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High-Flux Beam Source for Cold, Slow Atoms or Molecules

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We demonstrate and characterize a high-flux beam source for cold, slow atoms or molecules. The desired species is vaporized using laser ablation, then cooled by thermalization in a cryogenic cell of buffer gas. The beam is formed by particles exiting a hole in the buffer gas cell. We characterize the properties of the beam (flux, forward velocity, temperature) for both an atom (Na) and a molecule (PbO) under varying buffer gas density, and discuss conditions for optimizing these beam parameters. Our source compares favorably to existing techniques of beam formation, for a variety of applications.

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Cold, slow beams of atoms or molecules are of wide utility. A common use of such beams is as a source for loading into traps, where the particles can be further cooled and manipulated, e.g., to create Bose condensates or Fermi degenerate gases. Because the number of trapped particles is typically limited by the characteristics of the initial beam (flux, forward velocity, temperature, etc.), significant effort has been put into developing improved atomic beam sources [1–7]. Work on developing cold molecular sources has recently been a particularly active field of research [8–13]. As with atoms, one of the aims is to produce quantum degenerate gases, including those comprising strongly interacting electric dipoles [14–16].

For the purposes of loading traps, an ideal source would produce a large flux of any atom or molecule at temperatures less than the depth of the trap, T_t . For most currently used trap technologies, $T_t \lesssim 1$ K. For species amenable to laser cooling, this temperature is within the capture range of a magneto-optical trap [17]. For paramagnetic atoms and molecules, strong magnetic traps can have depths exceeding 1 K [18]. For polar molecules, electric field-based traps can reach similar depths [19,20].

We describe here a simple, robust source that can operate with nearly any atomic or molecular species, and which produces a beam at high flux with translational and rotational temperatures near 1 K. We demonstrate this source for both an atom (Na) and a polar molecule (PbO), and discuss the different regimes of beam formation. Our analysis makes it possible to estimate and optimize various characteristics of the beam source for general use. We believe this provides an attractive alternative to many beam techniques now in use.

A simple outline of the operation of our source is as follows. Atoms or molecules of the desired species, A , are first vaporized by laser ablation. This produces N particles of A per pulse, at a high temperature T_i . The ablation takes place inside a cryogenic cell filled with He buffer gas at low temperature T_b and density n_{He} . After a characteristic

number of collisions \mathcal{N} , the translational temperature T of A comes arbitrarily close to equilibrium with the buffer gas, such that $T \approx (1 + \epsilon)T_b$ when $\mathcal{N} = -\kappa \ln(\epsilon T_b/T_i)$ [21]. [Here $\kappa \equiv (m_A + m_{\text{He}})^2/(2m_A m_{\text{He}})$, and m_{He} (m_A) is the mass of He (A).] Rotational degrees of freedom are also cooled during these collisions. By allowing both He and A particles to exit the cell via a small hole, a beam of A is formed. The beam persists for a duration given by the diffusion lifetime of A in the cell, which is limited by the sticking of A particles upon contact with the cell walls.

The number of cold particles of A in the beam is determined by both n_{He} and the cell geometry. During thermalization, a particle of A typically travels a distance $R \approx \mathcal{N}/(n_{\text{He}}\sigma_t)$, where σ_t is a thermally averaged cross section for elastic collisions. Hence, for a cell with distance R_h from the ablation point to the hole, the particles of A will be efficiently thermalized before exiting the hole only if $R < R_h$. In addition, the purely geometric probability for a particle of A to escape in the beam is governed by the ratio d/R_h , where d is the diameter of the exit hole.

The forward velocity, v_f , of the thermalized beam of A particles also depends on both n_{He} and d . Specifically, v_f is determined by the ratio d/λ , where $\lambda = 1/(n_{\text{He}}\sigma_c)$ is the mean free path of A particles in the cell; here σ_c is the elastic cross section for cold A -He collisions. In the effusive limit ($\lambda \gg d$), v_f will be given approximately by the thermal velocity of cold A particles, i.e., $v_f \approx v_A \equiv \sqrt{2k_B T_b/m_A}$. By contrast, when $\lambda \ll d$ the A particles will become entrained in the outward flow of He, so that $v_f \approx v_{\text{He}} \equiv \sqrt{2k_B T_b/m_{\text{He}}}$. Since $m_A \gg m_{\text{He}}$ for most species of interest, v_f is much smaller in the effusive limit than for an entrained beam.

