Characterizing and optimizing a laser-desorption molecular beam source

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(Dated: 28 September 2017)

The design and characterization of a new laser-desorption molecular beam source, tailored for use in x-ray-free-electron-laser and ultrashort-pulse-laser imaging experiments, is presented. It consists of a single mechanical unit containing all source components, including the molecular-beam valve, the sample, and the fiber-coupled desorption laser, which is movable in five axes, as required for experiments at central facilities. Utilizing strong-field ionization, we characterize the produced molecular beam and evaluate the influence of desorption laser pulse energy, relative timing of valve opening and desorption laser, sample bar height, and which part of the molecular packet is probed on the sample properties. Strong-field ionization acts as a universal probe and allows to detect all species present in the molecular beam, and hence enables us to analyze the purity of the produced molecular beam, including molecular fragments. We present optimized experimental parameters for the production of the purest molecular beam, containing the highest yield of intact parent ions, which we find to be very sensitive to the placement of the desorbed-molecules plume within the supersonic expansion.

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I. INTRODUCTION

Laser desorption (LD) is a widely used technique to vaporize non-volatile organic molecules for gas-phase studies. The concept of LD is a rapid heating of the sample to be vaporized, at around $10^{10} - 10^{12}$ K/s, such that a fraction of molecules desorb intact instead of fragmenting. Later studies combined LD with pulsed molecular beams to directly cool the desorbed molecules, enabling the investigation of intact neutral molecules in the gas-phase at low vibrational temperatures. The main advantage of LD over other vaporization techniques, such as thermal vaporization, is the ability to introduce intact thermally labile organic molecules, including peptides and proteins, into a cold molecular beam. This has been demonstrated, e.g., for a pentapeptide (Ser-Ile-Val-Ser-Phe-NH$_2$) or the delta sleep inducing nonapeptide.

A first detailed characterization of a LD source coupled to a molecular beam (MB) valve was conducted nearly 30 years ago. Using anthracene, diphenylamine, and perylene, combined with resonance-enhanced multiphoton ionization (REMPI) spectroscopy, approximate vibrational temperatures of $< 15$ K and rotational temperatures of 5–10 K were determined. This demonstrated the ability of LDMB to gently vaporize large, thermally labile molecules and to efficiently cool them. Since then various spectroscopic techniques have been combined with LDMB sources and recent experiments have included resonance-enhanced multiphoton ionization studies, (far) infrared (IR)-ultraviolet (UV) double resonance techniques, IR multiphoton dissociation, and zero-kinetic-energy-photoelectron (ZEKE) spectroscopy.

In recent years, x-ray free-electron lasers (XFELs) have emerged as powerful tools for structure determination of gas-phase systems, with the potential to achieve atomic-resolution structures with femtosecond temporal resolution, recording so-called molecular movies. The ultrashort pulse duration available at XFELs enables the recording of a diffraction pattern from a molecule prior to destruction by the high intensity of the x-ray pulses. This diffraction-before-destruction paradigm, albeit still discussed, has recently also been demonstrated for isolated gas-phase molecules. Similar to the time-resolved nuclear dynamics that can be recorded at XFEL sources, modern laboratory based attosecond light sources allow the measurement of real-time electron dynamics in isolated molecules.

These experiments, however, are themselves inherently not species specific, i.e., all molecules within the interaction region will be probed. Therefore, combination of LD
with XFEL and attosecond experiments requires a pure molecular sample in the gas-phase. Furthermore, to be compatible with central facility light-sources, the laser desorption source needs to be translatable in three axes to adjust the molecular beam to the fixed XFEL beam. Additionally, the continuous measurement time should be as long as possible and the sample quickly exchangeable.

