



Separation and Detection of Charged Unilamellar Vesicles in Vacuum by a Frequency-Controlled Quadrupole Mass Sensor

Anatolii Spesyvyi,* Marek Cebecauer, Ján Žabka, Agnieszka Olżyńska, Michaela Malečková, Zuzana Johanovská, Miroslav Polášek, Ales Charvat, and Bernd Abel



Cite This: Anal. Chem. 2025, 97, 9131-9138



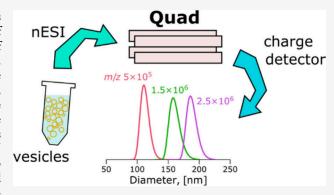
ACCESS I

III Metrics & More

Article Recommendations

Supporting Information

ABSTRACT: Extracellular vesicles (EVs) are membranous particles released by cells and are considered to be promising sources of biomarkers for various diseases. Mass spectrometry (MS) analysis of EVs requires a sample of purified and detergent-lysed EVs. Purification of EVs is laborious, based on size, density, or surface nature, and requires large amounts of the source material (e.g., blood, spinal fluid). We have employed synthetically produced large unilamellar lipid vesicles (LUVs) as analogs of EVs to demonstrate an alternative approach to vesicle separation for subsequent mass spectrometry analysis of their composition. Mass-to-charge ratio m/z separation by frequency-controlled quadrupole was employed to filter narrow-size distributions of LUVs from a water sample. Lipid vesicles were positively charged with nanoelectrospray and trans-



ferred into a vacuum using two wide m/z-range frequency-controlled quadrupoles. The m/z, charges, and masses of individual vesicles were obtained by the nondestructive single-pass charge detector. The resolving mode of the second quadrupole with m/zRSD < 10% allowed to separate size selected distributions of vesicles with modal diameters of 88, 112, 130, 162, and 190 nm at corresponding quadrupole m/z settings of 2.5×10^5 , 5×10^5 , 8×10^5 , 1.5×10^6 , and 2.5×10^6 , respectively with a rate of 20-100counts per minute. The distributions of bioparticles with masses between 108 and 1010 Da were separated from human blood serum in the pilot experiment. The presented approach for lipid vesicle separation encourages the development of new techniques for the direct mass-spectrometric analysis of biomarkers in MS-separated EVs in a vacuum.

xtracellular vesicles (EVs) are membranous particles released by diverse cell types. EVs participate in intercellular transport and communication.² Their cargo may include proteins, nucleic acids, lipids, and cell metabolites, which can be valuable biomarkers of human diseases or metabolic changes.^{3,4} Mass spectrometry (MS) analysis of the EVs' cargo composition requires isolated and lysed (membrane disintegration) vesicles.⁵ The EVs' isolation from biological samples (e.g., blood plasma, urine) is laborious and timeconsuming and includes ultracentrifugation, filtering, sizeexclusion chromatography, or affinity-based purification on functionalized beads. Established separation techniques usually require larger amounts of material, which can be limited in samples from children or in the case of cerebrospinal fluid. The size of isolated vesicles is then characterized by dynamic light scattering, nanoparticle tracking analysis, flow cytometry, or transmission electron microscopy, which further reduces the yield of EV separation. The size of the analyzed EVs is a critical parameter. There is a large variety of EVs produced by cells, and those within a smaller range (50-300 nm in diameter) are considered more informative about the cell state or the local environment than larger EVs.8 Large unilamellar vesicles (LUVs), i.e., synthetic liposomes of size

100-1000 nm with a single outer lipid bilayer and enclosed aqueous core, are an established model system of EVs, particularly because of size and shape similarity and the fact they mimic mechanical and structural properties of the membrane and its functionalization with proteins. 9-11 1palmitoyl-2-oleoyl-glycero-3-phosphocholine used for synthetic vesicle preparation for this study is a common lipid mimicking cellular membrane in model systems. 12 The characterization of charged liposomes based on their gasphase mobility was introduced as a nanoelectrospray gas-phase electrophoretic mobility molecular analyzer (nES GEMMA) and was applied as a separation stage for Raman and midinfrared spectroscopy. 13

Recently, joint efforts of Jarrold and Clemmer groups at Indiana University have yielded a new approach to studying

Received: October 24, 2024 Revised: April 4, 2025 Accepted: April 12, 2025 Published: April 22, 2025