Note that the conditions for efficient thermalization and for a slow beam are in conflict. Thermalization is most efficient for n_{He} above a threshold value, but effusive flow demands that n_{He} be less than a typically different threshold. The highest flux of cold, slow A particles is obtained

when R_h and d are chosen so that these thresholds in buffer gas density coincide (namely, when $R_h/d \approx \mathcal{N}\sigma_c/\sigma_t$), and n_{He} is set at this common threshold value ($n_{\text{He}}^{-1} = d\sigma_c = R_h\sigma_t$).

A schematic of the beam setup is given in Fig. 1. The buffer gas cell is a brass box ~ 10 cm on edge. The cell is mounted in vacuum, with the top face attached to the cold plate of a liquid He cryostat. A 0.8 mm long exit hole with $d = 3$ mm is centered on one side face. The bottom and other side faces are covered with windows for optical access. Several ablation targets are mounted on the top face at $R_h \approx 6$ cm. The ablation light consists of laser pulses of ~ 5 ns duration, with energy ~ 15 mJ, focused to a spot size ≤ 1 mm, at a wavelength of 532 nm. The ablation laser is typically fired at 10 Hz repetition rate. We produce Na atoms with sodium metal or NaCl targets, and PbO molecules with a vacuum hot-pressed PbO target. With our ablation conditions, typically $T_i \approx 1000$ K for both species.

Buffer gas continuously flows into the cell through a narrow tube that is thermally anchored to the cold plate. This ensures that the buffer gas is at the temperature of the cold plate upon entry to the cell. The cell walls, and thus the He gas, are typically at $T_b = 5$ K. Helium in the cell must be replenished as it flows continuously through the exit hole. A calibrated metering valve at room temperature is used to control the flow into the cell, and hence the density, of the buffer gas. We determine n_{He} to within a factor of 2. Good vacuum is maintained in the beam region by means of a coconut charcoal sorption pump with a pumping speed of ~ 1000 l s^{-1} .

For a typical elastic cross section $\sigma_c \approx 3 \times 10^{15}$ cm^{-3} , the crossover between effusive and entrained flow of A particles—i.e., the condition $d = \lambda$ —occurs for $n_{\text{He}} = n_c \approx 10^{15}$ cm^{-3} . This should be compared to the density $n_{\text{He}} = n_t$ required for full thermalization of A particles—

i.e., such that $R \approx \mathcal{N}/(n_{\text{He}}\sigma_t) = R_h$. Assuming $\sigma_t \approx \sigma_c$, we find that for Na in our cell, $n_t \approx n_c$. Thus, it should be anticipated that our cell is near the optimal geometry for producing a maximal flux of slow, cold Na. By contrast, the larger mass of PbO makes \mathcal{N} much larger than for Na, implying that our cell geometry is not optimal for PbO. We characterize the beam source for both species within a range of densities around the anticipated optimal condition for Na, namely $n_{\text{He}} \approx (0.2\text{--}5) \times 10^{15}$ cm^{-3} .

The beam source is monitored using laser spectroscopy. Doppler shifts and widths of the spectra are used to determine beam velocity profiles. Signal size and timing yield the column density and particle dynamics. To measure the longitudinal (transverse) velocity profile, a probe laser beam is sent collinear with (perpendicular to) the molecular beam. For Na, a third probe beam monitors atoms inside the cell.

We monitor Na atoms via absorption of a probe laser tuned to the $3s_{1/2} \rightarrow 3p_{3/2}$ transition at wavelength $\lambda_{\text{Na}} = 589$ nm. The probe laser frequency is continuously scanned over the entire absorption profile (a range of ~ 1 GHz), at a rate of 1 kHz, resulting in several complete spectral profiles of the Na atomic beam for each ablation pulse. The relatively slow scan of the laser ensures that we are sensitive only to nearly thermalized Na atoms; hot atoms traverse the detection region before a scan is complete.

