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ARTICLE

Ion Molecule Reactions in the $\text{HBr}^+ + \text{CH}_4$ System: A combined experimental and theoretical study

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Received 00th January 20xx,
Accepted 00th January 20xx

DOI: 10.1039/x0xx00000x

Reactions in the system $\text{HBr}^+ + \text{CH}_4$ have been investigated inside a guided ion-beam apparatus under single-collision conditions. The HBr^+ is vibrational and rotational state selected in the electronic $X^2\Pi_{1/2}$ state created by (2+1)-REMPI. Due to the excitation scheme employed different rotational states of the HBr^+ are accessible. Four reaction channels have been observed. The cross section, σ , for the exothermic proton transfer channel (PT) decreases with increasing collision energy, steeper than predicted by the Langevin model. The cross section also decreases with increasing rotational energy in the HBr^+ , with the effect of the rotational energy being stronger than that of translational energy. The cross section for the endothermic charge transfer (CT) increased with increasing collision energy. The energy dependence is well reproduced by a simple line of center (loc) model. Although the bromine transfer (BT) is exothermic the observed cross section increased with increasing collision energy due to an activation barrier on the potential energy surface (PES). Analysis by a modified loc model suggest the relevance of an angle dependence of σ . The cross section for the endothermic hydrogen atom abstraction (HA) exhibits a maximum at 2 eV $E_{\text{c.m.}}$. The measured cross sections are rationalized by means of reaction dynamics simulations which shows good agreement with the experimental cross sections. The dynamics simulations are carried out with a machine learning potential that is developed and benchmarked with *ab initio* molecular dynamics simulation. The absolute cross sections predicted by reaction dynamics simulations are well within the same order of magnitude while reproducing the trends over three different collision energies for all four reaction channels. Furthermore, the simulations demonstrate various reaction mechanisms for these reaction channels, including a very interesting HBr^+ orientation selectivity for the BT reaction channel.

Introduction

Methane is an abundant chemical species influencing earth's atmosphere as well as extra-terrestrial atmospheres. Earth's atmosphere contains an amount of methane in the order of 2000 ppb. In the last four decades the methane concentration in the atmosphere increased dramatically^{1,2}. Li et al. investigated the influence of halogen species on the CH_4 loss and lifetime of the CH_4 in the atmosphere by simulations. This led to the conclusion that the rate of CH_4 loss will decrease and therefore the total lifetime of CH_4 in the atmosphere will increase until the end of the century³. Methane is also detected in the interstellar medium⁴, in the atmosphere of Jupiter⁵, on the Saturn moon Titan⁵, on Enceladus⁶ and Mars⁷. The atmospheres of Uranus and Neptune also contain CH_4 ⁸. HBr has been detected in the upper troposphere of Jupiter and Saturn with upper limits of 3 ppb respectively 1.6 ppb⁹.

Since activation barriers in ion-molecule reactions are in general small, these reactions play a major role in the interstellar medium and in planetary ionospheres. Additionally long-range interactions like Coulomb or ion induced dipole interactions favor reactions of this type¹⁰. Ion-molecule reactions (IMR) were essential in successfully describing the particle densities observed in interstellar clouds¹¹. The chemistry of the interstellar medium is characterized by low particle densities and low temperatures ranging from 10 K to 100 K^{12,13}. Due to the often barrierless IMRs small changes in the energetics of such systems can influence the outcome of this reaction type sensitively¹².

IMR also form the basis for Chemical Ionization (CI) Mass Spectrometry. Munson and Field exploited the formation of CH_5^+ from methane and its subsequent proton transfer reactions for analyzing unknown compounds^{14,15}. Ultimately, this led to the development the Proton Transfer Reaction Mass Spectrometry (PTR MS) for on-line trace analysis down to the ppb level¹⁶.

The CH_5^+ ion is an intriguing chemical species and was subject of interest in numerous studies. Under laboratory conditions CH_5^+ ions can be formed by radiative association of CH_3^+ with H_2 or by hydrogen atom abstraction in the reaction $\text{CH}_4^+ + \text{H}_2$ ^{12,17}. Clearly, the CH_5^+ ion is a rather flexible molecular ion exhibiting a potential energy surface with shallow minima.

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Electronic Supplementary Information (ESI) available: [details of any supplementary information available should be included here]. See DOI: 10.1039/x0xx00000x

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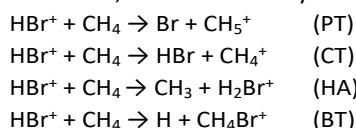


In fact, laboratory studies have been vital in reaching the current understanding of interstellar ion chemistry. Many studies aimed at measuring cross sections or rate constants either as a function of the temperature (in thermal ensembles) or as a function of the center of mass collision energy (typically in energy selected ensembles)^{18,19}.

Rather few studies have focused on the influence of the rotational quantum state of the reactants on the reaction dynamics. As a prominent example we mention the investigation of the reaction $\text{H}_2^+ (v=0,1 j=0,4) + \text{H}_2$ with a single and merged beam approach²⁰. The observed cross section decreased with increasing vibrational and rotational excitation of the H_2^+ ions, with the effect of the rotation exceeding that of the vibration²⁰. Viggiano et al. analyzed the $\text{Kr}^+ + \text{HCl}$ system in a SIFT apparatus. Here, the increase of the rotational temperature of the neutral target, HCl, increased the rate constant for charge transfer considerably²¹.

The role of ion rotation in IMR has been studied in a number of cases focusing on hydrogen halide ions, HX^+ with $x = \text{Cl}$ and Br , for which the rotational constants are large allowing to address rotational selectivity in ion preparation. Conceptually, rotational effects may be expected to be large in reactions involving hydrogen halide ions. It has been argued that there is potential relevance of the hydrogen halide ions in the upper atmosphere of earth as well as other astronomical objects²². Among the previous reaction systems investigated were the self-reactions of HCl^{23,24} and HBr²⁵. The authors groups also recently reported a combined experimental and theoretical study of the cross reaction system $\text{HBr}^+ + \text{HCl}$ ²⁶.

In the current work the study of rotational effects is extended to the system $\text{HBr}^+ + \text{CH}_4$. Here, four reaction channels are accessible, these are namely:



Experiment and Simulation

Experimental approach

The measured cross sections are analyzed using a guided ion beam setup. This was described in detail in earlier studies²⁷. Here, the apparatus was improved by implementing a second linear octopole stage to inhibit undesired side reactions²⁶. The setup is shown in Figure 1. The ions are created by a Laser as described below, then accelerated and by a focusing stage introduced into the first new linear octopole ion guide. The collisions occur in the second linear octopole ion guide.

Behind the second linear octopole ion guide the ions are

DOI: 10.1039/D4CP01121J

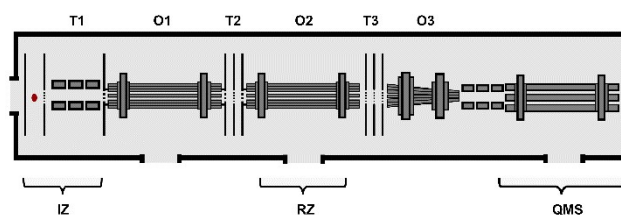


Figure 1. Guided ion beam setup of the apparatus. Ions are created between the first electrostatic lenses marked in red. O1 and O2 indicate linear octopoles, O3 a conical octopole. T1 T2 and T3 constitute electrostatic lens systems for transfer of the ions. IZ and RZ indicate the ionization zone and the reaction zone respectively.

transferred into a Quadrupole Mass Spectrometer (QMS) with the help of 2 lens assemblies and a conical octopole ion guide²⁸. HBr^+ was created in the $X^2\Pi_{1/2}$ electronic state by (2+1) resonance enhanced multiphoton ionization (REMPI) in its vibrational ground state^{29,30}. The REMPI transitions for ion creation were addressed with a tunable dye laser (CobraStrech, Sirah) which was pumped by a 20 Hz ND:YAG laser (INDI, Spectra Physics). The ion rotational energy was varied from 3.4 meV up to 46.8 meV. Experiments have been performed on the R(1), R(3), R(4), R(5), R(6) transition. The R(2) transition was not addressed since it overlaps with the S(0) transition. Ultimately, the REMPI excitation scheme allows to prepare HBr^+ ions with narrow rotational state distribution dominated by few rotational states. The measured rotational distribution of the ions prepared on the pump lines has been reported by Penno et al.³¹.

The collision energy in the center of mass frame was varied in the range from 0.25 eV up to 3 eV. The neutral reaction partner CH_4 was introduced at room temperature with a dosing valve. The ions were analyzed by their mass to charge ration (m/z) with a Quadrupole Mass Spectrometer (QMS) and detected by a Channeltron. The signals were recorded with a multichannel scaler card (FAST ComTec, P7888). The HBr pressure was set to $5 \cdot 10^{-6}$ mbar and the CH_4 pressure to $3.5 \cdot 10^{-5}$ mbar ensuring single collision conditions. The purity of the HBr gas was 3.5, that of the methane 4.5. The kinetic data in this study are based on the analysis of the ion species CH_4^+ , CH_5^+ , HBr^+ , H_2Br^+ and CH_4Br^+ .

