $0.5 < R_2 < 10.5$, covering the essential region where molecular dynamics occurs, and the time evolution of the nuclear wave function has been correspondingly carried out within this range. We have used a time step of $\Delta t = 1$ and a spatial step of $\Delta R_1 = \Delta R_2 = 0.013$, and convergence has been achieved with this grid parameter. Meanwhile, an absorption boundary is employed to prevent reflection of the nuclear wave packet from the grid border.

In order to extract the time-related information of this reaction, we employ the virtual detector method [19–22] to obtain the probability flux of the wave function at the position defining the formation of D_3^+ : $R_1 < 4$ a.u. and $R_2 = 5$ a.u. where the virtual detectors are placed right on the grid points. The probability flux at the position \mathbf{r}_d of a specific virtual detector can be calculated as

$$\mathbf{j}(\mathbf{r}_d, t) = -\frac{i}{2m} [\psi^*(\mathbf{r}_d, t)\nabla\psi(\mathbf{r}_d, t) - \text{c.c.}]. \quad (4)$$

Upon accumulating all detector signals, the distribution of the probability flux (projected onto the direction of the departing coordinate R_2) over time can be analyzed to determine the reaction time of the formation process.

3. Bohmian mechanics

Originally proposed by Louis de Broglie and later elaborated upon by David Bohm, BM serves as a complementary yet fully equivalent quantum approach to the TDSE, describing quantum evolution in terms of trajectorylike language [30]. In this approach, the nuclear wave function can be expressed in polar form as

$$\psi(\mathbf{r}, t) = R(\mathbf{r}, t)e^{iS(\mathbf{r}, t)}, \quad (5)$$

where $R(\mathbf{r}, t)$ represents the amplitude and $S(\mathbf{r}, t)$ denotes the phase of the nuclear wave function. Inserting this polar

form [Eq. (5)] into the TDSE [Eq. (3)] results in two coupled differential equations:

$$\frac{\partial S}{\partial t} + \frac{(\nabla S)^2}{2m} + V(\mathbf{r}) + Q(\mathbf{r}, t) = 0, \quad (6)$$

$$\frac{\partial \rho}{\partial t} + \nabla \cdot \left(\frac{\rho \mathbf{p}}{m} \right) = 0, \quad (7)$$

where $\rho(\mathbf{r}, t) = R^2(\mathbf{r}, t) = |\psi(\mathbf{r}, t)|^2$ is the probability density and $\mathbf{p} = \nabla S(\mathbf{r}, t)$ is the momentum of the nucleus. Equation (7) is the continuity equation, while Eq. (6) is referred to as the quantum Hamilton-Jacobi equation in BM.

Notably, in comparison to the classical Hamilton-Jacobi equation, Eq. (6) incorporates an additional term:

$$Q(\mathbf{r}, t) = -\frac{1}{2} \frac{\nabla^2 R(\mathbf{r}, t)}{R(\mathbf{r}, t)}, \quad (8)$$

known as the quantum potential in BM. This term introduces nonlocality, thereby incorporating quantum features distinct from classical trajectories.

Based on Eqs. (6) and (7), a Bohm-Newton equation of motion can be derived:

$$m\ddot{\mathbf{r}} = -\nabla[V(\mathbf{r}) + Q(\mathbf{r}, t)], \quad (9)$$

whose direct solution is, however, difficult. Instead, one usually resort to a much simpler equation of motion:

$$m\dot{\mathbf{r}} = \nabla S(\mathbf{r}, t), \quad (10)$$

where $S(\mathbf{r}, t) = \arg \psi(\mathbf{r}, t)$ is the time-dependent phase of the wave function obtained by solving the TDSE. Therefore, the Bohmian trajectories are guided by the pilot wave $\psi(\mathbf{r}, t)$ calculated by the TDSE. Finally, similar to the case of CTMC, time-related information can be extracted using this approach.