PbO is monitored via laser-induced fluorescence. This probe laser is tuned to the $X(v'' = 0) \rightarrow B(v' = 5)$ transition at $\lambda_{\text{PbO},e} = 406$ nm. Fluorescence is detected using a photomultiplier tube (PMT), with interference and colored glass filters to selectively observe the $B(v' = 5) \rightarrow X(v'' = 4)$ transition at $\lambda_{\text{PbO},f} = 460$ nm. A shutter with an opening time of ~ 7 ms is used to shield the PMT from the initial glow following each ablation pulse. Again, this technique ensures that only slow-moving molecules are detected. The signal is averaged over several (typically 10) shots of the ablation laser with the probe laser frequency fixed. A single spectral scan consists of frequency points separated by 10–30 MHz across a span of up to several GHz. Our quantitative determination of molecule number from fluorescence counts includes estimates of the detection efficiency and branching fraction for the detected transition [22]. We estimate the error in our absolute number measurement of PbO to be less than a factor of 2.

For Na, the in-cell probe beam was used to determine both N and σ_c . We find $N_{\text{Na}} \approx 10^{14}$ /pulse for both the Na and NaCl targets. We measure diffusion lifetimes of $\tau[\text{ms}] \approx 4 \times 10^{-15} \times n_{\text{He}}[\text{cm}^{-3}]$. From this we infer $\sigma_c \approx 3 \times 10^{-15}$ cm^2 . For PbO, previous work has measured an ablation yield of $\approx 10^{12}$ /pulse [23], and our measurements with an in-cell probe indicate a comparable yield.

In Fig. 2 we plot the number, N_h , of thermalized particles of species A exiting the hole as a function of n_{He} . We

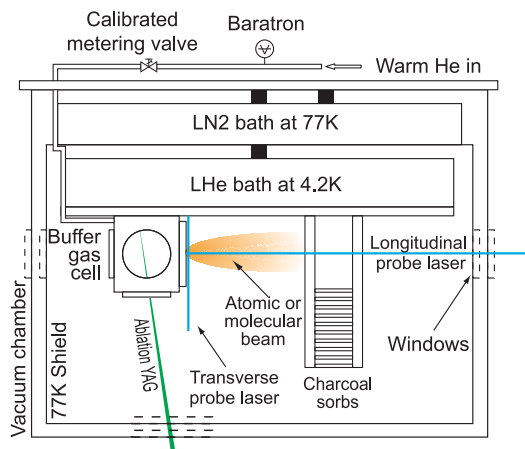


FIG. 1 (color online). A schematic of the beam apparatus. When detecting fluorescence, a lens (not shown) collimates a fraction of the fluorescence light and directs it out a window to a PMT.

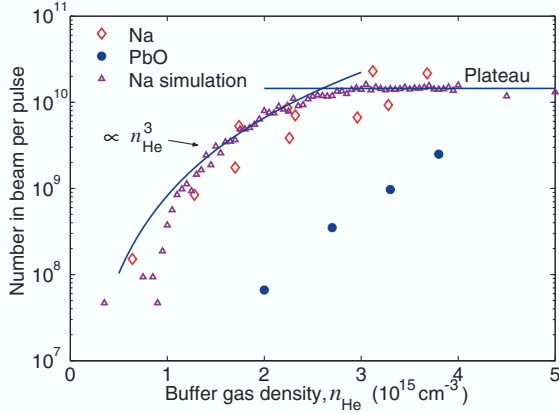


FIG. 2 (color online). The number of cold A particles, N_h , that emerge in the beam as a function of n_{He} . Curves with specific functional forms have been inserted to show the different scaling regimes. The intersection of the curves corresponds to the condition $R = R_h$, where a typical A particle is thermalized just as it reaches the hole.

also show the results of a Monte Carlo simulation of the beam formation process. The condition for full thermalization is apparent in both the experimental and the simulated data for Na, while as expected we do not appear to reach the condition of full thermalization for PbO.