The second-order rate constant k_i , where i indicates the reaction channel, is given in eq. (1) for the general case.

$$k_i = \frac{k'_{\text{tot}} \cdot f a_{\text{product}}}{1 - \exp(-k'_{\text{tot}} \cdot t)} \cdot \frac{1}{[\text{CH}_4]} \quad (1)$$

Here, k'_{tot} is the pseudo-first order rate constant of the total reaction, $f a_{\text{product}}$ is the fractional abundance of the respective product and $[\text{CH}_4]$ is the particle density of the neutral. Further details have been elaborated in the supplement by Plamper et al.²⁶

Rate constants can either be transformed into cross sections employing eq. (2) assuming the velocity of the neutral target is negligible compared to that of the ion under the conditions chosen, or directly via the approach described by Armentrout³². Here, the fact that the neutral target molecules are not at



rest but exhibit an isotropic velocity distribution in laboratory space does not affect the value of the center of mass collision energy but its distribution^{33,34}. This corresponds to an uncertainty in E_{cm} as discussed by Plamper et al.²⁶. As a consequence, reaction barriers may effectively be overcome at an E_{cm} nominally below that barrier. In the E_{cm} domain this uncertainty is on the order of ± 130 meV.

$$\sigma = \frac{k}{v_{HBr^+}} \quad (2)$$

Experimental cross sections are complemented by theoretical data, in part derived from analytical models, but ultimately also from sophisticated molecular dynamics calculations.

For the exothermic reaction channels the experimental data are compared to the Langevin-model. According to the Langevin-model, the cross section of an exothermic ion-molecule reaction is given by^{35,36}

$$\sigma_L(E_{cm}) = \pi \cdot \left(\frac{2 \cdot \alpha \cdot q^2}{E_{cm}} \right)^{0.5} \quad (3)$$

Here, q denotes the charge of the ion, α the polarizability of the neutral molecule. The Langevin cross section σ_L scales with $E_{cm}^{-0.5}$ and is considered as an upper limit (Langevin-limit) in the case of a polarizable target. Experimental data smaller than σ_L indicate that not each collision leads to reaction.

For target molecules with non-zero permanent dipole moment Su and Bowers developed a modified Langevin theory, the approximate dipole orientation (ADO) theory.³⁷ Since the permanent dipole moment of the CH_4 is very small (approx. 10^{-6} D at room temperature)³⁸, its justified to stay with the classical Langevin model of eq. (3). Empirically, the energy dependence of cross sections for exothermic reaction channels often deviates from the $(1/E_{cm})^{0.5}$ characteristic. To reflect this, the experimental data are modeled by eq. (4).

$$\sigma_{fit}(E_{cm}) = A \cdot \left(\frac{1}{E_{cm}} \right)^n \quad (4)$$

where A reflects a system-specific constant and n reflects the deviation from ideal Langevin behavior ($n = 0.5$). An overview on classical and quantum capture theory models can be found elsewhere³⁹.

For endothermic reaction channels, exhibiting an energetic threshold or a barrier to be overcome before reaction can proceed⁴⁰, the cross section can be modeled by eq. (5)⁴¹

$$\sigma(E) = A \cdot \frac{(E - E_0)^n}{E^m} \quad (5)$$

where E_0 denotes threshold for the reaction and A is scaling factor. If $n = m = 1$ this expression simplifies to the classical line of centres model (loc)^{40,42-45}. The loc model assumes that reaction can only occur if the collision energy along a line connecting the centres of masses is larger than an effective threshold barrier.^{46,40}

Throughout this work we will fix the value of n to $n=1$. The classical loc model would be expected to apply for e.g. atom-

atom reactions but also to reactions dominated by the center mass of two reactants.

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DOI: 10.1039/D4CP01121J

For more complex reactions it may be necessary to account for sterical effects. In the simplest case one can define two line of centers and the angle between these two lines. In the reaction system investigated in this work one loc is the HBr^+ axis, the second loc can be chosen to be the $Br - C$ axis. For such a situation Levine and Bernstein suggested eq. (6) which takes into account this effect of relative orientation in a parameterized form^{40,43,45}. Here, D is the critical separation, which can be smaller than the hard-sphere separation, E_0' is the negative derivative of the reaction energy profile with respect to the cosine of the angle between the two axis mentioned above⁴³. Eq. (6) will be applied to the bromine transfer channel.

$$\sigma(E) = \frac{\pi D^2 (E - E_0')^2}{4 \cdot E \cdot E_0'} \quad (6)$$

Alternative models for describing cross sections for endothermic reactions have been elaborated in the literature^{40,43,45,18,19,47}.

Potential Energy Profile Calculation

Although there have been studies on ion-molecule reaction involving HBr^+ ,^{26,48,49,50,51} a detailed potential energy profile of the $CH_4 + HBr^+$ reaction has not been reported. The relativistic effects experienced by the core electrons of Br could be accounted for by an appropriate basis set with an effective core potential (ECP)⁵², which replaces the explicit treatment of these electrons. Hundreds of combinations of computationally efficient methods (DFT and MP2) and basis sets with different ECPs were performed on a similar reaction ($HBr^+ + HCl$), where the frozen core MP2 (fc-MP2)⁵³ in general is more accurate in representing the benchmark potential energy profile computed at the CCSD(T)/cc-pVDZ//CCSD(T)/CBS level of theory⁴⁸. Therefore, fc-MP2 with five different ECPs for the bromine atoms (pseudopotential (PP)⁵⁴, LANL2DZ⁵⁵, LANL2DZdp^{55,56}, CRENL⁵⁷, Stuttgart RLC⁵⁸) and cc-pVTZ^{59,60} for carbon and hydrogen atoms have been examined to find an appropriate method for the stationary point geometry search. The heat of potential reactions is summarized in Table 1 and compared with experimental values. The latter are available for the PT, the CT and the HA reaction channels based on the 0 K heat of formation for HBr^+ (${}^2\Pi_{3/2}$) (1097.83 kJ/mol)⁶¹, Br (${}^2P_{3/2}$) (117.91 kJ/mol)⁶¹, CH_4 (-66.55 kJ/mol) CH_5^+ (921.98 kJ/mol), CH_3 (149.87 kJ/mol), CH_4^+ (1150.68 kJ/mol), H_2Br^+ (924.20 kJ/mol), HBr (-27.85 kJ/mol). It is important to note that neither fc-MP2 nor CCSD(T) accounts for spin-orbit (SO) coupling effect in HBr^+ or Br, whereas the experimental heat of formation of these species are measured at ground SO state. Therefore, the SO coupling energies reported by Fujioka et.al.⁴⁸, are used to calibrate the experimental heat of formation of HBr^+ (${}^2\Pi_{3/2}$) or Br (${}^2P_{3/2}$) to obtain the "spin-free" experimental heat of reaction. For HBr^+ , the four-fold degenerate spin free ${}^2\Pi$ state splits into two double-degenerate (${}^2\Pi_{1/2}$ and ${}^2\Pi_{3/2}$) states, thus the spin free



energy state lies halfway between the two SO states. Hence, half of the SO coupling energy (31.65 kJ/mol)^{62,63} is added to the experimental heat of formation of HBr⁺ (²P_{3/2}) to obtain the ²Π spin free energy of HBr⁺. For Br, the six-fold degenerate SO states split into double degenerate ²P_{1/2} state and quadruple degenerate ²P_{3/2} state. The SO states are split in 1:2 ratio and hence, the spin free energy state ²P is obtained by adding one-thirds of the SO coupling energy (43.64 kJ/mol) which is added to the experimental heat of formation of Br (²P_{3/2}). This protocol offers a fair comparison of the heats of reaction obtained between calculations and experiments^{24,48,49}. Among the five ECPs tested, only PP gives good agreement (RMSE = 6.32 kJ/mol) with the experimental heat of the reaction. Note the calculation without any ECP on Br gives comparable result (RMSE = 7.44 kJ/mol) as PP, but it is 5 times slower. Therefore, PP is selected as the ECP and fc-MP2/cc-pVTZ-PP is employed to explore the potential energy surface of the reaction. The transition states search is carried out with nudged elastic band

(NEB)⁶⁴ calculations and their connections to intermediates are confirmed with intrinsic reaction coordinate (IRC)⁶⁵ calculations. The identity of critical points is confirmed by harmonic frequency calculations with 3N-6 (N is the number of atoms in the system) positive frequencies for the intermediates, and 3N-7 positive frequencies for the transition states with one imaginary frequency along the reaction coordinate. The critical points identified at the fc-MP2/cc-pVTZ-PP level of theory is reoptimized with coupled cluster with single and double excitations and perturbative triples with explicitly correlated F12 method (CCSD(T)-F12)⁶⁶⁻⁶⁸ for better accuracy. As shown in Table 1, the RMSE associated with the CCSD(T)-F12/cc-pVDZ-PP-F12 level of theory is only 6.65 kJ/mol. The CCSD(T)-F12/cc-pVDZ-PP-F12 potential energy profile is used as the benchmark to screen computationally efficient method for dynamics simulations.

Table 1. Heat of various reactions. All numbers are given in kJ/mol, numbers in brackets are in eV. Zero-point energy is included and computed at the same level of theory.