4. Backpropagation

The backpropagation method [31–34] involves a full quantum propagation of the nuclear wave function forward in time, followed by a classical backpropagation of the target portion of the wave function to pinpoint the formation time of D_3^+ . During the forward phase, akin to the TDSE method, virtual detectors are positioned at $R_1 < 4$ a.u. and $R_2 = 7$ a.u., where all quantum flux is outward, representing the eventual formation of D_3^+ . This portion of the wave function is then transformed into classical trajectories and propagated backward in time to extract their temporal information. The local momentum required to initiate backpropagation is computed by substituting Eq. (5) into Eq. (4):

$$\mathbf{p}(\mathbf{r}_d, t) \equiv \nabla S(\mathbf{r}_d, t) = \frac{m\mathbf{j}(\mathbf{r}_d, t)}{\rho(\mathbf{r}_d, t)}. \quad (11)$$

Subsequently, the resulting phase-space density $\rho(\mathbf{r}, t)$ is propagated backward in time with classical trajectories $[\mathbf{p}(t), \mathbf{r}(t)]$ until $R_2 = 5$ a.u. The resulting distribution of the formation times of D_3^+ is then compiled.

III. ULTRAFAST REACTION DYNAMICS

To investigate quantum effects in chemical reactions and molecular dynamics, we present in Fig. 3 the normalized distribution of the formation rate of D_3^+ starting from the T-type initial configuration of the neutral dimer, calculated by different theoretical methods: CTMC [panel (a)], BM [panel

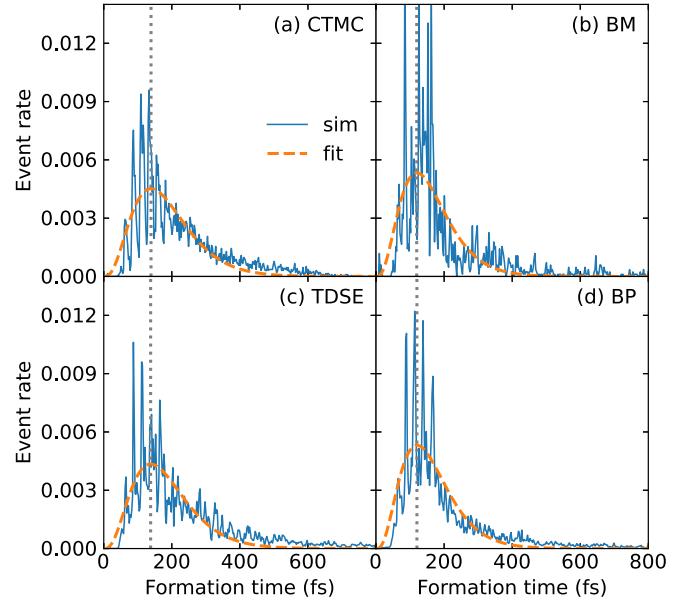


FIG. 3. Histogram of the D_3^+ formation time calculated by (a) classical-trajectory Monte Carlo (CTMC), (b) Bohmian trajectory (BM), (c) time-dependent Schrödinger equation (TDSE), and (d) backpropagation (BP) methods.

(b)], TDSE [panel (c)], and BP [panel (d)]. While CTMC is a purely classical method, BM and TDSE are purely quantum methods, and BP is a hybrid quantum-classical method. The distributions of the formation time obtained from these methods are subsequently fitted with the Gamma distribution given by

$$f(t) = \alpha t^{k-1} e^{-t/\tau}, \quad (12)$$

which is frequently used to model time distributions in Poisson processes, where α is an overall scaling factor, $k = 4$ is the shape parameter for the T-type configuration, and τ is the scale parameter. The gray dotted lines in each panel in Fig. 3 indicate the peak position of the fitted curves, representing the peak reaction time.

In Fig. 3, the peak formation time obtained using CTMC is 138.8 fs, while the peak formation time is 119.2 fs as obtained from BM. Comparing Eqs. (1) and (9), we note that the difference between the two equations lies in the presence of the quantum potential $Q(\mathbf{r}, t)$. Therefore, the small disparity in the reaction time calculated by CTMC and BM methods can be attributed to the quantum effects introduced by the quantum potential $Q(\mathbf{r}, t)$.