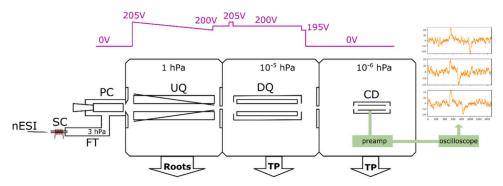


Figure 1. SELINA schematic representation with the main components denoted: nanoelectrospray emitter with the sample (nESI), 20 mm sampling capillary with resistive heating (SC), flow tube (FT), plenum chamber (PC), upstream quadrupole with slanted wires (UQ), downstream quadrupole (DQ), charge detector (CD), turbomolecular pumps (TP). The electric potential levels are shown at the top, and the exemplary signals from the CD, recorded by an oscilloscope, are on the right; the x-axis for signals is time in μ s, and the y-axis is voltage in mV.

intact EVs (i.e., EVs with structurally undamaged membranes). 14,15 In particular, the intact EVs were characterized using charge detection mass spectrometry (CDMS) with nanoelectrospray. 14 The precisely measured charge and mass-to-charge ratio m/z of the EVs transported into the vacuum and characterized by the charge detector provided exact mass and charge distributions, with the last one reflecting structural and surface information. CDMS enabled distinguishing keratinocytic exosomes in diabetic and nondiabetic mice based on the charge distribution subpopulations retrieved by a Gaussian mixture model. 15 The CDMS instrument used in these studies consisted of multipole nonresolving ion guides, acting simultaneously as a high-pass filter to transfer charged particles generated in nanoelectrospray from atmospheric pressure toward a multipass charge detector in vacuum. 14,16

In other investigations, charged particles with $m/z > 10^5$ were also analyzed using electrodynamic traps, including several Paul's designs¹⁷ and linear quadrupole traps.¹⁸⁻ The former were shown to be suitable for the analyses of highmass microparticles, e.g., polystyrene particles and cell particles.^{21–24} It is noteworthy that MS using transmission linear quadrupoles is generally limited to $m/z < 10^5$ with usually subunit resolution.²⁵ The resolving power can reach thousands, albeit at the cost of ion transmission. For even higher m/z, the quadrupole driver electronics must feature extraordinary specifications, and a few special approaches to the quadrupole operation were introduced, i.e., the frequencyswept,²⁶ frequency and amplitude scanned,²⁷ and digitally driven^{28–30} modes. All rely on the frequency variation of the frequency and amplitude scanned, 27 and digitally periodic waveform applied to pairs of quadrupole rods. The 'frequency scanned' mode utilizes a constant or semiconstant ratio of AC and DC components at a selected frequency to establish a stability zone for a certain m/z. In the digitally driven mode, the stability zone is defined by varied frequency and duty cycles of square waveforms without a DC component. In the case of linear ion traps, frequency of an axial or radial excitation periodic potential may be scanned to resonantly eject ions with different m/z.^{31,32}

Here, we use the term 'frequency-controlled' because for the present application only the frequency of AC voltage is changed, and the resolution is defined by the ratio of constant AC and DC components. While quadrupoles are used in this work at fixed m/z settings, a frequency scan is necessary to obtain a mass spectrum.

The Selected Ice Nanoparticle Accelerator (SELINA) instrument was originally designed to produce mass-selected

beams of highly charged water ice nanoparticles with 50–1000 nm diameter mimicking space dust found in the vicinity of several icy moons in our Solar system. Impact mass spectrometry of such particles accelerated electrostatically to hypervelocity is supposed to be the best terrestrial analogue experiment for past and future space missions with impact ionization MS instrumentation onboard. Moreover, it is the first step toward developing novel, high-resolution space probes based on impact ionization MS.³³ The SELINA design is also suitable for the production and mass measurement of submicrometer solid particles from liquid samples nebulized by electrospray. Special care must be taken to fully desolve the particles. This procedure was successfully carried out for 300 and 600 nm polystyrene beads during the charge detector validation experiments.³⁴

In this Technical Note, we showcase the operation of the SELINA apparatus when the resolving mode of the frequencycontrolled quadrupole in the range of m/z 10^5-10^6 is used to identify different mass (size) of LUV fractions upon electrospraying a liquid sample of LUVs with a broad distribution. After quadrupole separation, positively charged LUVs were detected using a single-pass charge detector, which provided m/z, charge, and mass of an individual vesicle. Finally, in a proof-of-concept experiment, the technique was adopted to investigate a sample of the human blood serum, aiming to assess the underlying mass distribution fractions of bioparticles from the present heterogeneous sample. Considering the current advances in CDMS technology, i.e., electrostatic ion trap designs with a single elementary charge detection limit and superior resolution of m/z, charge, and mass, we deliberately use the 'quadrupole mass sensor' name for our setup to underline its application as a technique primarily used for vesicles separation from a sample, not for their thorough characterization, which could be otherwise possible by application of the state-of-the-art CDMS.