Method	PT (p1): CH ₅ ⁺ + Br	HA (p2): CH ₃ + H ₂ Br ⁺	CT (p3): CH ₄ ⁺ + HBr	BT (p4): CH ₄ Br ⁺ + H	RMSE
Experimental at 0 K, SO ground state	8.61 (0.089)	42.79 (0.443)	91.55 (0.949)	-	-
"spin free" experimental	7.09	27.83	75.74	-	-
CCSD(T)-F12/cc-pVDZ-F12	5.08	30.71	86.71	25.75	6.65
-	15.63	27.62	82.61	8.08	7.44
PP	17.34	26.85	83.49	16.30	6.32
fc-MP2/cc-pVTZ-					
LANL2DZ	67.42	30.74	142.74	37.50	52.08
LANL2DZdp	38.19	38.01	112.20	21.44	28.28
CRENBL	110.40	-10.92	133.10	1.55	71.79
Stuttgart RLC	60.91	32.78	133.64	40.54	45.72

Reaction Dynamics Simulation

The bimolecular reactions of interest are also studied using *ab initio* molecular dynamics (AIMD) simulations, in which the trajectories are propagated classically using the energy gradients computed on the fly with the FC-MP2/6-311G(d,p)^{53,69} level of theory. FC-MP2/6-311G(d,p) has been shown to accurately represent the dynamics of similar reactions^{24,26} and its accuracy is further justified in this reaction system (see Results). Separated by 8.0 Å (center of mass distance) with random orientations, HBr⁺ is fixed at its ground rotational and vibrational state and the rotational and vibrational states of CH₄ are selected from a Boltzmann distribution at 298 K to mimic the experimental conditions. Three out of the five collision energies studied in the experiments have been simulated with AIMD, e.g., 0.5, 2.0 and 3.0 eV. A chemical dynamics software VENUS⁷⁰ is employed to sample the initial conditions and to propagate the trajectory, which is interfaced with NWChem⁷¹ for *ab initio* energy gradients. Velocity verlet⁷² with a time step of 0.15 fs is used to propagate the trajectories, and for those having energy jumps greater than 1 kcal/mol, the time step is reduced and restarted with the same initial conditions. The

simulation is stopped when the two molecules (either reactive or unreactive) are separated by a center of mass distance of 12 Å. The impact parameter (b) is sampled discretely with a fixed stride (Δb). Firstly, b_{max} (maximum impact parameter beyond which no reactive trajectories are observed) is identified by sampling 100 trajectories at each b with Δb of 0.5 Å from 0 to 6 Å. For collision energies of 0.5 eV, 2.0 and 3.0 eV, the b_{max} are 4.5 Å, 4 Å and 4 Å respectively. After detecting b_{max}, 50 (for 0.5 eV collision energy) or 60 (2.0 and 3.0 eV collision energy) AIMD trajectories are carried out at impact parameters separated by Δb of 0.25 Å between 0 and the b_{max} at each collision energy, yielding a total of 950, 1020 and 1020 trajectories for collision energies of 0.5, 2.0, and 3.0 eV, respectively.

The AIMD trajectories are further used as the training set to develop a machine learning (ML) potential to carry out simulations at a much faster rate. 171,955 unique geometries (selected by comparing distance matrices, made of the reverse of pairwise distances between all atoms in the system, of configurations from AIMD trajectories with a threshold of 0.001 Å⁻¹) and their energy and gradients are employed as the initial training (75%) and validation (25%) set. Several iterations of active learning were carried out – a preliminary trained ML-PES was used to propagate MLMD



trajectories to potentially obtain novel geometries, which (along with their energy and gradients) were added to the training set and produce a new ML-PES. Schnetpack^{73,74} v1.0 is used for all ML-PES training and predictions. The MLMD trajectories are simulated using ASE's velocity Verlet integrator with a timestep of 0.15 fs and stopped when any two atoms are more than 20 Å apart. The MLMD trajectories are simulated starting from $b = 0$ with fixed increment Δb of 0.1 Å to b_{\max} . The estimated b_{\max} is 4.5, 3.8 and 3.8 Å for 0.5, 2.0 and 3.0 eV collision energies. Trajectories are rejected and restarted if the vibrational energy is lower than the zero-point energy for any species. The number of trajectories simulated at collision energies of 0.5, 2.0, and 3.0 eV, are 10350, 8200 and 8200, respectively.

Results

Experimental results

Experiments have been performed with HBr^+ prepared in the upper SO state. The question whether the SO energy (0.328 eV^{62,63}) is available to the reaction coordinate, will be discussed below.

The analysis of the mass spectra results in the observation of four different reaction channels. These are namely the proton transfer (PT), the hydrogen abstraction (HA), the charge transfer (CT) and the bromine transfer (BT). The mass spectra is provided in the ESI. In Figure 2 the cross section of the total reaction and all observable reaction channels is shown as a function of the collision energy for an ion rotational energy of 3.4 meV.

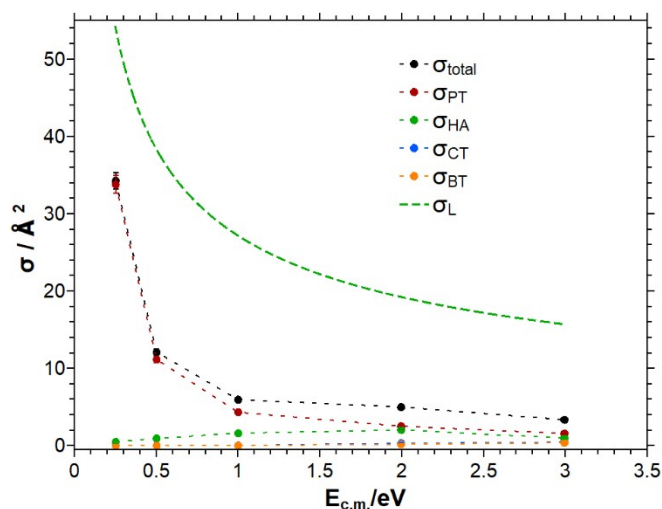


Figure 2. Cross section of the total reaction and all observable reaction channels as a function of the collision energy $E_{c.m.}$. The data is provided for the ion rotational energy of 3.4 meV. For comparison the cross section according to the Langevin model σ_L is shown. Note, that the CT data are difficult to distinguish from the BT data plotted in front.

The total cross section decreases with increasing collision energy as does the PT reaction cross section, which dominates the total cross section for collision energies below 2 eV. At collision energies above 2 eV the HA reaction becomes

comparable to the PT reaction. The CT and BT reactions are the least efficient reactions and both of the same order of magnitude. Below 1 eV collision energy the latter channels do not proceed to a measurable extent. The total cross section is significantly below the Langevin prediction indicating that approximately every 4th collision leads to a reactive process.

Tichy et al. studied the reaction of HBr^+ with CH_4 in a SIFDT apparatus. The cross sections measured by Tichy et al. are larger than the numbers measured in this work, but in the same order of magnitude⁷⁵. One difference between the work of Tichy et al. and this work concerns the ion preparation: in this work the HBr^+ is prepared in selected rovibronic states, whereas Tichy et al. employed electron impact ionization, presumably leading to a broader distribution of ionic states. In both studies, the PT reaction is the most efficient reaction at low collision energies. At collision energies around 1 to 2 eV center of mass the PT channel becomes comparable to the HA channel. The CT reaction is in both studies the least efficient reaction. In the present study this reaction channel has a threshold of around 0.5 eV, comparable to the earlier study by Tichy et al.⁷⁵

In Figure 3 the collision energy dependence of the PT reaction is presented. The different traces correspond to different ion rotational energies. The cross section decreases with increasing

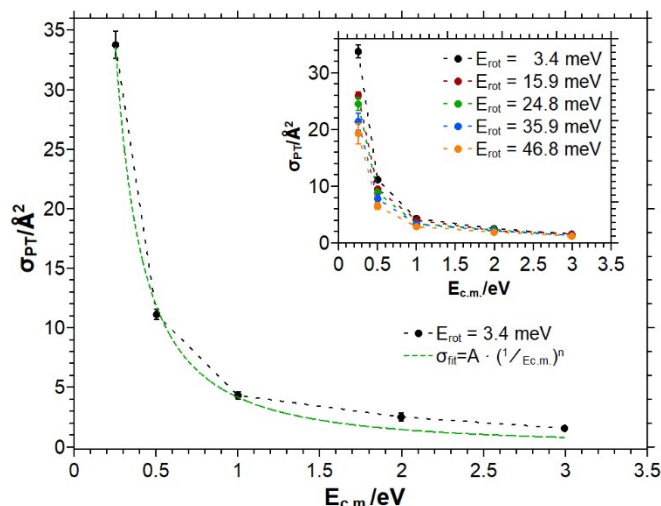


Figure 3. σ_{PT} as a function of the collision energy $E_{c.m.}$ for different ion rotational energies. The green dotted curve is the Langevin fit to the data for $E_{rot}=3.4$ meV according to eq. (4).

collision energy. Table 1 indicates a slightly positive heat of reaction for the PT channel referenced to the SO ground state of the HBr^+ . Here, the experimental observation of a cross section significantly decreasing with increasing collision energy suggests that the SO energy of the HBr^+ ion prepared in the $X^2\Pi_{1/2}$ electronic state is available to the reaction, turning it effectively into an exothermic channel.

In Table 2 the fit parameters for the PT reaction are assembled. Within the error margins the exponent n emerges constant. The decrease is much steeper than predicted by the Langevin model. With increasing rotational energy, the cross section



$E_{\text{rot}}/\text{meV}$	$A/\text{\AA}^2\text{eV}^{m-1}$	E_0/eV	m
3.4	1.81 ± 0.16	0.16 ± 0.04	0.70 ± 0.12
15.9	2.03 ± 0.19	0.18 ± 0.04	0.83 ± 0.14
24.8	1.95 ± 0.01	0.15 ± 0.01	0.72 ± 0.01
35.9	2.11 ± 0.41	0.06 ± 0.12	0.60 ± 0.23
46.8	1.92 ± 0.04	0.09 ± 0.01	0.62 ± 0.03

Table 2. Fit parameter A and n according to eq. (4) for the PT reaction

decreases significantly, which is recognizable in Figure 5 and indicated by the parameter A .