Results from both BM and TDSE methods are based on the quantum wave function. Compared to BM, the formation time extracted from TDSE is a little longer, at 137.7 fs. As mentioned earlier, the D_2^+ ion approaches the D_2 molecule while vibrating around the new equilibrium internuclear distance. Until D_3^+ is formed with the departure of the neutral D atom, an oscillation in the distance between the nuclei of the D_2^+ ion persists, implying that, at certain time instances, the distance R_2 may get close to or even exceed 5 a.u. during the oscillation before the formation of D_3^+ . Leveraging the trajectorylike language in the BM method allows us to directly determine the formation time by observing the last

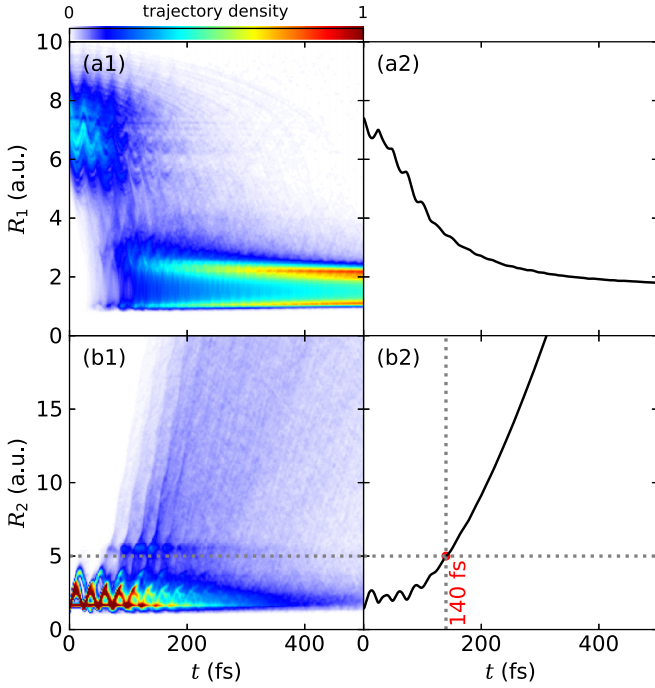


FIG. 4. Ultrafast reaction dynamics of the T-shape configuration. Column (1) displays the classical trajectories leading to the formation of D_3^+ . Column (2) indicates the average value of these trajectories on the R_1 and R_2 coordinates at each moment.

instance when Bohmian trajectories cross the $R_2 = 5$ a.u. boundary. In contrast, in TDSE, wave packets approaching $R_2 = 5$ a.u. before the formation of D_3^+ are indiscriminately detected by virtual detectors, leading to additional signals and consequently distinct peak formation times compared to the BM method.

BP provides an alternative solution to this issue. The BP method comprises two stages: forward evolution of the nuclear wave packet and the backpropagation of classical trajectories. The forward evolution aligns with the TDSE method, with the only difference being the placement of virtual detectors further away to ensure the complete formation of D_3^+ . Consequently, the corresponding reaction has already concluded before detection, significantly reducing interference between the signals. When the virtual detectors located at $R_2 = 7$ a.u. detect a signal, it is converted into a classical trajectory and backpropagated. The first instance when the particle reverses its path and reaches $R_2 = 5$ a.u. is considered the D_3^+ formation time of this trajectory. Using this approach, the reaction time obtained by the BP method is determined to be 120.1 fs, very close to the results from BM. This demonstrates the existence of quantum effects, albeit small, on the motion of atomic nuclei.

Considering the significantly longer computation times of BM and BP methods compared to TDSE, TDSE remains a reasonable choice for quantum computations within an acceptable margin of error. For highly accurate studies of quantum effects, however, BM and BP methods offer superior precision.