EXPERIMENTAL SECTION

SELINA Instrument. A detailed description of SELINA (shown schematically in Figure 1) has been published previously.³³ Briefly, charged particles from a nanoelectrospray pass through a 20 mm long sampling capillary (SC) heated to approximately 200 °C for vesicle desolvation and continue moving through an aerodynamic system consisting of a flow tube with a plenum chamber (PC) by gradually losing their initial momentum in collisions with gas molecules. The particles are subsequently dragged by the jet of air into a

first upstream quadrupole (UQ). This quadrupole is equipped with slanted wires between rods to impose an electric drift field $(\sim 0.4 \text{ V cm}^{-1})$ at a pressure of 1 hPa of lab air flowing through the PC. The charged particles are further thermalized in collisions with air molecules and drift in the UQ toward an orifice with a diameter of 1 mm into a differentially pumped chamber (5 \times 10⁻⁵ hPa), which accommodates a linear quadrupole trap operated as a downstream quadrupole filter (DQ). The third differentially pumped chamber contains a single-pass charge detector (CD) to measure the particle's charge and m/z. The charged particles leave the DQ through a 2 mm ID orifice and fly about 5 cm toward the 2 mm ID differential pumping orifice into the charge detector vacuum chamber and then an additional 5 cm toward the 2 mm ID entrance orifice into the charge detector. The potentials set were 205 V for the UQ, 195 V for the slanted wires, 205 V for the entrance DQ electrode, 200 V for the DQ and its enclosure, and 195 V for the exit DQ electrode.

Charge Detector. The same CD as in ref 33 and ref 34 was utilized here, with the previously done charge calibration being 38 elementary charges (e) for 1 mV of the entrance peak amplitude of preamplifier response transient output for square pulse input. The procedure and corrections related to the shape of the transient signal and particle velocity are described in detail in ref 33. Briefly, the signal with amplitudes down to a few millivolts was pulsed through the calibration capacitor (0.8 pF), resulting in a minimum accessible injected charge of 20000 e. The linear dependence of the CD response on the injected charge is assumed valid down to the lower inaccessible values. This 'low charge' region was verified by polystyrene 300 nm beads measurement, where detected charges were in the range of 1000-5000 e. The CD electronics placed in a vacuum amplifies and shapes an image charge induced by a particle flying through a 50 mm long charge pick-up stainless steel tube. The resulting transients are recorded by an oscilloscope (PicoScope5000, Pico Technology, UK) at 16-bit resolution and sent to PC software for processing and storage. For vesicle measurement, the trigger was set to 20 mV, and the sampling rate and range were 50 kS/s and 1600 µs. Processed transients allow assessment of particle velocity (detector length divided by the time difference between entrance and exit peaks) and charge (average of amplitudes of entrance and exit peaks multiplied by 38 e/mV and velocity correction coefficient) and thus calculation of m/z (from known acceleration potential and velocity) and mass (as product of charge and m/z). An acceleration potential was defined by the potential of the DQ and was kept constant at 200 V.

The CD charge measurement error obtained from the blank signal is 328 e (see details in Figure S1), while the noise root-mean-square (RMS) is 131 e. Thus, for a particle with charge 600 e and m/z 2.5 × 10^5 , the relative standard deviation (RSD) is 66% for mass and 22% for diameter using charge measurement error 328 e. When noise is used, RSD values are 38% and 13%, respectively. The limit of charge detection of used CD is approximately 600 e, which is considerably higher than the usual value of 250 e for single-pass detectors. This is not an issue for the measurement of highly charged ice nanoparticles in SELINA, but just at the detection limit for vesicles.