$E_{\text{rot}}/\text{meV}$	$A/(\text{\AA}^2\text{eV}^n)$	n
3.4	4.16 ± 0.47	-1.52 ± 0.09
15.9	3.93 ± 0.38	-1.37 ± 0.07
24.8	3.55 ± 0.39	-1.4 ± 0.09
35.9	3.3 ± 0.39	-1.36 ± 0.09
46.8	2.68 ± 0.43	-1.43 ± 0.12

Table 3. Fit parameter for the HA reaction according to eq. (5), using $n=1$.

Figure 5 illustrates σ_{PT} as a function of the ion rotational energy. The cross section is highly dependent on this degree of freedom. By using the Langevin fit for the rotational energy of 3.4 meV the cross section decreases by the same amount in collision energy by 8.4 \AA^2 whereas the same amount of rotational energy at 0.25 eV collision energy leads to a decrease of 13 \AA^2 . Thus, the rotational motion of the ion has a larger influence on the reaction dynamics than the center of mass collision energy and is not only an additive contribution to the total excess energy.

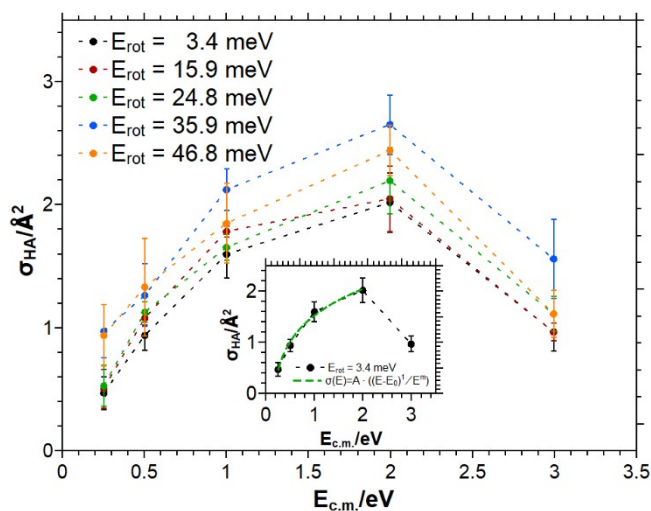
As indicated in Figure 4 the cross section of the HA reaction increases with increasing collision energy as expected for an endothermic reaction channel. The threshold behavior of σ_{HA} is modelled by eq. (5) fixing $n = 1$ in the range from $E_{\text{cm}} = 0.25 \text{ eV}$ up to 2 eV (i.e., the maximum of σ_{HA}). The fit parameters A , E_0 and m are summarized in Table 3.

Note, that the center of mass collisional energy as well as the rotational energy may help in overcoming the chemical threshold for the reaction. This fact is reflected in the observation, that the effective E_0 fitted to the data decreases with increasing rotational energy (cf. Table 3). The decrease in E_0 is in fact comparable to the concomitant increase in the rotational energy. The reaction barrier E_0 derived (ca. 0.16 eV) matches well the heat of reaction assuming the SO energy of HBr^+ is available to the reaction (0.443eV-0.328 eV).

At 2 eV E_{cm} , the cross section has a distinct maximum. At even higher collision energies the cross section decreases significantly. The increase up to 2 eV can be modelled by the classical loc model suggesting that no angle dependence of the

HA reaction is operative. This appears in line with the intuitive expectation, because the transition state for the HA reaction should not involve steric requirements. As complementation, a plot of the cross section for HA as a function of the rotational energy is presented in the SI.

In Figure 6 the collision energy dependence of the CT reaction is presented. The green dotted line is a fit to the data for $E_{\text{rot}} = 3.4 \text{ meV}$ according to eq. (5) with n and m equaling unity and a scaling factor A which reflects the line of centers (loc)



model.

Figure 4. σ_{HA} as a function of the collision energy shown for the investigated ion rotational energies. The applied fit model is discussed in the text below and summarized in eq. (5).

All parameters obtained in the analysis of the CT reaction

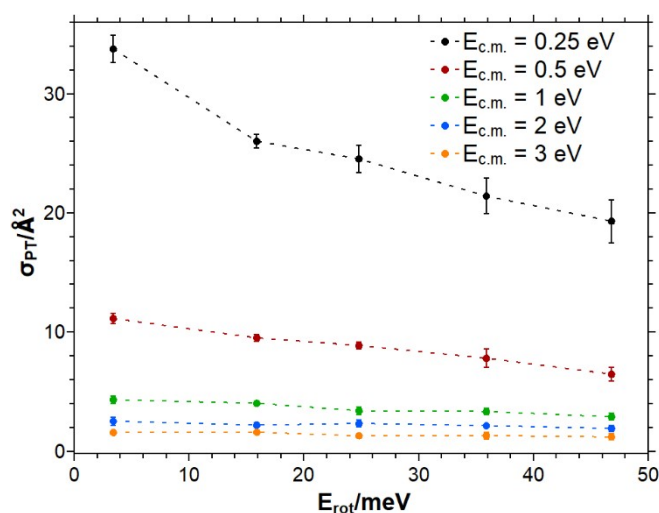


Figure 5. σ_{PT} as a function of E_{rot} .

channel are listed in Table 4.



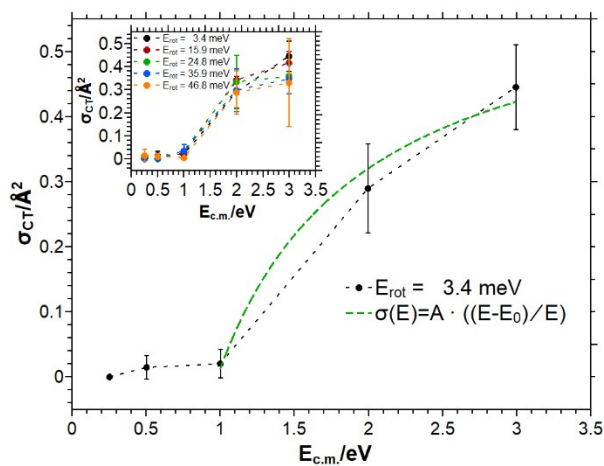


Figure 6. Collision energy dependence of the CT reaction. Different traces correspond to the different investigated ion rotational energies. The green dotted line is a line of centres model fit as indicated.

Table 4. Fit parameter for the collision energy dependence of the CT reaction according to the loc model.

$E_{\text{rot}} / \text{meV}$	$A / \text{\AA}^2$	E_0 / eV
3.4	0.63 ± 0.05	0.98 ± 0.06
15.9	0.63 ± 0.03	0.96 ± 0.03
24.8	0.55 ± 0.06	0.92 ± 0.08
35.9	0.52 ± 0.03	0.93 ± 0.05
46.8	0.52 ± 0.04	0.98 ± 0.06

The model fits the experimental data quite well. The threshold of the reaction, E_0 , is found to be 0.95 ± 0.03 eV for all investigated rotational energies. The almost exact agreement between this threshold and the experimental heat of reaction given in table 1 is fortuitous. In principle the true molecular threshold could be higher than the numbers given in table 1 due to the smearing out of center of mass collision energies³⁴. For a step like model for the cross section this could lead to observing products some 0.25 eV below the true threshold. For the rather shallow increase of σ with E_{cm} as operative in the loc model (and also in the modified loc model), convolution with the thermal velocity distribution of the neutral target reveals an effective additional smearing out which is below 50 meV. Given the limited number of data points we would be reluctant to overinterpret the threshold values. On the other hand, it suggests that the SO energy deposited in the HBr^+ ion upon preparation is not available for the reaction in this channel. Evaluating the PES of the CT reaction (cf. Figure 8 (c)) there is no indication of additional activation barriers. As complementation, a plot of the cross section for CT as a function of the rotational energy is presented in the SI.

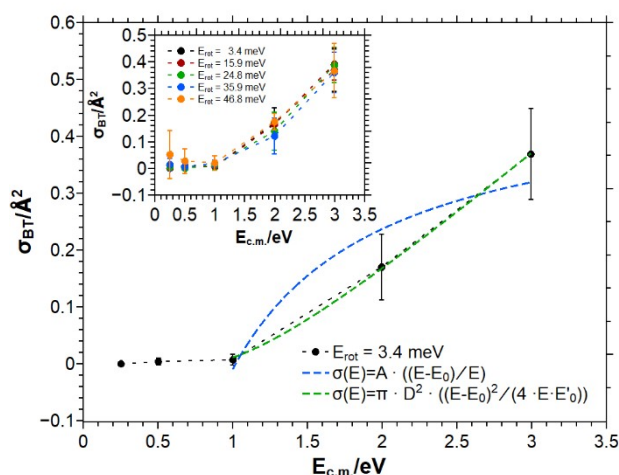


Figure 7. Experimental cross section of the BT reaction. The main graph shows the $E_{\text{rot}}=3.4\text{meV}$ data together with two fits. The blue dotted line is a fit according to the loc model, whereas the green dotted line originates from the improved loc model. The inset shows traces corresponding to additional rotational energies.

In Figure 7 the collision energy dependence of the BT reaction is presented. Overall, the efficiency of the BT reaction is comparable to that of the CT reaction. However, the characteristics of the variation of σ with E_{cm} is distinctly different.

Below 1 eV collision energy no reaction is observable. Above this threshold, the cross section for BT has been fitted both by the simple loc model and the modified loc model. Quite obviously the modified loc model fits significantly better to the experimental data shown in Figure 7. We conclude that the BT channel must be subject to steric restriction. Note, that in the modeling, the critical distance D was chosen to be 2.53 Å using the covalent radii of H, Br and C. An overview over the fit parameter obtained for all rotational energies of the HBr^+ is given in Table 5. As complementation, a plot of the cross section for BT as a function of the rotational energy is presented in the SI.