Here, we detail the characterization, and optimization of our novel LDMB source design, constructed to be compatible with central facilities, such as XFELs or attoscience centers. Using the dipeptide Ac-Phe-Cys-NH$_2$ as a prototypical labile biological molecule, which has first been laser desorbed and studied by the Rijs group,\textsuperscript{21,22} we characterize the created beam using strong-field ionization with a femtosecond laser pulse. This allows us to monitor all species present in the interaction region, including the carrier gas of the supersonic expansion. We show the optimization of experimental parameters to reduce fragmentation, to improve cooling of desorbed molecules and, thereby, to maximize the phase-space density of intact parent molecules in the interaction region. The created molecular beams are well-suited to further manipulation and purification, e.g., using electrostatic deflection techniques,\textsuperscript{23} an important step towards recording temporally and spatially resolved nuclear and electronic dynamics of isolated biomolecules.

II. EXPERIMENTAL SETUP

The mechanical design and construction of this LD source is based on compatibility with large-scale-facility-based photon sources. The laser-desorption source consists of a single central mechanical unit containing all necessary parts (molecular beam valve, sample bar with motors, and desorption laser optics). It is mounted on a three axis manipulator on a single flange for independent motion in the source chamber, which is pumped with a turbo molecular pump (Pfeiffer Vacuum HiPace 700P) to operating pressures typically around $10^{-5}$ mbar. It contains a cantilever piezo valve\textsuperscript{24} operated at 6 bar backing pressure of argon. The valve has a 300 $\mu$m orifice, followed by a conical opening of 4 mm length and 40° opening angle. Conical nozzle shapes are well known to produce molecular beams with more efficient translational cooling and greater directionality, and hence densities, than simple pinhole sources\textsuperscript{25} A graphite (Poco EDM-1) sample bar (80 mm long, sample channel width 1.2 mm) is placed approximately 200 $\mu$m in front of the valve, see Figure 1 The sample
FIG. 1. (a) Experimental setup for characterizing a laser desorption source. The source chamber contains the laser desorption source and is separated from the detection chamber by a 2 mm conical skimmer. The detection chamber houses a velocity-map imaging setup and microchannel plate detector. (b) Experimental timing diagram. The master trigger is given by the ionization laser, a kHz-repetition-rate femtosecond-pulse laser system that cannot trivially be externally triggered, and the molecular beam valve is triggered relative to this with delay $t_{\text{valve}}$. The desorption laser trigger is defined relative to the valve with delay $t_{\text{desorption}}$, while the energy of the desorption laser can be changed by modifying the Q-switch timing $t_{\text{energy}}$.

Bar height ($y$ axis) can be translated using an in-vacuum two-phase stepper motor (Owis SM.255.V6). To replenish the molecular sample, the sample bar can be moved along the $x$-direction using an in-vacuum linear piezo-stage (SmarAct SLC-24120-S-HV), typically operated at 0.02 mm/s. This results in measurement times of around 70 min per sample bar. For longer measurement periods the sample bar can be quickly exchanged with a load-lock system, pumped by a Pfeiffer Vacuum HiCube 80 Eco pumpstand (typical turn-around time 10 minutes). The entire molecule source (valve, sample-bar holder with motors, and desorption-laser optics) is placed on a three axis manipulator and can furthermore be adjusted for tip and tilt angle, allowing independent five-axes motion of the device within the vacuum chamber, as required for experiments at XFEL facilities. It is generally useful for operation of the source in molecular-beam setups were accurate alignment of the source is crucial, e.g., multi-skimmer setups or electrostatic manipulation devices. Detailed drawings of the source and individual components are given in the supplementary information.
Molecules on the sample bar (see below for sample preparation procedures) are desorbed by pulses from a fiber-coupled, diode-pumped Nd:YAG laser at 1064 nm (Innolas Spitlight Compact DPSS10), operating at 20 Hz with a pulse duration (full width at half maximum) of 9 ns and pulse energy up to 0.8 mJ. This is coupled into a multimode fiber (CeramOptec WF 400/440P) with core diameter 400 µm and numerical aperture of 0.22. The fiber is coupled into the vacuum chamber with a custom-made Swagelock connection and polytetrafluoroethylene (PTFE) plug. Inside the chamber the fiber is out-coupled with a custom-made vacuum compatible fiber collimator and the laser beam is focused to a spot size of approximately 0.6 mm on the sample bar. Custom mounting of the collimator allows variation of the laser spot size on the sample, as well as translation of the laser beam along the x and z axes, and tilting in the yz plane.