Quadrupoles. The mechanical assemblies of the ionguiding UQ and the mass-resolving DQ were repurposed from commercial mass spectrometers (Thermo Scientific). The UQ has 250 mm long hyperbolic rods with a 6 mm field radius

(distance from the axis to the rod), and the DQ is a part of the linear quadrupole trap assembly, which has 60 mm long hyperbolic rods and a 4.75 mm field radius. The DQ is enclosed and has entrance and exit electrodes. Its inner volume is filled at a flow rate of 1 sccm with lab air, which results in a pressure of 10^{-3} – 10^{-2} hPa in the enclosure. Both quadrupoles are powered by two dedicated voltage sources (Janas Card, Czechia) that can float at a potential of up to ± 1 kV. A direct digital synthesis chip is used to generate a harmonic signal with 0.03725 Hz resolution. The choice of harmonic potential simplifies the design of the sources and is advantageous in that m/z settings are calculated for the first stability region of the Mathieu diagram. The sources have slightly different operational parameters. The UQ source provides harmonic voltages in the 1-300 kHz range with a constant amplitude of 50 or 100 V, ensured via the automatic gain control loop. The DQ source operates in the 0.5-200 kHz with a constant nominal amplitude of 100 or 200 V corrected with the internal calibration table of peak-to-peak amplitudes depending on the frequency from 200.594 V at 500 Hz to 206.099 V at 200 kHz for the nominal amplitude 100 V, and from 401.862 V at 500 Hz to 400.437 V at 100 kHz for the nominal amplitude 200 V. The calibration table is sufficient for the quadrupole application presented here. However, high-precision measurement of the peak-to-peak voltage at each frequency would be necessary to obtain an exact m/z value of transmitted ions.³⁶ Both sources provide variable DC components of 0 to \pm 35 V with 16-bit resolution. The voltage parameters for certain m/zare calculated by a software (see below) accounting for the calibration table if necessary and sent to the voltage sources via USB. The m/z settings accessible for the present combination of quadrupole physical dimensions and voltage source characteristics allow to set m/z 1 × 10³-4 × 10⁷ for UQ and $m/z \times 10^3 - 4 \times 10^8$ for DQ. The quadrupoles were used in two modes of operation: the nonresolving one (DC component for UQ and DQ was 0 V) and the resolving one (DC component for UQ was 0 V and \pm 30.07 for DQ). For the sake of clarity, we will further mention only the m/zsettings of DQ (m/z_{DO}) , assuming that the UQ is always set to the same m/z with the DC component equal to 0 V. The DC component of the DQ for the resolving mode is calculated from the apex of the first stability region of the Mathieu diagram, being ~0.168 of the AC amplitude and taken at the 90% percent tune. The voltage and frequency values for all m/zsettings used are provided in Table S1 in the Supporting Information.

LUVs Preparation. The suspension of LUVs was prepared as described below. The 100 μ L aliquot of chloroform (Merck, Darmstadt, Germany) solution of POPC (1-palmitoyl-2-oleyl-sn-glycero-3-phosphocholine; Avanti Polar Lipids, Alabaster, AL) was transferred to a glass test tube. The solvent was evaporated under a stream of nitrogen, and the lipid film was kept under a vacuum overnight. The dry lipid film was hydrated with Mili-Q water (Millipore, USA). After 4 min of continuous vortexing, the suspension of multilamellar vesicles (MLVs) was extruded with a mini extruder (Avestin, Ottawa, Canada) through a polycarbonate membrane (Whatman; Little Chalfont, UK) with a nominal pore diameter of 200 nm. The lipid concentration was 1 mM, and the LUVs suspension was used at different dilutions.

The size distribution of LUVs suspension was measured using dynamic light scattering (DLS) setup Zetasizer Nano ZS (Malvern Instruments, Worcestershire, UK) consisting of a

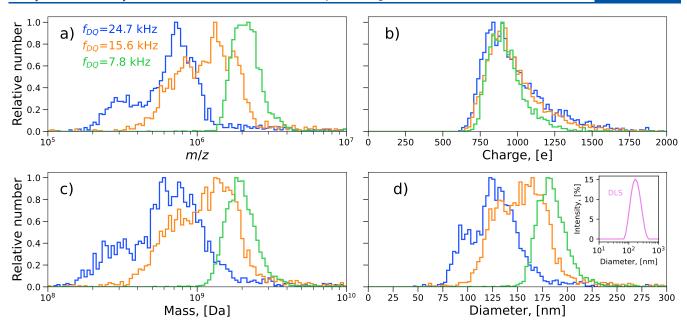


Figure 2. Histogram distributions of m/z (a), charge (b), mass (c), and diameter (d) of vesicles as measured by the charge detector at three different frequencies (color coded) of DQ in the nonresolving mode. The DLS measurement is plotted in panel (d) inset for comparison. The numbers of CD detection events used to plot histogram distributions are 2454, 2585, and 3141 for f_{DQ} values of 24.7, 15.6, and 7.8 kHz, respectively; bin number in each histogram is 100 with equal widths (b, d) and varying widths spaced evenly on a log scale (a, c).