Table 5. fit parameter according to eq. (6) for the BT reaction.

$E_{\text{rot}}/\text{meV}$	E_0/eV	E_0'/eV
3.4	0.78 ± 0.04	22.26 ± 0.85
15.9	0.86 ± 0.01	19.49 ± 0.2
24.8	1.06 ± 0.08	16.11 ± 1.59
35.9	1.16 ± 0.07	15.56 ± 1.46
46.8	0.7 ± 0.01	24.25 ± 0.21

The data shows the typical behavior of a reaction with a threshold. The *ab-initio* potential energy surface (cf. Figure 8 (d)) shows an activation barrier for the spin-free BT reaction channel lying around 0.54 eV. Since the calculation of that PES is based on a spin-free Hamiltonian, the true spin-orbit energies for HBr^+ in the $^2\Pi_{3/2}$ and $^2\Pi_{1/2}$ would correlate with barriers being approximately 0.16 eV higher or lower respectively. The experimental finding of an effective threshold around 1 eV possibly indicates that the spin orbit energy of the



HBr^+ is not available for the BT reaction coordinate, similar to the CT.

As mentioned above the collision energy dependence of CT and BT reaction channels exhibits a characteristic difference. While σ_{CT} shows a right-curved characteristic above the threshold, well represented by the classical loc model, σ_{BT} shows a left-

curved characteristic above the threshold only represented by the modified loc model. As a consequence, it is concluded that the BT reaction channel is angle-dependent in contrast to the CT reaction channel, where steric orientation does not appear to play a role.

The Potential Energy Profile

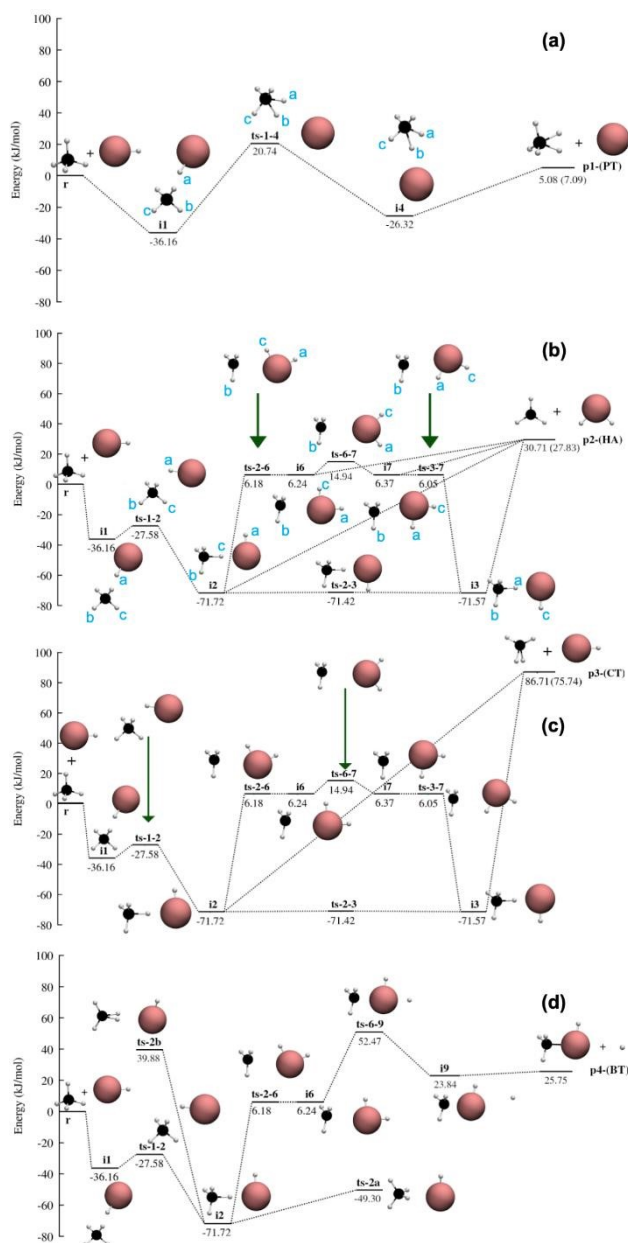


Figure 8. Potential energy profile of PT (a), HA (b), CT (c) and BT (d) reaction channels at CCSD(T)-F12/cc-pVDZ-PP-F12 level of theory with ZPE included. The atoms are represented as black (C), maroon (Br) and white (H). The energies in parentheses are experimental heats of reaction. The coordinates of these stationary points are given in the Supporting Information.

The potential energy profile of the proton transfer (PT) pathway is shown in Figure 8 (a), where the proton from HBr^+ is



transferred to CH₄ forming Br (²P) atom and CH₅⁺ (carbonium ion). The barrierless association between CH₄ and HBr⁺ (**r**) forms a van der Waals (vdW) complex **i1** (-36.16 kJ/mol with respect to separated reactant, **r**) where the proton from HBr⁺ (H_a) is pointing towards the CH₄ molecule. **i1** isomerizes to another vdW complex **i4** (-26.32 kJ/mol) *via* a transition state **ts-1-4** that is 20.74 kJ/mol above **r**. Following the IRC of **ts-1-4**, Br pivots around CH₅⁺ and moves from coordinating with H_a in **i1** to H_b (\angle C-H_b-Br = 180°) in **i4**. This pivoting mechanism is a result of the 3 centered 2 electron bond of H_a-C-H_b where H_a and H_b possess most of the partial positive charges. C_s symmetry is maintained throughout the path where Br-H_a-C-H_b-H_c are in the same plane. **i4** can dissociate without a transition state to form **p1** (Br (²P) + CH₅⁺). The PT reaction is endothermic by 5.08 kJ/mol with an overall barrier of 20.74 kJ/mol (**ts-1-4**). The structure and reactivity of CH₅⁺ has been extensively studied in the literature^{76–86} since its discovery in the 1950s which show that CH₅⁺ is highly fluxional with many different local minimum structures⁸⁷. But the global minimum among them is the C_s(I) structure⁸⁸ as depicted in the Fig. 1.a. It is important to note that experimental and previous dynamics study of CH₅⁺ have shown its tendency to dissociate into CH₃⁺ and H₂ molecule⁷⁷, which is out of the scope of the present work.

The potential energy profile of the hydrogen abstraction (HA) pathway is shown in Figure 8 (b), where HBr⁺ abstracts one hydrogen atom from CH₄ to form planar CH₃ radical (D_{3h}) and H₂Br⁺ (C_{2v}). The vdW complex **i1** (-36.16 kJ/mol, see the PT pathway) is connected to a submerged barrier **ts-1-2** (-27.58 kJ/mol), where HBr⁺ (instead of just Br as seen in the PT pathway) pivots around CH₄ to form **i2** (-71.72 kJ/mol). **i2**, stabilized by the hydrogen bond between H_c and Br, possesses a 'staggered' conformation where the dihedral angle of H_a-Br-C-H_b is 180°. **ts-2a** (-49.30 kJ/mol) (Fig. 1.d) is a self-isomerization transition state of **i2**, where CH₄ tumbles, but does not lead to the products. CH₄ in **i2** could also rotate with respect to C-H_c-Br axis to form the 'eclipsed' conformer **i3** (-71.57 kJ/mol), where the dihedral angle of H_a-Br-C-H_b is 0°. The corresponding transition state of this rotation, **ts-2-3** (-71.42 kJ/mol), is well less than 1 kJ/mol compared to **i2** and **i3**. The H atom in the CH₄ group of **i2** and **i3** can transfer to Br, forming **p2** without a transition state. **p2** could also be formed without a transition state *via* vdW complexes **i6** (6.24 kJ/mol) and **i7** (6.37 kJ/mol), where the H has already transferred from CH₄ to HBr, but the newly formed H₂Br⁺ has not dissociated. **i6** ('staggered') and **i7** ('eclipsed') are roaming complexes formed from its corresponding H-bond complex **i2** and **i3**, which can also isomerize to one another *via* transition state **ts-6-7** (14.94 kJ/mol). The C-Br distance changes from 2.92 Å (**i6**) → 3.21 Å (**ts-6-7**) → 2.92 Å (**i7**) in this process on a relative flat potential energy profile. The HA reaction is endothermic by 30.71 kJ/mol.

The potential energy profile of the charge transfer (CT) pathway is shown in Figure 8 (c), where an electron is transferred from CH₄ to HBr⁺. The CT reaction pathway is very similar to the HA pathway, except **p3** is only formed *via* a barrierless dissociation of **i2** and **i3** (not **i6** and **i7**). CH₄ changes from T_d point group to C_{2v} point group after losing an electron, while the H-Br bond

length decreased from 1.43 Å to 1.40 Å after gaining an electron. The CT reaction is the most endothermic (86.71 kJ/mol) among all the reaction pathways.