Following desorption, molecules are picked up by the supersonic argon jet from the valve and rapidly cooled down. The resulting molecular beam is skimmed with a 2 mm diameter skimmer (Beam Dynamics Inc. Model 50.8), located approximately 5 cm downstream the valve. Following the skimmer the molecular beam enters the differentially pumped (Pfeiffer Vacuum HiPace 2300) detection chamber, maintained at pressures around $3 \times 10^{-7}$ mbar. The detection chamber contains a velocity-map imaging (VMI) setup with a classic Eppink and Parker 3-plate design. The distance from the molecular beam valve to the interaction point is around 45 cm. For the results presented here, the VMI setup was operated as a time-of-flight mass spectrometer, with typical mass resolution $m/\Delta m \approx 100$.

The molecular beam is probed via strong-field ionization using a femtosecond Ti:Sapphire laser system (Spectra Physics Spitfire Ace) with a central wavelength of 800 nm, a pulse duration of 40 fs and pulse energies up to 300 µJ. It is focused into the vacuum chamber with a $f = 800$ mm lens to a spotsize (FWHM) of 80 µm in the interaction region between the VMI electrodes.

The timing scheme for our experimental setup is shown in Figure 1. The femtosecond ionization laser is used as the master trigger in the experiment. Its native repetition rate of 1 kHz is electronically reduced to provide a trigger signal at 20 Hz. The molecular beam valve trigger is defined, relative to this fs-laser trigger, by the delay $t_{valve}$. This delay defines which (longitudinal) part of the molecular beam we are probing with the ionization laser. The desorption laser is now triggered relative to the valve trigger and defined by the delay $t_{desorption}$, which controls where within the gas pulse the desorbed molecules are placed. This
FIG. 2. Time-of-flight mass spectrum of laser desorbed APCN following strong-field ionization. Peaks resulting from carbon or carbon clusters are labeled with *. 1 corresponds to the parent ion, and 2–6 to molecular fragments. A small signal from APCN dimer is also observed, 7. Grey shading marks the peak from argon carrier gas, which appears much smaller than molecular fragments due to the much higher ionization potential.

setup enables us to change $t_{\text{valve}}$ without changing $t_{\text{desorption}}$. We note that the timing values given should be seen as relative, not absolute values, as they are susceptible to electronic delays within the valve and laser controls used.

The velocity of the molecular beam is measured by recording the temporal profile of the beam, i.e., scanning $t_{\text{valve}}$, for different longitudinal positions of the valve, i.e., its distance from the first skimmer. We then evaluate the beam velocity from the temporal shift in the peak of the parent ion for different valve positions to be approximately 670 m/s.

The dipeptide Ac-Phe-Cys-NH$_2$ (APCN, 95% purity, antibodies-online GmbH) is used in this study without further purification. The sample powder is mixed with graphite powder (0.44:1 by weight) and ground with pestle and mortar to a fine powder. The top surface of the graphite sample bar is roughened with sand paper and pushed into the prepared sample mixture. Gentle force is used to ensure the mixture sticks to the sample bar and an even sample layer is formed.
III. RESULTS AND DISCUSSION

A. Molecular fragmentation

A measured time-of-flight mass spectrum of laser desorbed and strong-field ionized APCN is shown in Figure 2. It shows clear signals from parent ions (1 in Figure 2), fragments ions (2–6) and parent dimer (7). Furthermore, we observe several peaks from carbon and carbon clusters consisting of up to 11 atoms (highlighted with an asterisk in Figure 2). These are present due to their direct desorption from the graphite matrix material within which the sample is embedded, as well as due to the formation of higher-order carbon clusters within the desorption plasma created by the laser pulse. The molecular fragments originating from the APCN sample identified in the spectrum are shown in Figure 2. Strong-field ionization is a non species-selective method and thus allows the identification of all species present within the molecular beam. This approach, therefore, allows us to optimize the yield and fraction of intact parent molecules contained within the molecular beam. For further analysis of the contributing parameters for laser desorption, we identify four characteristic fragments; the APCN parent ion ($m/z = 309$, peak labeled 1), the $C_{10}H_{12}NO$ fragment ion ($m/z = 162$, 3), the argon ion peak ($m/z = 40$), and the carbon peak ($m/z = 12$). The particular molecular fragment (3) is chosen as it provides the largest-intensity clean signal, i.e., it does not overlap with a carbon cluster fragment.