He—Ne laser (633 nm) and an avalanche photodiode detector (APD). To get the optimal scattering intensity, the sample was diluted to the 0.2 mM final lipid concentration, transferred to a plastic disposable cuvette (Brand, Wertheim, Germany), and measured after 2 min of equilibration time at 298 K. Scattered light was collected at the angle of 173° and intensity-weighted size distribution was obtained using Zetasizer Software 7.11.

Nanoelectrospray Emitters Preparation and Vesicles Measurement. Nanoelectrospray emitters were pulled with the P-2000 laser-based micropipette puller (Sutter Instrument, CA) from the 10 cm length 1 mm OD, 0.78 mm ID borosilicate glass with filament (Sutter Instrument, BF100-78–10). The puller settings resulted mostly in a 4 mm taper with a closed tip. After the sample was loaded, the emitter was conditioned by gently touching the metal surface of the sampling capillary (SC) until the signal could be registered and no corona discharge was visible at a 1.5-2 mm distance from the SC. After the experiments, the emitters were inspected with a light microscope (Olympus FluoView 1000 MPE), as shown in Figure S2. The tip ID varied from 3 to 20 μ m. The electrostatic potential was supplied to the platinum wire (0.05 mm diameter) within the emitter that was filled with the sample liquid. The emitters were first loaded with 300 nm latex beads in 100 mM ammonium acetate solution to find the suitable operating parameters of the nanoelectrospray (i.e., electric potential, distance toward SC, desolvation temperature, and the tip diameter) and to get rid of the solvent nanodroplets (larger than 50 nm in diameter). For 300 nm particles, the following parameters were found: the distance between the emitter and SC of 1.5-2 mm, the emitter potential of +1.2 kV and the SC temperature of 200 °C. Vesicle samples were then measured at identical conditions. After the latex beads sample was consumed, the sample with vesicles was loaded with a syringe into the emitter without changing its position and sprayed at the same potential. In this manner, polystyrene beads in a 100 mM ammonium acetate solution sample acted as a negative control, ensuring the

absence of detectable solvent droplets at the defined nanoelectrospray conditions. The measurement procedure was repeated for three different emitters with freshly prepared vesicle samples each time (200 mM ammonium acetate aqueous solution and original vesicle sample 1:1 by volume). The transients recorded from the charge detector at different quadrupole settings were processed using Python with Numpy,³⁷ Scipy,³⁸ and Matplotlib³⁹ libraries, and details are provided in the caption of Figure S4. Postprocessing was necessary to exclude invalid transients (e.g., when signals of two or more particles overlapped). While in the nonresolving mode with a high count rate, 49% of the transients had to be discarded, only 14% of transients were found invalid in the resolving mode having substantially lower count rates (see Table S2 for details). Processed results for all three emittersample couples are also provided in the SI (Figures S6-S9). Since the resulting m/z, charge, mass, and diameter distributions belonging to the different emitters did not differ substantially, we focus on the data acquired with the second emitter sample in the next section.

Blood Serum Measurement. The 6 mL of blood was sampled from the veins of the volunteer into a blood-collection tube containing sodium citrate (final concentration of 3.8% w/V). The sample was left to separate erythrocytes (red blood cells) and platelets from serum by standing still at room temperature for 3 h. The upper yellow-brown supernatant was transferred into a new 15 mL tube with a conical bottom and centrifuged for 10 min at 1500g at room temperature to remove debris and white blood cells. The supernatant (blood serum) was transferred into 3 new 1.5 mL Eppendorf tubes and stored in a fridge at +8 °C for 5 days in a vertical position.