The potential energy profile of the bromine transfer (BT) is shown in Figure 8 (d), which forms CH₄Br⁺ (C_s) and H. As noted in the method section, although the experimental heat of formation of CH₄Br⁺ is not known to date, there is one ion-beam experiment tentatively speculating its formation⁸⁹. Theoretical investigation by Chistyakov et. al.⁹⁰ reported the geometries and energies of CH₄Br⁺ as intermediates of the CH₄ + Br⁺ → BrH + CH₃⁺ reaction. While the level of the theory (MNDO/PM3) in Chistyakov et. al. is not considered as state of the art anymore, its reported structure (Figure 8 (d)) agrees with CCSD(T)-F12/cc-pVDZ-PP-F12 geometry optimization in this study. Electron density analysis shows that the net positive charge is mostly concentrated on Br. The BT reaction pathway follows the reaction pathways of HA and CT until **i6**. The H₂Br in **i6** could recombine with CH₃ and form a vdW complex, **i9** (23.84 kJ/mol), after crossing a barrier (**ts-6-9**) of 52.74 kJ/mol. According to MP2/cc-pVTZ-PP, several vdW complexes similar to **i9** have been identified with the H atoms at different positions with respect to CH₄Br⁺, suggesting a roaming region in the potential energy surface. However, these stationary points could not be confirmed with CCSD(T)-F12/cc-pVDZ-PP-F12 level of theory, thus they are not reported in Figure 8. Further **i9** dissociates to **p4** without a transition state. The BT reaction is endothermic (25.75 kJ/mol) with an overall barrier of 52.47 kJ/mol (**ts-6-9**).

Computationally Efficient Method for AIMD Simulations

The potential energy profile is informative for a conceptual understanding of the mechanism of the reaction, but it relies on the assumptions that the reaction behaves statistically (sufficient lifetime of each intermediate) and strictly follows the IRC. Numerous examples have shown that these assumptions are not necessarily true in bimolecular reactions^{24,26,49,91–94}, thus *ab initio* molecular dynamics (AIMD) simulations are carried out. In AIMD, the positions of the atoms are propagated by the classical equations of motion with the potential energy gradients computed using quantum chemistry on the fly. AIMD demands an enormous number of gradient calculations, often in the order of millions to sample an ensemble of trajectories (of different orientations, impact parameters, etc.) that represent the experimental conditions. Therefore, it is imperative to screen different computationally efficient methods for one that can accurately characterize the potential energy profile. In the current study of CH₄ + HBr⁺ bimolecular reaction, the collision energies in the experiments range from 48.25 kJ/mol (0.5 eV) to 289.46 kJ/mol (3.0 eV). According to the CCSD(T)-F12/cc-pVTZ-PP-F12 potential energy profile, all stationary points reported in Figure 8 are within the energy range thus should be included in selecting the quantum chemistry method for AIMD.

Both frozen core and non-frozen core MP2 has been screened in this study due to our previous experiences with similar



reaction systems^{24,26,49}. 15 different basis sets, including 6-311G(d,p)^{95,96}, aug-cc-pVTZ-PP^{59,60,97}, aug-cc-pVTZ^{59,60,97}, pc-1^{98,99}, pc-2^{98,99}, aug-pc-1^{98,99}, aug-pc-2^{98,99}, cc-pVDZ^{59,60}, cc-pVDZ-PP^{59,60}, cc-pVTZ^{59,60}, cc-pVTZ-PP^{59,60}, def2-SVP¹⁰⁰, def2-SVPD¹⁰⁰, def2-TZVP¹⁰⁰ and def2-TZVPD¹⁰⁰, are screened. Each method/basis set attempts to identify the stationary points found by CCSD(T)-F12/cc-pVDZ-PP-F12. An appropriate method/basis set for AIMD simulation should be able to identify all stationary points in Figure 8 and have a small root mean square deviation (RMSD) with the benchmark PES. The RMSD of a candidate method/basis set (A) is computed as:

$$RMSD(A, ref) = \sqrt{\frac{1}{K} \sum_{i=1}^K (E_i^A - E_i^{ref} - \Delta E(A, ref))^2} \quad (7)$$

$$\Delta E(A, ref) = \arg \min_{\Delta E} \left(\frac{1}{K} \sum_{i=1}^K (E_i^A - E_i^{ref} - \Delta E)^2 \right) \\ = \bar{E}_A - \bar{E}_{ref} \quad (8)$$

K ($K=20$) is the number of stationary points (including reactants and products). The energy shift $\Delta E(A, ref)$ is difference between the average potential energy computed from

candidate method A (\bar{E}_A) and CCSD(T)-F12/cc-pVDZ-PP-F12 (\bar{E}_{ref}). i is the index of stationary points. Table 6 contains the RMSD of the screened candidate method/basis set for AIMD simulations. It is important to note that Table 6 only assesses the performance of these methods on the $\text{CH}_4 + \text{HBr}^+$ bimolecular collision and should not be taken as a comparison of their general performance. The same convergence criteria such as maximum (1.5×10^{-5} Hartree/Bohr) and RMSD (1.0×10^{-5} Hartree/Bohr) change in energy gradient, maximum (6.0×10^{-5} Bohr) and RMSD (4.0×10^{-5} Bohr) change in coordinates are applied to all candidate methods. fc-MP2/def2-SVPD, MP2/aug-cc-pVTZ, MP2/aug-cc-pVTZ-PP, and MP2/cc-pVTZ fail to optimize **i6**, **i7**, and **i9**, hence their RMSDs are marked as **n/a** and are not considered further. The results show that both fc-MP2 and MP2 have fairly low RMSD with all tested basis set, with the lowest (9.1 kJ/mol) and largest (16.1 kJ/mol) RMSD found in FC-MP2/cc-pVTZ-PP and MP2/pc-1, respectively. Considering the cost of MP2 method scales as N^5 ⁴⁸ where N is the number of basis functions which ranges between 52 (fc-MP2/cc-pVDZ-PP) and 220 (MP2/aug-pc-2), the cost of these candidate method/basis sets varies dramatically. Generally speaking, the level of theory with an RMSE of ~ 10 kJ/mol or smaller could be used for trial runs, where the energy conservation of the trajectory, the stability of molecular orbital, and the converging speed will be used to further select a quantum chemistry method for AIMD simulations.

Table 6. RMSD (kJ/mol) of each candidate method with respect to benchmark method. n/a indicates at least one of the optimal structure in Figure 8 could not be optimized.

fc-MP2	RMSD (kJ/mol)	MP2	RMSD (kJ/mol)
6-311G(d,p)	10.7	6-311G(d,p)	11.2
aug-cc-pVTZ	9.9	aug-cc-pVTZ	n/a
aug-cc-pVTZ-PP	10.5	aug-cc-pVTZ-PP	n/a
aug-pc-1	15.1	aug-pc-1	15.1
aug-pc-2	10.3	aug-pc-2	10.9
cc-pVDZ	11.4	cc-pVDZ	11.0
cc-pVDZ-PP	10.8	cc-pVDZ-PP	10.9
cc-pVTZ	9.4	cc-pVTZ	n/a
cc-pVTZ-PP	9.1	cc-pVTZ-PP	9.3
pc-1	15.6	pc-1	16.1
pc-2	12.4	pc-2	13.5
def2-SVP	12.4	def2-SVP	13.9
def2-SVPD	n/a	def2-SVPD	10.4
def2-TZVP	11.2	def2-TZVP	13.0
def2-TZVPD	11.5	def2-TZVPD	11.2



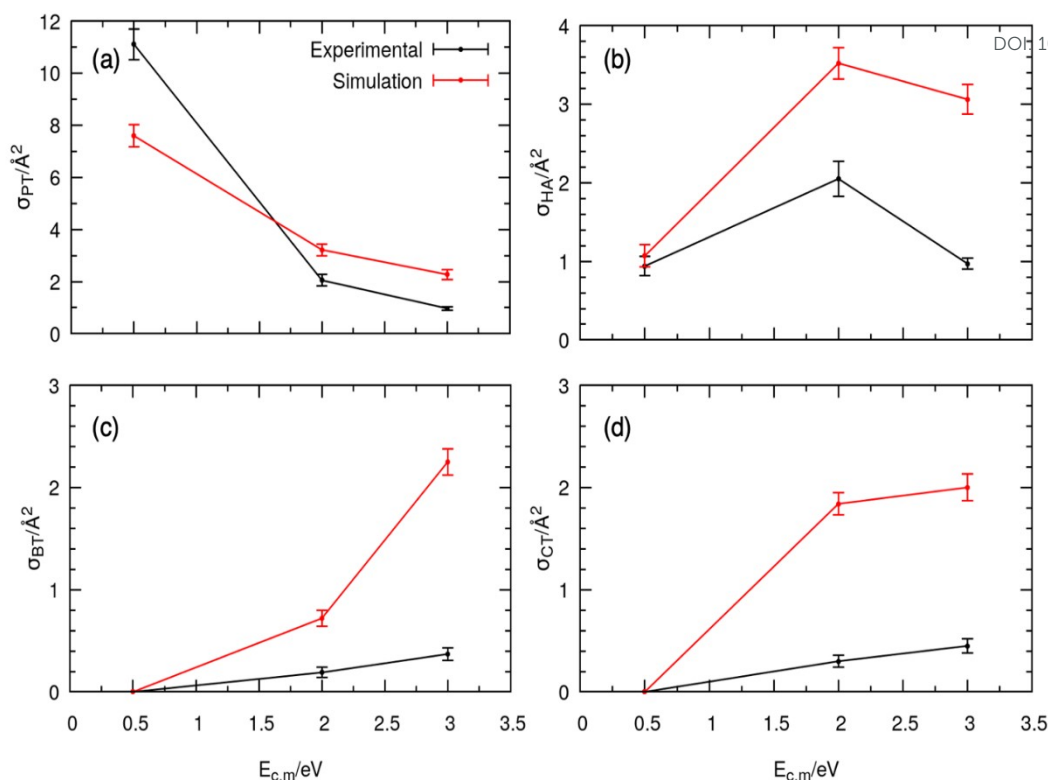


Figure 9. Reactive cross sections of the four product channels i.e., PT (a), HA (b), BT (c), and CT (d) measured from the experiments (black) and reaction dynamics simulations (red) at three collision energies.