In the Na data and simulations, we find that N_h increases rapidly (approximately $\propto n_{\text{He}}^3$) up to a critical value of n_{He} , above which N_h is roughly constant at its maximum value $N_{h,\text{max}}$. The low-density scaling is consistent with a simple picture in which A particles are distributed uniformly over a volume of characteristic length $L_t \approx \mathcal{N}\lambda$, when they have thermalized to near T_b . (This broad distribution arises, in our simulations of the thermalization process, from the spreads both in the number of collisions required to thermalize and in the free path between collisions.) At high density, the condition $N_h \approx N_{h,\text{max}}$ arises in our simulations when $R < R_h$ for essentially all A particles. In this regime, the fraction of A particles escaping, $f_{\text{max}} = N_{h,\text{max}}/N$, is given roughly by the area of the hole to the area of a hemisphere at radius R_h , i.e., $f_{\text{max}} \approx d_h^2/(8R_h^2)$. In our geometry, $f_{\text{max}} \approx 3 \times 10^{-4}$.

The simulated beam data in Fig. 2 are scaled to match the experimental Na data by adjusting the values of N_{Na} and σ_t . The resulting value, $N_{\text{Na}} \approx 5 \times 10^{13}$, is in reasonable agreement with the determination from the in-cell probe. The fitted thermalization cross section, $\sigma_t \sim 1 \times 10^{-15} \text{ cm}^2$, is somewhat smaller than σ_c . This is reasonable, since elastic cross sections are typically smaller at higher collision energies [24].

Figure 3 shows the average forward velocity, v_f , of the beams of A particles as n_{He} is varied. For both Na and PbO, the data show a nearly linear increase of v_f with n_{He} , with the velocity of the lighter species always larger. This behavior is consistent with the following simple picture.

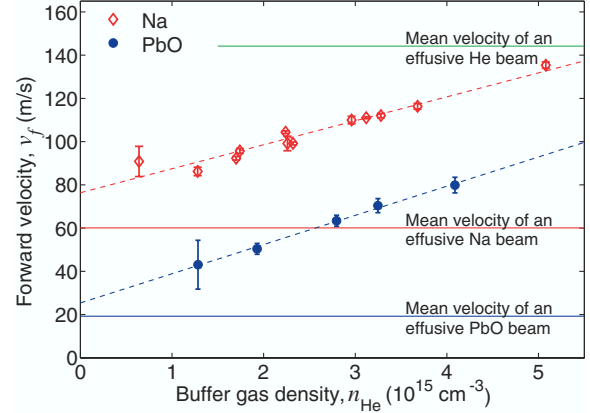


FIG. 3 (color online). PbO and Na beam mean forward velocities, v_f , as a function of n_{He} . Extrapolation of the data to zero buffer gas density is illustrated by best-fit lines (dashed lines).

A slowly moving particle of A takes a time T_e to exit the hole, where $T_e \sim d/v_A$. During this time, it undergoes N_e collisions with fast, primarily forward-moving He atoms, where $N_e \sim n_{\text{He}}\sigma_c v_{\text{He}} T_e$. Each collision imparts a momentum transfer $\Delta p_A \sim m_{\text{He}}\sigma_c v_{\text{He}} T_e$. This results in a net velocity boost Δv_A , given by $\Delta v_A \sim v_A d/\lambda \propto n_{\text{He}}$. This picture should be roughly valid for densities below the regime of full entrainment, where $v_f \sim v_{\text{He}}$. The velocities we measure for Na are approximately reproduced by modeling of the beam formation process with our measured value of σ_c .

This picture also predicts that the behavior of v_f , when extrapolated to $n_{\text{He}} = 0$, should yield the velocity of an effusive beam of A particles at temperature T_b . To make this comparison, it is critical to note that our detection technique is sensitive to molecules within a roughly cylindrical volume, of diameter D_d , and with length L extending from the exit hole. Under our conditions, where $D_d \sim d < L$, it can be shown that an effusive beam will exhibit a velocity distribution close to $f(v) \propto v^2 e^{-m_A v^2/(2k_B T_b)}$ and a mean velocity $\bar{v}_{\text{eff}} \approx 1.13v_A$. Our extrapolated data are within $\sim 25\%$ of this prediction for both species.