The 20 μ L of serum was taken from the upper half of the vial and mixed with 800 μ L of 100 mM ammonium acetate solution. The emitter distance from SC was about 2 mm and nanoelectrospray potential +1.3 kV. The size distribution of the blood serum sample was measured using the same DLS setup and experimental conditions as those used for the LUVs

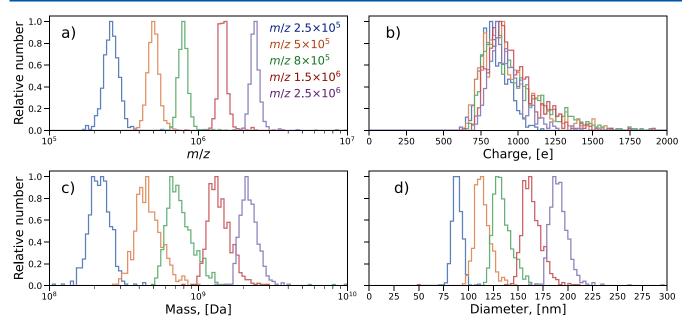


Figure 3. Histogram distributions of m/z (a), charge (b), mass (c), and diameter (d) of vesicles as determined by the charge detector with the DQ quadrupole in a resolving mode at five different m/z settings (color-coded). Numbers of CD detection events used to plot histogram distributions are 646, 705, 897, 748, and 503 for $m/z_{\rm DQ}$ 2.5 × 10⁵, 5 × 10⁵, 8 × 10⁵, 1.5 × 10⁶, and 2.5 × 10⁶ respectively; bin number in each histogram is 100 with equal widths (b, d) and varying widths spaced evenly on a log scale (a, c).

suspension. Before being transferred to a plastic disposable cuvette, the blood serum was diluted with water in a 1:1 (v/v) ratio.

RESULTS AND DISCUSSION

LUVs Sample. After loading the emitter with the vesicle sample, the DQ in nonresolving mode was used to transfer all charged vesicles from nanoelectrospray toward the charge detector. To transmit the full range of LUVs with initially unknown m/z distribution, the DQ frequencies were set to 24.7, 15.6, and 7.8 kHz. These values correspond to $m/z_{\rm DO}$ settings 2×10^5 , 5×10^5 , and 2×10^6 with 0 DC component, respectively, and comply with the lower m/z limit in the stability diagram. Each frequency setting was repeated in cycles with 30 s dwell time for 40 min to collect a sufficient number of detection events while keeping enough amount of the sample in the emitter for the subsequent measurements in the resolving mode. Upon passage through the CD, the charged vesicle generates a transient signal. Once digitized, the signal allows us to obtain the charge and the velocity; the latter provides m/z for a known accelerating potential (i.e., DQ potential of 200 V). The mass was calculated as a product of the mass-to-charge ratio and the charge; the diameter of the vesicle was derived from the mass, assuming the spherical shape of LUVs and a density of water (997 kg/m³). All experimental and derived values of m/z, charge, mass, and diameter at each nonresolving DQ setting are plotted in Figure 2 as histogram distributions. Figure 2a shows that the vesicle m/z distributions roughly span the range from 2×10^5 to 3×10^5 10⁶. Because of the close similarity of the three charge distributions (Figure 2b), the shape of the corresponding mass distributions resembles that of m/z (Figure 2c). In comparison with the DLS measurement (inset in Figure 2d), the diameter distributions are slightly shifted to lower values, but most abundant diameters are in the same range between 100 and 200 nm.

To acquire separate narrower mass distributions of LUVs, the DQ was used in resolving mode, and five specific m/z_{DO} were selected from the range obtained in the nonresolving mode. The resulting m/z, charge, mass, and diameter distributions are shown in Figure 3. Again, due to the similarity of charge distributions (Figure 3b), the mass distributions (Figure 3c) are determined primarily by that of m/z (Figure 3a). The m/z resolving power calculated as a ratio of the m/z peak position and its Full Width at Half Maximum (fwhm) is about 5 and did not change substantially at higher DQ resolution settings (see Figure.S10). This means, in turn, that the peak fwhm is defined not by the quadrupole but by the error of the m/z measurement with the charge detector. The relative standard deviation (RSD) can be estimated from Gaussian fits to m/z distributions, and it ranges from 12% for $m/z \ 2.5 \times 10^5$ to 7% for $m/z \ 2.5 \times 10^6$.