Simulation results

All four reaction channels observed in experiments are detected in simulations and their cross sections are compared in Figure 9. First, it is interesting to note that trace amount of hydrogen exchange (HE, hydrogen from HBr^+ ion is exchanged with one of the hydrogens from neutral CH_4 molecule) reactions is detected in the simulation (e.g., cross section $\sim 0.01 \text{ \AA}^2$), which cannot be distinguished from the non-reactive collisions in the guided-ion beam experiments. Second, for the PT (proton from HBr^+ ion is transferred to CH_4 molecule, forming neutral Br atom and CH_5^+ carbonium ion) reaction, the cross section obtained from the simulation agrees well with the experiment. While the simulations in general overestimate the cross section of HA (hydrogen from CH_4 is transferred to HBr^+ ion, forming H_2Br^+ ion and neutral CH_3 radical), BT (the Br from HBr^+ is transferred to CH_4 molecule, forming CH_4Br^+ ion and hydrogen atom), and CT (single electron is transferred from neutral CH_4 molecule to HBr^+ ion, forming neutral HBr molecule and CH_4^+ ion) reactions, they qualitatively reproduce the trend observed in the experiment, including the local maximum observed in the HA reaction at 2.0 eV collision energy. The simulations also reproduce the relative branching ratios of the products, for example, at 0.5 eV collision energy, nearly $88 \pm 10\%$ of the reactive trajectories yield PT products and the remaining $12 \pm 2\%$ are HA products; no CT or BT products are observed. This is in good agreement with the experimental where the reaction cross section of CT

and BT products are negligible, while the cross section of HA is about one order of magnitude smaller than the PT. At higher collision energies (i.e., 2.0 eV and 3.0 eV), the agreement becomes only qualitative due to the overestimate of the cross section for HA, BT, and CT reactions. Nonetheless, the simulations agree with the experiments that PT is the most predominant reaction, followed by HA.



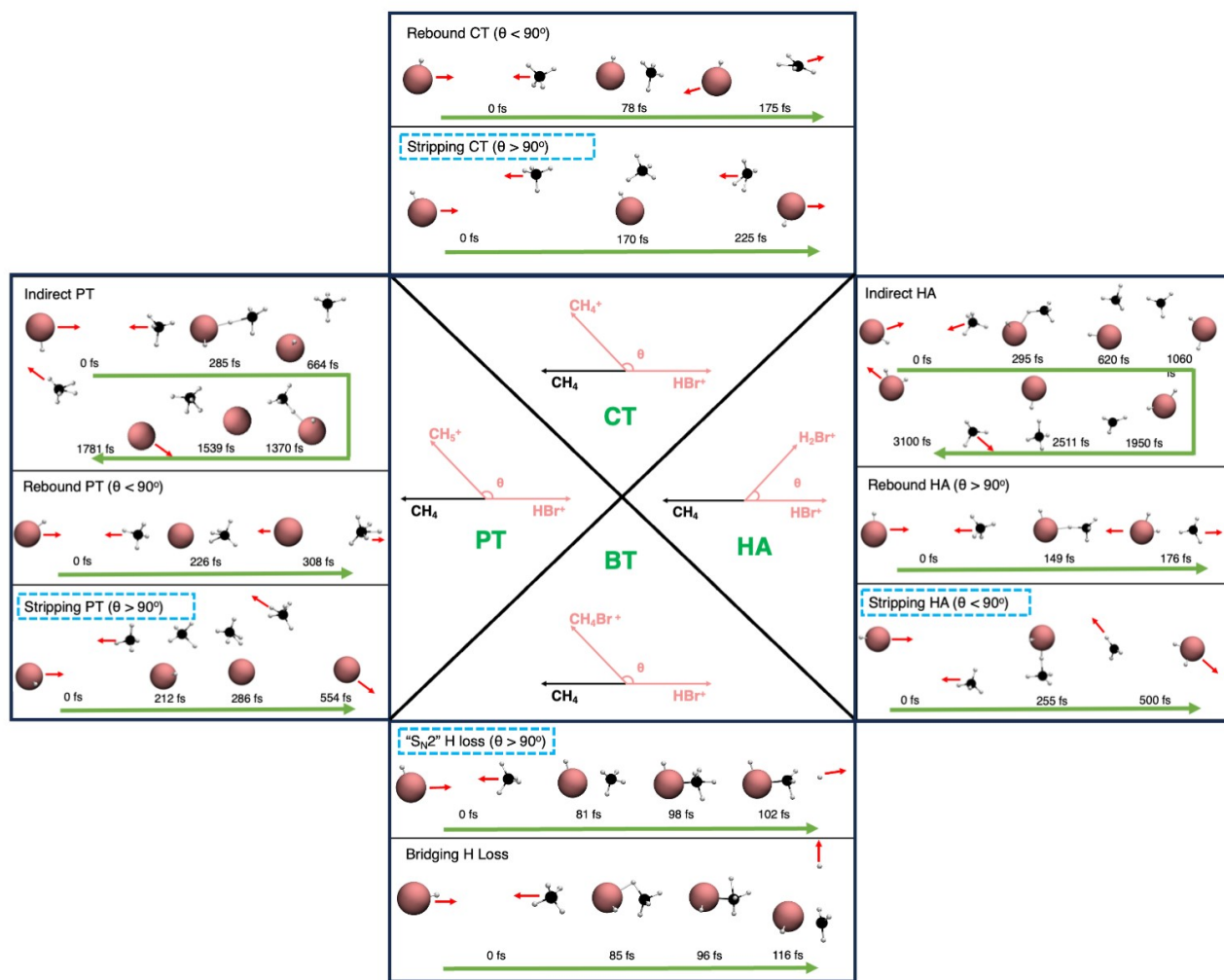


Figure 10. Illustration of the definition of scattering angle θ , which is defined as the angle between the velocities of the reactant and product ions (pink vectors). \vec{v}_i and \vec{v}_f denote the velocities before and after collision. Snapshots from representative trajectories of different reaction mechanisms are provided in the box. Blue dotted box denotes the dominant mechanism for each reaction channel.

ion-molecule collisions yield products *via* direct and indirect mechanism^{26,48,49,93,94,101–105} with their scattering angle measured according to the illustration in Figure 10. Reactive trajectories which do not form long-lived intermediates are classified as direct reactions. Following the convention of other ion-molecule reactions, direct reactions are classified as direct rebound (DR) and direct stripping (DS), where DR is mostly observed at small impact parameters and direct stripping (DS) is mostly observed at large impact parameters. The scattering

angle distribution of direct reactions of each reaction will be discussed in detail later in the manuscript. Reactive trajectories with significant lifetime of the collision complex are classified as indirect reactions, whose collision complex is long-lived (e.g., longer than the time for intramolecular vibrational energy redistribution) before dissociating into products, and as a result, yielding a near isotropic scattering angle distribution.



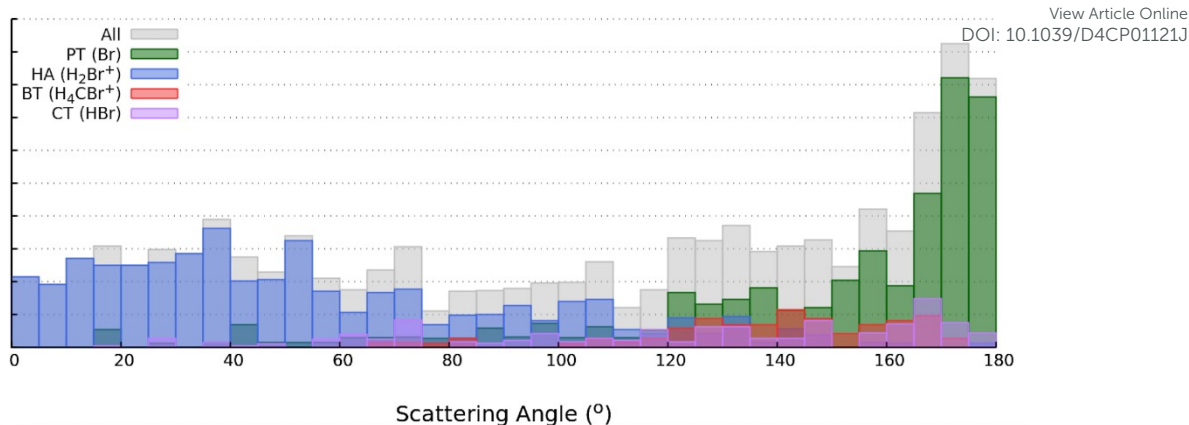


Figure 11. The scattering angle distribution (overall, gray; PT, green; HA, blue; BT, red; and pink, CT) from the simulation for 2.0 eV.