The distributions of formation rates obtained from all methods exhibit a common feature: multiple peaks can be

observed in the early stage of the histogram. This observation could be hypothesized to be related to the vibration of D_2^+ upon photoionization of D_2 [36]. To substantiate this conjecture, the classical trajectories leading to the formation of D_3^+ are plotted in Figs. 4(a1) and 4(b1). As depicted in Fig. 4(b1), D_2^+ , formed via single ionization of D_2 , vibrates while approaching D_2 . Whenever the average internuclear distance of D_2^+ reaches a local maximum, the probability of its dissociation to form D_3^+ also peaks subsequently. Combining this observation with the PES in Fig. 2(a), one may conclude that, as R_1 decreases, the neutral D_2 molecule depresses the PES of D_2^+ , leading to dissociation. Hence, when R_2 vibrates to a maximal value, R_1 of the complex simultaneously reaches a minimal value, resulting in a relatively high probability of D_3^+ formation.

Figures 4(a2) and 4(b2) display the average values of the trajectories on the R_1 and R_2 coordinates at each moment, reflecting the average reaction process in the T-shape configuration. The intersection of the two dashed gray lines in Fig. 4(b2) indicates the average time of D_3^+ formation, which is very close to the formation time calculated by the CTMC method. Consequently, analyzing the internal nuclear motion of the reaction can be achieved by tracking the average positions of the trajectories over time.

IV. INFLUENCE OF INITIAL NUCLEAR MOMENTUM

Up to now, we have assumed the initial momentum of the nuclear trajectory to be zero. We proceed to explore the possible influence of a distribution of initial nuclear momentum on the subsequent molecular dynamics. From the neutral ground state of the nuclear wave function $\psi_0(R_1, R_2)$, in principle, a four-dimensional Wigner quasiprobability distribution $\rho_0(R_1, P_1, R_2, P_2)$ needs to be constructed, where P_1 and P_2 are the conjugate momenta of R_1 and R_2 , respectively. This four-dimensional distribution is heavy to construct. However, the procedure can be simplified if one considers the dimensions of R_1 and R_2 to be independent of each other, or $\psi_0(R_1, R_2) = \psi_0(R_1) \otimes \psi_0(R_2)$, where $\psi_0(R_1) = \psi_0(R_1, R_2 = R_2^{(0)})$ is assessed at the equilibrium position of R_2 and $\psi_0(R_2) = \psi_0(R_1 = R_1^{(0)}, R_2)$ is assessed at the equilibrium position of R_1 . This can be justified by the fact that the dynamics in R_1 and R_2 are largely decoupled, which can be seen also from the contour plot of the initial-state distribution in Fig. 2, whose symmetric shape indicates it is roughly a direct product of the individual distributions in R_1 and R_2 . Thereby, the Wigner quasiprobability distribution can be approximated by $\rho_0(R_1, P_1, R_2, P_2) = \rho_0(R_1, P_1) \otimes \rho_0(R_2, P_2)$, where $\rho_0(R_i, P_i)$ is the individual quasiprobability distribution for coordinate R_i .

Shown in Figs. 5(a) and 5(b) are the Wigner quasiprobability distributions of $\rho_0(R_1, P_1)$ and $\rho_0(R_2, P_2)$, respectively. Clearly, the initial momentum is no longer simply zero. Rather, the trajectories are now sampled according to the quasiprobability distribution. With this Wigner sampling, we carry out CTMC simulations and extract the formation time distribution as shown in Fig. 5(c). Through fitting, the most probable reaction time is determined to be 136.4 fs, close to that obtained by CTMC simulations with zero initial velocity. It is thus clear that the initial distribution of nuclear