The charge distributions at different $m/z_{\rm DQ}$ settings are generally similar, with their modal values in the range of 750-900 e, and various tailings toward higher charges. The detection limit of the present CD is about 600 e, as mentioned in the Experimental section, and careful signal processing was required, especially when several charged particles were recorded simultaneously in one transient (see Figure S4). Additionally, heatmaps of the charge versus mass dependence are shown in Figure S12 for both nonresolving and resolving modes of operation. All charges measured are smaller than the Rayleigh charge limit for water, and their number does not show any substantial increase with the diameter (mass). On the other hand, distinct charge distributions may indeed be observed with CDMS. This was the case in the studies of lipoproteins and exosomes, where different particle structures, geometries, and molecular cargos located on the surface resulted in several distinct charge subpopulations. 15,40 The reduction of a charge of electrosprayed spherical-shaped bacterium Staphylococcus aureus (500-1000 nm) with increasing sampling capillary temperature was explored by Shao-Yu Liang et al. using a charge-sensing particle detector.

They discovered that at 200 °C, the bacterium was fully desolvated from methanol and bears a mean charge of 1046 e, which is compatible with values obtained here.

Figure 4 shows the time series of the vesicle detection count rate in the resolving mode for five m/z_{DO} settings. Though a

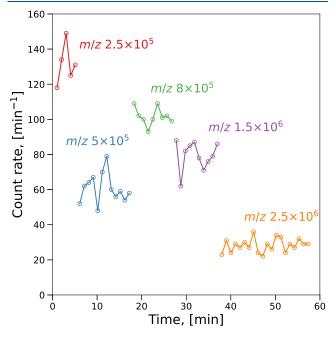


Figure 4. Measurement time series of the vesicle count rate as detected by the charge detector at various m/z settings of DQ in resolving mode.

gas-filled DQ focuses charged vesicles on the instrument axis, some divergence is expected downstream so that the number of detection events is lower compared to the number of vesicles leaving the DQ. Assuming an average flow rate in the nanoelectrospray being 50 nL min⁻¹, we can roughly estimate the total number of vesicles emitted in front of the SC as 10⁸ per minute (see SI for the calculation details). It is, however, not clear how many of them were sampled into SELINA.

Blood Serum Sample. The measurement of the blood serum was done in the same manner as that for the LUVs samples. First, the DQ was used in the nonresolving mode to get broad m/z distributions of charged particles at $f_{\rm DQ}$ values of 24.7 kHz, 15.6 kHz, and 7.8 kHz. The m/z values of detected particles during these initial measurements with different emitters were roughly in the 10^5-10^7 range (see Figure S14). The detection of particles with $m/z > 10^6$ at $f_{\rm DQ} = 7.8$ kHz was unstable and occurred in short bursts, probably affected by some processes in the nanoelectrospray ion source. Unlike LUVs, the DLS analysis of the serum sample shows a much wider, multimodal size distribution with diameters from 4 nm to 6 μ m featuring three peaks around 10, 40, and 500 nm (see Figure S13).

In the resolving mode, DQ was set to six different values of $m/z_{\rm DQ}$ (i.e., 2×10^5 , 3×10^5 , 5.5×10^5 , 7×10^5 , 1.2×10^6 , 3×10^6) based on m/z distributions obtained in the nonresolving mode. While Figure 5 shows histograms for m/z and mass obtained with the CD at four $m/z_{\rm DQ}$ values (i.e., 2×10^5 , 5.5×10^5 , 1.2×10^6 , 3×10^6), all of them are depicted in Figure S15. The diameter distributions are not evaluated here, as the nature of the detected particles and thus their shape and densities are not known. Instead masses of spherical water

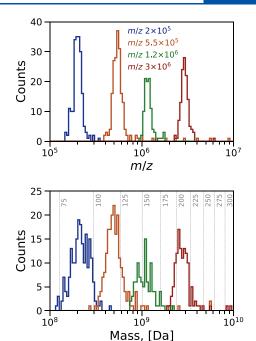


Figure 5. Histogram distributions of m/z (top) and mass (bottom) for serum sample particles as detected by CD at $m/z_{\rm DQ}$ 2×10^5 , 5.5×10^5 , 1.2×10^6 , 3×10^6 . In the bottom plot, the gray vertical lines denote mass values equivalent to spherical particles with a density of 997 kg m⁻³ for different diameters in nanometers. The bin number in each histogram is 100, with varying widths spaced evenly on a log scale.

particles with various diameters are depicted in Figure 5 for the sake of comparison. The count rates in the resolving mode were lower than in the LUVs experiment and covered the range from 6 min $^{-1}$ at $m/z_{\rm DQ}$ 3 \times 10 6 to 30 min $^{-1}$ at $m/z_{\rm DQ}$ 2 \times 10 5 . We also observed that after about 30 min, a visible film was formed near the orifice of the sampling capillary and partial clogging of the latter started after an even longer period of operation.