Figure 4 shows the temperature of the beams vs n_{He} , as measured by the velocity spreads in the longitudinal and transverse directions, as well as the rotational population distribution (for PbO). For transverse temperature we fit to a distribution of the form $f_t(v) \propto e^{-m_A v^2/(2k_B T)}$. For longitudinal temperature we use $f_l(v) \propto e^{-m_A (v-v_l)^2/(2k_B T)}$. Fits of the longitudinal data to the effusive distribution $f_{l,\text{eff}}(v) \propto v^2 e^{-m_s v^2/(2k_B T)}$ were poor, consistent with the partial entrainment of A in the helium flow. For the rotational temperature, we use Clebsch-Gordan and Hönl-London factors to determine the relationship between fluorescence intensity and initial state population for various rotational lines. The ratio of initial state populations for 2–3 rotational lines determine the temperature corresponding to a Boltzmann distribution. Our data indicate complete

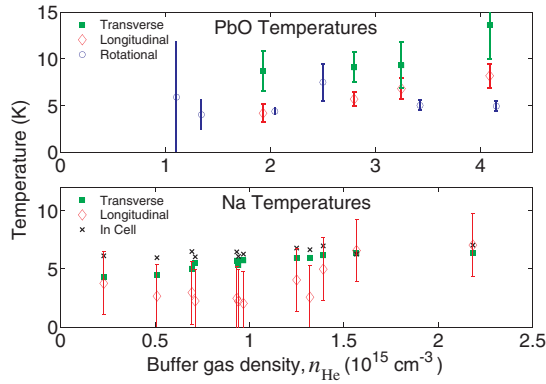


FIG. 4 (color online). PbO and Na temperatures as a function of n_{He} as determined by fits to spectral profiles. “Transverse,” “longitudinal,” and “in cell” refer to the location of the probe laser (see Fig. 1). Typical errors for the transverse and in cell Na temperatures are comparable to the marker size.

thermalization of all detected particles. Note that we observe no additional cooling below T_b , as would be expected for entrainment in a fully supersonic He flow.

The beam source described here can be readily adapted to the needs of a wide range of experiments. For example, it can be used for a wide variety of species with performance similar to that described here. The total flux depends linearly on the ablation yield N , which for any given target is difficult to predict *a priori*. However, for nearly every species we have tried (here and in many related experiments), it has been possible to achieve large values of N by a suitable choice of precursor material. For example, under operating conditions similar to those used here, we have obtained $N = 10^{12}$ – 10^{14} for a variety of metal atoms and $N = 10^{11}$ – 10^{13} for many species of diatomic molecules (including radicals) [25–28]. In addition—subject to limitations of cooling power or gas load—the source could be run at lower temperatures, higher repetition rates, or with higher extraction efficiency (e.g., by using several separated exit holes). Our discussion of the beam formation mechanism makes it straightforward to determine the effects of such changes.

A key possible improvement to the source would be the addition of a guide, either magnetic (for paramagnetic species) or electric (for polar molecules). In both cases, He is unaffected by the guide potential and will exit through the sides of the guide, allowing extraction of the A beam into a region of ultrahigh vacuum. Under the conditions described here, this beam source could be used to load a peak flux of up to $2 \times 10^{11} \text{ s}^{-1}$ Na atoms into a simple permanent magnet guide such as the type described in Ref. [7]. An electrostatic guide such as that

described in Ref. [12] could be loaded with a peak flux of $\sim 10^9 \text{ s}^{-1}$ PbO molecules in the $J = 1$ rotational state, adequate for loading into a microwave [19] or electrostatic [29] trap. The high fluxes from our source could result in substantial improvements in atomic and molecular trapping experiments that depend on large initial numbers.

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