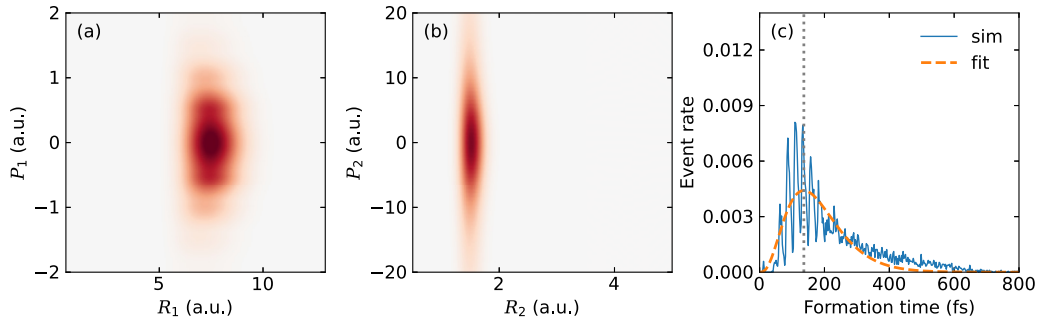


FIG. 5. Wigner quasiprobability distribution (a) $\rho_0(R_1, P_1)$ and (b) $\rho_0(R_2, P_2)$, and (c) the formation time distribution using CTMC with Wigner sampling.