In order to evaluate the effect of the femtosecond ionization laser on the observed molecular fragmentation, we scan the laser pulse energy between 60 µJ and 220 µJ. The observed integrated ion intensities for the 4 characteristic peaks are shown in Figure 3 b. The solid lines are a power-law fit of the form $A \times x^n + c$. Additionally, in panel (a) we plot the ratio of observed parent ions to the selected fragment ions for the probed intensity region. This nearly constant ratio indicates that the strong-field ionization process has little influence on the fragmentation of the parent ion. This is in agreement with previous studies that indicated that the fragmentation of complex molecules in intense laser fields is very sensitive to the laser pulse duration, but not the total energy of the pulse. Therefore, increasing the ionization laser intensities leads to larger ion signals, but has little effect on the fragmentation patterns observed. Hence using strong-field ionization with ultrashort laser pulses is a powerful tool for the full characterization of molecular beams containing complex molecules and fragments.
FIG. 3. Measured ion intensity for parent APCN, the characteristic fragment at \( m/z = 162 \), carbon and argon as a function of ionization laser pulse energy (a,b) and desorption laser pulse energy (c,d). Solid lines are power law fits to the data. The top panels (a,c) show the ratio of parent to fragment ions observed. Throughout the manuscripts, shown error bars correspond to 1 standard error (std. err.).

To elucidate the effect of the high-power desorption laser on the molecular sample on the graphite sample bar, we record mass spectra for various desorption laser energies in the range \( \sim 350–800 \) µJ. In Figure 3 d, we plot the recorded integrated ion intensities for the four characteristic masses as a function of desorption laser energy, with solid lines corresponding to a power law fit to the data. As can be seen from these data, all intensities increase with increasing laser energy. This includes, somewhat un-intuitively, the argon seed gas signal observed in the interaction region, which will be discussed later on. The carbon signal shows the steepest dependence on desorption laser pulse energy, which is consistent with the formation of isolated carbon atoms and clusters within a laser-induced plasma. Figure 3 c shows the ratio of parent to fragment signal. A decrease in the parent-to-fragment ratio is observed as the desorption energy is increased. Thus the desorption process can cause
fragmentation of the sample and the highest fraction of parent ion within the molecular beam is obtained at the lowest desorption energies, albeit at the expense of density within the beam. We furthermore note that the actual fractional yield of parent ions within the molecular beam is significantly smaller than the numerical values shown in Figure 3 since these only take into account a single characteristic fragment. Additionally, any charged fragments produced during the desorption process will not arrive in the detection region due to the static fields applied to the time-of-flight electrodes. All studies below were conducted with around 670 µJ desorption laser pulse energy and 140 µJ ionization laser power.

B. Molecular beam properties

To probe the longitudinal (or temporal) profile of the produced molecular beam, we scan the timing between the valve trigger and the ionization laser, \( t_{\text{valve}} \), see Figure 1, probing different portions of the molecular beam. A typical temporal profile is shown in Figure 4a for the pure argon beam emerging from the valve with (purple) and without (blue) the graphite sample bar in place. Without the sample bar, we observe a single sharp peak with a full width at half maximum of \( \sim 75 \, \mu s \) (corresponding to a speed ratio of \( \sim 10 \)), and a small shoulder at longer times due to the rebounce of the piezo within the valve. When the sample bar is placed in front of the valve, as shown in the picture inset in Figure 4, the argon gas flow is significantly disturbed. The overall gas-pulse is significantly broader, the main peak intensity is decreased by a factor of \( \sim 3 \), and more intensity is observed at later times. We attribute these observations to the disturbance of the argon flow by the sample bar and possible turbulences in the flow-field within the dead volume behind the sample bar.