Snapshots of representative trajectories of different reaction mechanisms in all pathways are provided in Figure 10. Taking the dynamics of the reaction in the case of 2.0 eV collision energy as an example, a strong correlation between the reaction pathway and the scattering angle is observed. Figure 11 shows the scattering angle distributions for all four reaction pathways. As noted earlier, PT and HA are the predominant reactions. The PT reaction shows large scattering angles, where the product ion leaves in the opposite direction as the reactant ion, indicating the dominance of the DS mechanism (Figure S6 (b)). The DS mechanism is also the dominating mechanism for the HA reaction (Figure S6 (c)), but in this case, resulting in small scattering angle, where the product ion leaves in the same direction as the reactant ion. Overall, the dynamics of the PT and HA reactants are in accordance with what have been reported in similar ion-molecule bimolecular collisions, e.g., $\text{HBr}^+ + \text{HCl}$ and $\text{HCl}^+ + \text{HCl}$.^{24,26}

CT and BT are two minor reaction pathways observed in this reaction (Figure 9). CT reaction takes place only *via* direct mechanism across all impact parameters. This result is similar to what Luo et. al.²⁴ reported in the study of the $\text{HCl}^+ + \text{HCl}$ bimolecular collision, although the strong selectivity of the entrance channel complex (e.g., only those trajectories where chlorine collides with chlorine result in CT) is absent in the current system. More than 70% of the CT trajectories follow the DS mechanism (Figure S6 (a)), leading to large scattering angles. Similar to CT, all the BT products are formed *via* direct mechanism and the initial contact determines the reaction mechanism (Figure 10): (1) the bridging H loss: if the hydrogen in CH_4 is directly colliding onto the bromine of HBr^+ , it acts as a bridging atom between the C and Br, which is squeezed out immediately by these two heavy atoms to form the C-Br bond in CH_4Br^+ . In this case, the hydrogen atom (and the accompanying CH_4Br^+) traverses perpendicular to the C-Br axis, resulting in a scattering angle of near 90 degrees. This is evident from the scattering angle distribution in Fig. S6 (d). (2) the “ $\text{S}_{\text{N}}2$ ” H loss: if the CH_4 is orientated in a direction that allows for the bromine to directly form a bond with carbon, the H atom of CH_4 on the opposite side of the bromine will pop away. In this case,

the hydrogen atom (and the accompanying CH_4Br^+) traverses parallel along the C-Br axis, resulting in a near 180-degree scattering angle Figure S6 (d). Combining the scattering angles from all reactions, the overall scattering angle of the $\text{HBr}^+ + \text{CH}_4$ is shown in Figure 11.

Discussion

In this section, the results presented are put into the context of previous work. Four different reaction channels have been observed for the reaction system $\text{HBr}^+ + \text{CH}_4$, proton transfer (PT), hydrogen atom abstraction (HA), charge transfer (CT) and bromine cation transfer (BT).

The PT reaction shows a monotonically decreasing cross section with increasing collision energies implying that no significant barrier is operative on that reaction coordinate. The SO energy seems to be available to the reaction coordinate. The cross section for PT also decreases with increasing rotational energy. A similar trend was observed in earlier studies $\text{HBr}^+ + \text{HBr}$ ²⁵ and $\text{HBr}^+ + \text{HCl}$ ²⁶. In the reaction system $\text{HBr}^+ + \text{HCl}$ the influence of the rotational energy on the PT cross section was more pronounced than that of collision energy²⁶. The data presented in this work exhibit the same trend.

The cross section for the endothermic hydrogen abstraction can be fitted to the simple loc model up to $E_{\text{cm}} = 2$ eV. Beyond this collision energy the cross section decreases significantly. This can be rationalized by the switching-on of competing reactions, in particular the CT and the BT channel, which is observed also in the AIMD simulations. The PES prompt similar reaction pathways for the HA and BT channels with identical transition state followed by splitting up into different pathways. In the experiment the BT pathway appears significantly enhanced at the largest collision energies employed, while the HA channel appears suppressed at the largest collision energies. This observation comes along with the finding that the BT channel is the only channel exhibiting a quadratic increase of the cross section with increasing E_{cm} , which can be modelled by the modified loc model suggesting an angle-dependence of the



reaction, which is in line with the simulation results qualitatively.

In contrast, the cross section for CT can be fitted by the simple loc model which implies no angle dependence. The thermochemical threshold for the CT reaction pathway, as obtained from the calculated PES, is higher than the corresponding activation barrier of the BT reaction. This observation is consistent with experimental results, which also indicate a lower threshold for the BT reaction channel.

The cross sections for BT as well as CT are basically independent of the rotational energy of the ion, the cross section for HA has only a minor dependence on the ion rotation. In contrast, the cross section for PT markedly decreases with rotational energy of the ion, at least for the smallest E_{cm} investigated. To be effective, the PT channel requires the hydrogen atom pointing in the direction of the CH_4 . A higher rotational velocity will likely disfavor the passage through the transition state geometry.

For the HA and the CT reaction orientational requirements do not appear to pose a major restriction to the efficiency of those reaction channels. Consequently, the cross sections appear less affected by the rotational motion. However, the BT channel requires the Br atom of HBr^+ ion oriented towards CH_4 for the reaction to proceed.

Finally, the threshold for the respective reaction channels suggests that the spin-orbit energy of HBr^+ employed in the experiment may not automatically be available to the reaction. The SO energy in the HBr^+ ion appears to be available for the reaction for the PT and the HA channel, but not for the CT channel. We note, that earlier studies by Paetow et al. clearly showed that the spin orbit energy of the $X^2\Pi_{1/2}$ state is also available for the PT reaction in $HBr^+ + CO_2$ ^{27,106}.

It is of interest to note the behavior of the cross section of the PT reaction in AIMD simulations, which is slightly endothermic (5.08 kJ/mol) and has a non-negligible barrier (20.74 kJ/mol). Normally one would expect its cross section to increase as the collision energy increases but the simulation shows that the cross section decreases monotonically from 0.5 eV to 3.0 eV. We currently do not have a thorough explanation to this phenomenon, but we note the previous statement relies on maintaining a statistical ensemble that follows the intrinsic reaction coordinate. In the current system, the excess energy overwhelms the barrier and the heat of the reaction – the lowest collision energy simulated is 48.24 kJ/mol (0.5 eV). As a result, 47% percent of PT and 7% of HA trajectories are direct, as they simply glide over the intermediates and barrier reported in the potential energy profile (Fig. 8) and directly form the product. The ratio of direct reaction increases to over 90% when the collision energy is 289.44 kJ/mol. In these cases, an increase in the collision energy results in the reactants having even less time to interact with each other, thus the reaction probability decreases. A thought experiment can be laid out to make the point – even for an exothermic reaction, if the collision energy is infinite, the molecules would simply be passing through each other and no reaction can be observed. We note, that a similar observation has been arrived at with the $HBr^+ + CO_2$ reaction system⁴⁹. Considering a well-benchmarked ML potential has

been developed in this work, more AIMD simulations, below 0.5 eV will be conducted to verify the impact of the collision energy when it is below the barrier of the reaction and detect the turning point after which the cross section decreases with the increase of collision energy.”

Collision Energy (eV)	Direct trajectories	PT	Direct trajectories	HA
0.5	47%		7%	
2.0	84%		77%	
3.0	87%		91%	

The agreement between the simulations and experiments is qualitative, as shown in Fig. 9. The disagreement can be attributed to the lack of SO-coupling effect in the simulations and the differences between the SO-free PES used in the simulation and the true SO-free PES. However, it is important to realize that (1) the true SO-free PES is not accessible except for the heat of the reaction and (2) the PES for the simulations is selected from the benchmark, which only mimics the true PES. As shown in Table 6, there is a finite difference between the PES of the simulation and the benchmark thus only qualitative agreement can be expected. Nonetheless, it is our opinion that the level of agreement reported in this manuscript is better than other AIMD simulations of similar reactions, where only the trend of the cross section (e.g., normalized cross section) vs. collision energy is compared to the experiment^{24,49,26,107,108}. In other cases where the absolute cross sections are reported, they are off by one order of magnitude from the experiment^{109,110}. In particular there is almost quantitative agreement between the experimental and AIMD calculated cross sections for the dominating PT channel. The fact that both are lower by a factor of 3 compared to Langevin theory consistently indicates that on the average every third collision is reactive.

Summary

Four reaction channels in the system $HBr^+ + CH_4$ were investigated experimentally and analyzed. The PT exhibits characteristics of an exothermic reaction taking into account the SO energy in the HBr^+ . The HA and the CT are endothermic reaction with no dominant activation barriers observed in the PES. The BT pathway is dominated by an intermediate barrier. All measured cross sections are below the Langevin limit. The absolute values are observed between 34 \AA^2 and 0.1 \AA^2 .

The cross section of the PT reaction decreases with increasing collision energy as expected but steeper than predicted by the Langevin model. The monotonic decrease of σ_{PT} suggests that no effective barrier is operative in the PT reaction. σ_{PT} decreases with increasing rotational energy.

The HA cross section exhibits a maximum at 2 eV collision energy and decreases beyond this to the primal cross section value. The HA cross section is independent of the ion rotation with the exception at the point of similar angular speed of the reactants, where there is a maximum.



The CT reaction exhibits characteristics of a typical endothermic reaction. Experimentally the thermodynamic threshold was determined to be 0.95 ± 0.03 eV.

The cross section of the BT reaction fit quite well with a modified loc model, suggesting an angle-dependence to be operative. In contrast no angle-dependence appears operative for the CT reaction. For both reaction channels the cross sections are independent of the ion rotational motion in the range of collision energies investigated.

A machine learning potential for the title reaction system has been devised, which allowed the successful rationalization of experimentally measured cross sections as well as prediction of mechanistic aspects not directly accessible to the experiment.

Associated Content

There is supporting information available which contains the discussion of single collision conditions, the mass spectra and the data treatment.

Conflicts of interest

There are no conflicts to declare.

Author contributions

The experiments discussed have been conducted by D.P., the AIMD calculations and machine-learning potential have been worked out by K.F. and A.V.. The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

Funding Sources

Parts of this work have been supported by a grant from the Deutsche Forschungsgemeinschaft (We 1330-11). DP has been supported by a scholarship from the *Fonds der Chemischen Industrie* (FCI). Parts of this work has been supported by the National Science Foundation (USA) under Grant No. 2144031. The authors appreciate the information technology service (ITS) from the University of Hawai'i, Manoa for the computational resources.

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View Article Online
DOI: 10.1039/D4CP01121J