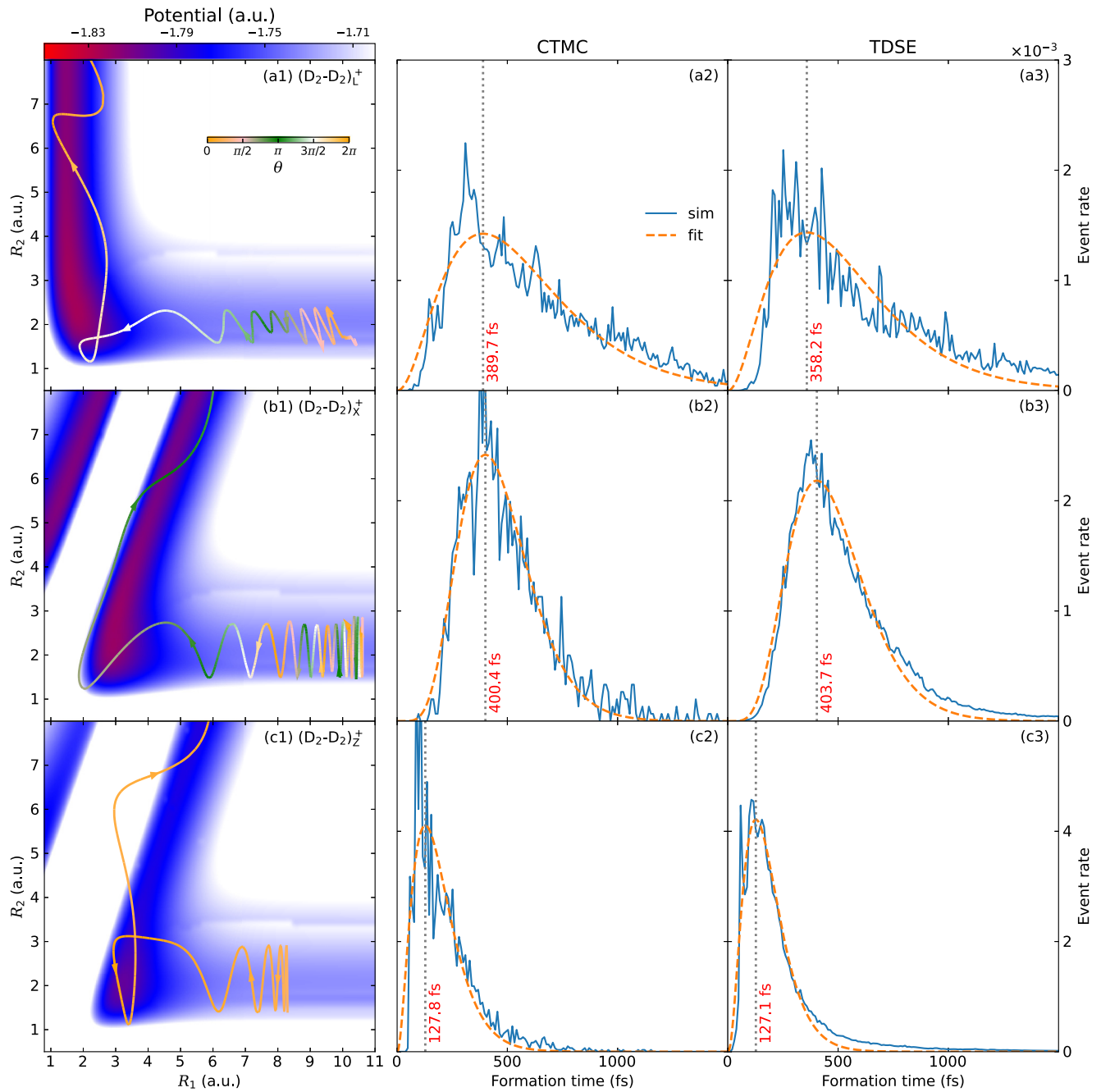


FIG. 6. Results starting from different initial configurations, including L-type [row (a)], X-type [row (b)], and Z-type [row (c)]. Column 1: PES of cationic $(D_2-D_2)^+$ at $\theta = 0$. Column 2: Histogram of the formation time calculated by CTMC. Column 3: Histogram of the formation time calculated by TDSE.

momentum has only minor influences on the subsequent molecular dynamics.

V. DIFFERENT INITIAL CONFIGURATIONS

In this section, we carry out a systematic exploration of the chemical reaction starting from various initial geometric configurations of the neutral dimer. For the sake of simplicity, we perform only CTMC and TDSE calculations for the L, X, and Z configurations, besides the T configuration discussed above. The PES at $\theta = 0$ and the corresponding histogram of the formation time for different configurations are presented in Fig. 6.

In the X and Z shapes, the definition of R_1 is the distance between the center of mass of D_2 and D_2^+ , which differs from that of the T and L shapes. A trajectory in X or Z shapes leading to the formation of D_3^+ exhibits a slope close to 2 in the R_1 - R_2 plane. Fitting the data to the Gamma distribution [Eq. (12)], we can obtain the peak reaction times for different initial configurations, as marked by the dashed gray lines.

As evident from Fig. 6, the calculated results are heavily dependent on the initial geometric configuration. The reaction times in L and X shapes are longer than those in T and Z shapes partially due to the larger equilibrium distance between two reactants in L and X shapes. Another important reason is the rotation of the molecule for L and X types (discussed below). Consequently, it takes more time for the two molecules to come together and react, forming D_3^+ .

In shapes other than T, rotation of the molecule is necessary to form D_3^+ , complicating the dynamics of the trajectories. To demonstrate the rotation dynamics, we use the color code to indicate the variation in the angular coordinate θ as the reaction progresses, as shown in Figs. 6(a1), 6(b1), and 6(c1). For L and X configurations, the molecule may rotate for a few rounds before D_3^+ could be possibly formed, leading to long formation times. In contrast, a rotation in the angle θ is not a must for the Z configuration, as demonstrated by the near uniform color of the typical trajectory in Fig. 6(c1). Thereby, the peak formation time is short for the Z configuration.

When both vibration and rotation are considered simultaneously, the multiple peaks in the time distribution, as observed in the T shape, become less distinct. This occurs because only when the rotation of the molecule reaches an appropriate dissociation angle can the dimer react to form D_3^+ . However, R_2 is not necessarily at a local maximum during this process, which makes it difficult to manifest the vibration of D_2^+ and significantly reduces the yield in L, X, and Z shapes.

The initial positions of the classical trajectories are sampled based on the ground-state nuclear wave function of the neutral dimer, inherently considering anharmonic effects to avoid the generation of nonphysical fast trajectories. This approach ensures the accuracy of the present calculations, leading to longer reaction times in all configurations compared to those reported in previous studies [17].

Notably, the formation times calculated by both CTMC and TDSE methods are comparable across all configurations, demonstrating the general applicability of both classical and quantum methods in the bimolecular reaction. Despite the disturbance induced by interference, our calculations pave

the way for integrating quantum methods into the studies of nuclear dynamics during chemical reactions.