By changing the relative timing of the desorption laser and valve trigger (\( t_{\text{desorption}} \) in Figure 1), we can now place the plume of desorbed molecules at different positions within the argon beam. Changing this relative timing has significant effects on the observed intensities of argon, parent, and fragment ions alike, as shown in Figure 4b. We show temporal molecular beam profiles for argon (diamond markers), fragment 3 (triangles), and parent ions (circles) at two different time delays \( t_{\text{desorption}} \). The relative timing of the desorption laser and molecular beam valve significantly affects the intensity of fragment and parent ions, and the ratio between the two. We attribute this effect to changes in the cooling efficiency as the hot desorbed molecules are placed within the argon expansion at different
FIG. 4. Integrated ion intensity as a function of valve delay $t_{\text{valve}}$, i.e., temporal (longitudinal) profiles of the molecular beam. (a) argon profile with (purple squares) and without (torquiose pentagons) sample bar in front of the valve. The inset shows a photograph of the sample bar in front of the valve at the optimized position; see text for details. (b) Temporal profile for APCN parent, characteristic fragment and argon for two different desorption laser delays, $t_{\text{desorption}}$. 

times. An efficient cooling process is required to quench the excess energy of the desorbed molecules and prevent further fragmentation. Comparing the relative integrated intensities of parent and fragment signals at the two timings shown in Figure 4 b, we observe that the combined intensity is approximately identical at the two time points, however the ratio
FIG. 5. Mass spectrum for the two different desorption laser timings shown in Figure 4(b), red and dark blue are the absolute intensities measured, the light blue trace has been scaled to the parent ion intensity of the red ($t_{\text{desorption}} = 80 \, \mu s$) trace. A significant increase in fragment ion yield is observed for $t_{\text{desorption}} = 40 \, \mu s$.

between the two differs significantly. This suggests that while approximately the same density of molecules, including fragments, is present within the initially desorbed plume, the less efficient cooling in the less-dense front of the gas pulse at the delay of 40 µs leads to significant fragmentation occurring before or during the argon gas pulse, i.e., the cooling occurs too late and fragmentation has already taken place.

Further to its influence on the molecular signals the timing of the desorption laser clearly has an effect on the observed argon signal. An increase in argon signal is observed at the timing where desorbed molecules (parent or fragments) are present within the beam. We suspect the observed increase in the argon signal is not due to more argon arriving at the detector, but due to signals of molecular fragments or carbon clusters overlapping with the argon mass at 40 u in a very crowded spectrum shown in Figure 2.

The dependence of the observed fragmentation on the relative position in the gas pulse is, furthermore, evident when comparing mass spectra recorded at different $t_{\text{desorption}}$, as shown in Figure 5. These spectra have been recorded with $t_{\text{valve}}$ optimized for parent ion signal and
are plotted normalized with respect to the observed parent intensity. This shows that all molecular fragments are significantly more abundant at $t_{\text{desorption}} = 40 \mu s$, indicating a much higher internal temperature of the desorbed molecules, due to the less efficient cooling, and correspondingly increased fragmentation.

To identify the optimum placement of the desorbed plume within the gas pulse, we have repeated these measurements for several delay points, shown in Figure 6, where the curve with higher intensity always corresponds to the fragment and the curve with lower intensity to the parent ion. For better comparison the parent intensity for each $t_{\text{desorption}}$, as well as the ratio of parent to the characteristic fragment, is shown in Figure 7.

For our setup and under the given experimental conditions, we identify a delay of 80 $\mu s$ as providing the highest total intensity of parent signal, as well as the best parent-fragment ratio. This ratio is very sensitive to the relative timings, and changes of 10 $\mu s$ can change this ratio by a factor of $\sim 2$. This is due to the gas pulse directly after the valve being significantly shorter (opening time of the piezo is around 25 $\mu s$) than in the detection region located $\sim 0.5$ m downstream where the measurements were taken. This is also reflected by the steep falloff of signal for later desorption laser timings.