VI. CONCLUSION

In summary, we have investigated the ultrafast bimolecular reaction of the D_2 - D_2 dimer upon photoionization, leading to the formation of D_3^+ . By exploring various initial geometric configurations (T, L, X, and Z shapes), we have provided a comprehensive understanding of nuclear motion from both classical and quantum perspectives. Several theoretical methods, including classical trajectory Monte Carlo, time-dependent Schrödinger equation, Bohmian mechanics, and the backpropagation method, were employed to dissect the reaction dynamics. By defining primary reaction coordinates, the potential energy surface was constructed for detailed analysis of the relevant reaction dynamics. We compared classical and quantum methods, affirming the reliability of these techniques. Notably, the quantum effects play an overall minor role in the reaction time. Our study further extends beyond the T configuration, venturing into the L, X, and Z shapes, providing a holistic view of the reaction dynamics. This systematic approach uncovers configuration-dependent reaction dynamics, demonstrating the important impact of geometric variations on reaction times. Anharmonic effects, essential in avoiding nonphysical trajectories, were inherently considered, ensuring the accuracy of the results. Our study on ultrafast bimolecular reactions not only enhances our understanding of detailed molecular dynamics but also lays the foundation for future studies, promising deeper insights into intricate quantum processes guiding chemical reactions.

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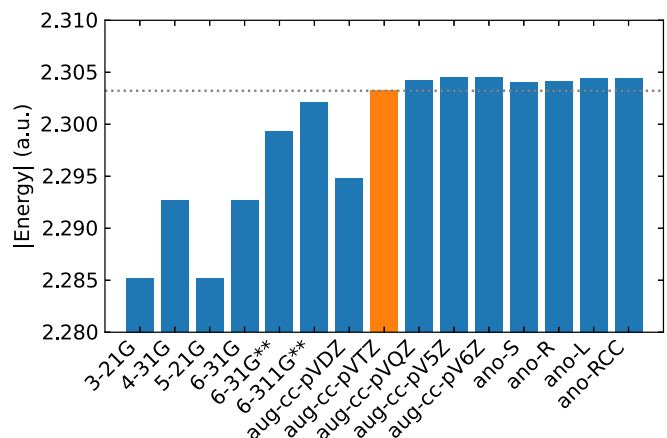


FIG. 7. The potential energy of the T-type D_2 - D_2 dimer at $R_1 = 7.131$ a.u. and $R_2 = 1.427$ a.u. using different basis sets.