As shown above, the sample bar has a large effect on the supersonic expansion and hence the produced molecular beam. To investigate this further we have taken data for different heights of the sample bar, shown in Figure 8. Here, we recorded the intensity of argon, carbon, fragment and parent ions for different heights of the sample bar, as well as the ratio of detected parent to fragment ions. We note that all this data was taken at identical timing of $t_{\text{valve}} = -900 \mu s$ and $t_{\text{desorption}} = 80 \mu s$ and that the height of the sample bar does change the desorption laser focusing conditions (since the focusing lens is fixed relative to the valve).

It is evident from this data that the sample bar height changes not only the peak intensity, but also the parent to fragment ratio, as shown in Figure 8(a). The observed intensities are very sensitive to the height of the sample bar, with parent, fragment and carbon ions showing maxima at different positions. This sensitivity was used in all previous measurements to fix the height for each new sample bar measured; it was optimized prior to taking data to obtain maximum signal from the parent ion. However, we note that due to the preparation method, slight differences in height can persist across a single sample bar, and affect the measurement and especially the comparability between data sets. This could be the reason for the large deviation of the data shown for 75 $\mu s$ in Figure 7. This sensitivity can
be explained by a number of contributing factors; (i) the efficiency with which desorbed molecules are picked up and carried by the argon beam. Assuming that the slight differences in mass and size between parent and fragments are negligible given the very large number of collisions with the carrier gas, this should be comparable for all species within the beam, i.e., a lower pressure of argon should affect all species to a comparable extent. (ii) The degree to
FIG. 7. Dependence of the parent ion intensity and the parent to fragment ratio on the desorption laser timing $t_{\text{desorption}}$; error bars correspond to 1 standard error. Optimum conditions are observed for $t_{\text{desorption}} = 80$ µs. The data at 75 µs shows a significantly lower intensity than expected, which could be due to irregularities in sample bar preparation; see text for details.

which the argon expansion is disturbed by the presence of the sample bar. In [Figure 4]a we have shown that the sample bar influences the molecular beam speed and distribution; at the fixed $t_{\text{valve}}$ conditions used here this will change the observed intensities. It has furthermore been shown already that the sample bar changes the directionality of the molecular beam.[31]

(iii) Changes in the cooling efficiency of desorbed species lead to differences in the parent to fragment ratio in the beam. (iv) The spotsize of the desorption laser on the sample bar changes with different sample bar heights and thus the intensity of the laser and the number of molecules interacting with the laser is influenced. From the collected data we cannot comment on the relative importance of these different mechanisms, however, since (i) and (ii) should influence the parent and fragment molecules nearly identically, the observed changes in the parent to fragment ratio indicate a dependence of the cooling efficiency on the sample-bar height. The cooling efficiency should be best within the densest part of the molecular beam, which is on the axis of the 300 µm nozzle orifice. Therefore, a sample bar height just below this position, i.e., covering slightly less than half the nozzle opening, should lead to the densest plume of desorbed molecules being entrained in the densest part of the molecular beam. This simple consideration is consistent with our observations of the maximum parent signal and parent to fragment ratio occurring at the position labeled 1.3 mm. While we cannot disentangle all the different effects of changing the sample bar height, it is clear that this, and the associated influence on the supersonic expansion and molecular
FIG. 8. Measured ion intensity for parent APCN, the characteristic fragment at $m/z = 162$, carbon and argon as a function of sample bar height. The top panel shows the ratio of parent to fragment ions observed.

beam properties, is a crucial parameter for laser desorption entrainment of molecules into supersonic expansions. This could be further investigated by either measuring the gas flow from the nozzle directly, for example through direct visualization of gas densities or by measuring spatial argon profiles through strong-field-ionization mass spectrometry at various distances from the nozzle.

Despite the wide use of laser desorption sources, very few studies have looked into the fundamental underlying processes. Furthermore, the vastly different source designs in use, e.g., different desorption laser wavelengths, intensities, pulse durations, different models of supersonic valves, etc., make comparison to previous studies difficult. While we believe this is the first study of laser desorption using strong-field ionization, previous experiments have utilized electron impact ionization and have similarly observed a large amount of neutral fragments produced by the desorption processes. Our finding that the molecular packet
of desorbed molecules is much shorter than the envelope of the seeding-gas pulse is also consistent with previous measurements. Lastly, we point out that the use of a fiber-coupled desorption laser has been demonstrated before, albeit without refocusing inside the vacuum chamber that we have introduced here for greater control.