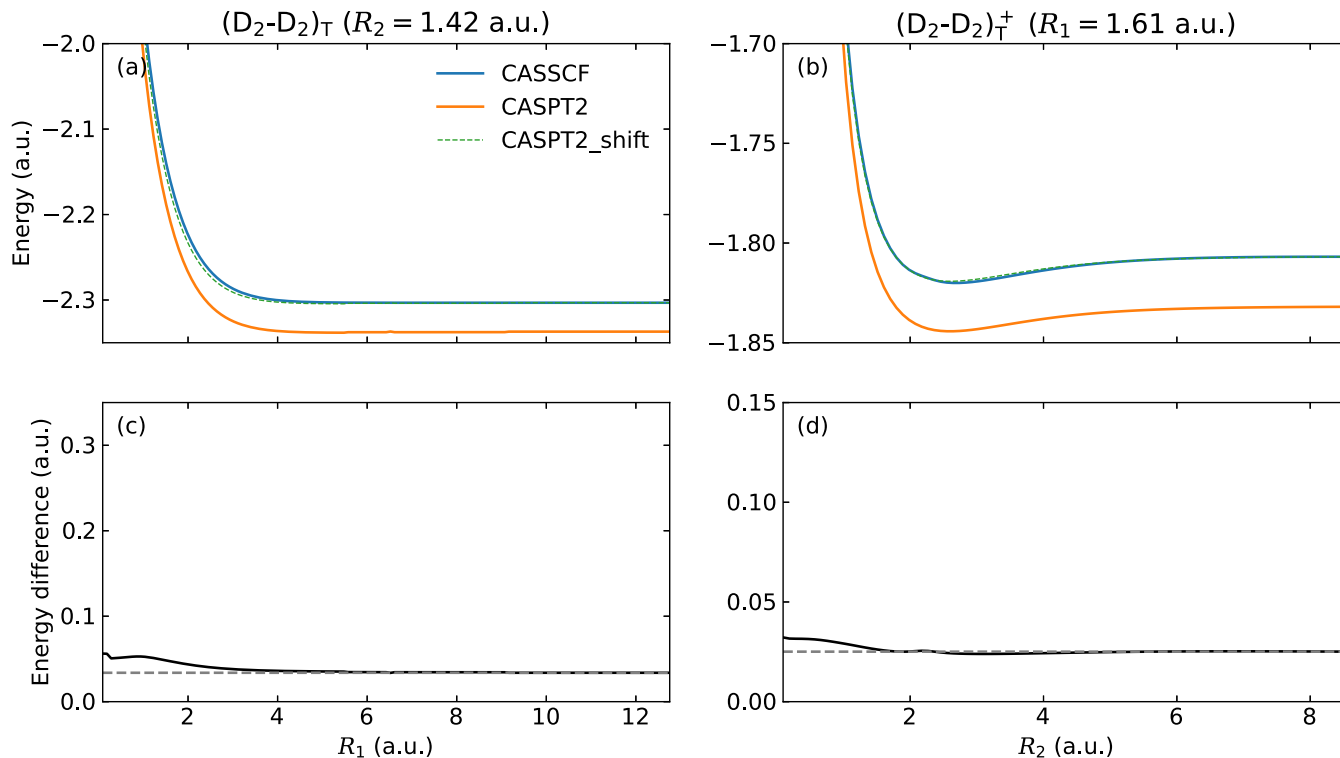


FIG. 8. The PES of the (a) neutral $(D_2-D_2)_T$ dimer at $R_2 = 1.42$ a.u. and (b) cationic $(D_2-D_2)_T^+$ dimer at $R_1 = 1.61$ a.u. calculated by CASSCF and CASPT2 methods. The black solid curves in panels (c) and (d) are the corresponding difference between the potential energy curves obtained by the two methods in panels (a) and (b).

APPENDIX: ACCURACY OF THE POTENTIAL ENERGY SURFACE

An accurate PES is important for the computation of the molecular dynamics. To check the accuracy of our calculations, we have performed the potential energy calculations of the neutral dimer in the T configuration at $R_1 = 7.131$ a.u. and $R_2 = 1.427$ a.u. using various basis sets, as shown in Fig. 7. It is clear that aug-cc-pVTZ is sufficient for the accuracy. Considering the calculation time, the aug-cc-pVTZ basis set is an ideal choice that balances speed and accuracy.

In addition, the PES is calculated using the CASSCF method. We have compared it with the more accurate method of complete-active-space second-order perturbation theory (CASPT2) in Fig. 8. Figure 8(a) is the calculated PES of the neutral $(D_2-D_2)_T$ dimer at $R_2 = 1.42$ a.u. and Fig. 8(b)

is that of the cationic $(D_2-D_2)_T^+$ dimer at $R_1 = 1.61$ a.u. Figures 8(c) and 8(d) are the corresponding difference in energy obtained by different theoretical methods. It is clear that CASPT2 gives an overall lower energy than CASSCF. The difference in energy is, however, a largely constant value. It is important to note that it is the spatial gradient of the potential, namely force, that governs the nuclear motion, as shown in Eq. (1). Thus, the overall shift in the potential energy is not important. Although the energy difference is not flat for very small R_1 and R_2 coordinates, the gradient in the potential itself is very large there, leading to strong repulsions, and the trajectories can hardly reach these regions. Hence, using the CASSCF PES only leads to minor difference from using the CASPT2 PES. Therefore, we have chosen CASSCF, which is computationally much less heavy, for the calculation of the PES.

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