IV. CONCLUSION

We have presented a novel laser-desorption setup designed for use in advanced imaging experiments of ultrafast molecular dynamics and we have carefully characterized and optimized the laser-desorption and molecular-beam-entrainment conditions. The setup consists of a single central mechanical unit containing all necessary parts (molecular beam valve, sample bar with motors and desorption laser optics) that is mounted on an XYZ manipulator on a single flange for independent motion. Furthermore, we have presented a detailed characterization of our new laser-desorption source as well as molecular beams produced using laser desorption in general. By utilizing strong-field ionization, we were able to probe all species contained within the beam. Under normal operating conditions we found that the molecular beam contains, in addition to parent molecules, significant amounts of molecular fragments, as well as carbon clusters from the desorption process. We investigated the role of the desorption laser fluence, the relative timing of valve opening and desorption laser, the sample bar height, and which part of the molecular packet is probed. While increased desorption laser fluence leads to more molecules contained within the molecular beam, it was found to induce fragmentation of the sample and leads to enhanced contamination of the beam with carbon and its clusters. The placement of the desorbed plume of molecules within the gas pulse from the supersonic expansion has a profound effect on the cooling efficiency, and thus the fragmentation observed. The best timing was found to be approximately in the center of the gas pulse, and is quite sensitive compared to the gas pulse duration in the detection region. The relative height of the sample bar in front of the valve orifice significantly affects the molecular beam expansion conditions, and hence the intensity of observed signals, as well as the parent to fragment ratio. However, finding the optimum position for the sample bar height is difficult, due to the number of competing effects taking place, and every sample bar being unique. Furthermore, parameters might be dependent on the employed molecular-beam nozzle, and our exact findings are specific to the used conical
nozzle shape.

From our detailed investigation we found that the optimal settings for building a laser desorption source very much depend on the planned experimental scheme. While some parameters, such as the relative timing of desorption laser and the molecular beam valve, should always be optimized as shown here, other parameters are not critical. For example the pulse energy of the desorption laser should be chosen according to the application. For techniques that are only sensitive to intact parent molecule signal, like resonance-enhanced multiphoton ionization, the pulse energy of the desorption laser should be kept high because this increases the number density of parent molecules in the interaction region. But for non-species specific techniques, such as x-ray diffraction, the pulse energy should be reduced to minimize the contamination with fragments and carbon. However, even at the lowest desorption energy used here, we still observe a significant amount of molecular fragments and carbon clusters in the beam. While the former originate to some extent from the strong-field-ionization probing, the carbon and carbon clusters are certainly in the beam due to the desorption process. In order to produce a pure beam of intact parent molecules in the gas-phase, one can consider coupling a laser desorption source with other species separation techniques for neutral molecules, such as electrostatic deflection or alternating gradient focusing [23,34] and such experiments are currently underway in our laboratory.

SUPPLEMENTARY MATERIAL

See supplementary material for detailed 3D drawings of the laser desorption source and individual components.

ACKNOWLEDGMENTS

We thank Anouk Rijs for helpful discussions regarding the design of our new source, Horst Zink and the DESY FS electronics workshop for expert support with the setup, and Ortwin Hellmig and Andreas Bick for help with the fiber vacuum feedthrough.

In addition to DESY, this work has been supported by the European Research Council under the European Union’s Seventh Framework Programme (FP7/2007-2013) through the Consolidator Grant COMOTION (ERC-614507-Küpper), by the excellence cluster “The
Hamburg Center for Ultrafast Imaging – Structure, Dynamics and Control of Matter at the Atomic Scale” of the Deutsche Forschungsgemeinschaft (CUI, DFG-EXC1074), and by the Helmholtz Gemeinschaft through the “Impuls- und Vernetzungsfond”. We gratefully acknowledge a Kekulé Mobility Fellowship by the Fonds der Chemischen Industrie (FCI) for Nicole Teschmit.

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