CONCLUSIONS AND OUTLOOK

We have demonstrated that LUVs with a 100-200 nm diameter can be charged in nanoelectrospray from aqueous solutions, transferred into the vacuum, and thermalized to a defined electric potential in the SELINA instrument. The combination of two wide m/z-range frequency-controlled quadrupoles, where the first one was used in nonresolving mode while the second one in resolving mode, facilitates the mass filtering of vesicles based on their m/z. As a result, we have succeeded in separating a broad initial distribution (fwhm of 200 nm, see Figure 2d) into distinct LUVs fractions having fwhm of 20-30 nm. The charged LUVs were mass characterized by a single-pass charge detector operated in the current study near its detection limit of 600 e. In future investigations, we will apply an improved design of the charge detector based on electrostatic trapping with a detection limit down to tens of elementary charges. 42 It will enhance the precision of the mass measurements and allow us, for instance, to investigate smaller vesicles.

It is well established that real biological samples containing EVs feature a high degree of particle (vesicle) heterogeneity. Therefore, both m/z and the mass correspondence may not be unique due to the broad charge and density distributions of the

bioparticles under study. Nevertheless, quadrupole preseparation performed in this work can be used to select a relatively narrow mass fraction for a detailed mass spectrometric characterization of the content carried by the selected vesicles. If needed, a deflection stage synchronized with the charge detector can be added to single out particles with a desired mass or charge as it was applied in dust particle accelerators.⁴³

The present pilot investigation with blood serum demonstrated that quadrupole-based mass separation of particles from a highly heterogeneous sample is indeed feasible. We found out that separation efficiency was not compromised due to the broader charge distribution of the blood serum sample with respect to LUVs. Currently, the main limitations are the stability of the nanoelectrospray source and a low count rate of the separated particles in resolving mode. In particular, for larger particles the count rate achieved was only 6 min⁻¹. The modifications of the ion source, atmospheric pressure interface, and use of additional ion optics could help to increase the count rate at least several times. Moreover, lower quadrupole resolution settings could further increase the transfer rate of vesicles, although the resulting size distribution will be broader.

Comparing our findings to those obtained by other separation techniques so far, the obvious advantage of the presented approach is its selectivity since the mass and charge are measured for each individual particle. However, we consider the short (millisecond) time scale needed for separation and mass characterization of an individual vesicle to be the main benefit of our approach. It enables a direct and contamination-free selection of vesicles that can be subjected to additional analyses. Particularly, in an electrodynamic trap placed downstream, selected vesicles could be accumulated and irradiated by short mid-IR laser pulses inducing the desorption of (charged) proteins and peptides from the vesicle membrane,⁴⁴ which can then be analyzed using e.g., a TOF mass spectrometer. A dedicated power supply allowing simultaneous storage of both the vesicles and desorbed species would be necessary for such a trap. 19 For example, to trap ions with m/z from 200 to 2×10^6 in a linear quadrupole trap with a 0.475 cm effective radius, a harmonic voltage with frequency of 1 MHz and an amplitude of 400 V would be required. 45 To this end, the count rate of 26 min⁻¹ achieved for the serum sample at $m/z_{\rm DQ}$ 5.5 \times 10⁵ makes us confident that such trapping experiment coupled with desorption MS analysis may be worth trying. Our future development of the current concept will therefore aim at the separation of individual EVs from real biomedical samples, with the main objective of obtaining EVs' signatures (e.g., mass and charge distributions) together with the mass spectra of cargo proteins.

ASSOCIATED CONTENT

Data Availability Statement

The data that support the findings of this study are openly available in the National Data Repository at 10.48700/datst. hqpbx-8ws91.

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acs.analchem.4c05730.

Histogram plots for charge measurement error; charge detector measurement error; table with quadrupole settings; microphotography of nanoelectrospray emitters; DLS analysis of 200 nm vesicle sample; examples of the charge detector signal postprocessing; table with

samples comparison and summary of the validation; histogram distributions for 300 nm polystyrene beads; histogram distributions for vesicles in nonresolving mode; density histogram distributions of vesicles diameter in nonresolving mode; histogram distributions of vesicles in resolving mode; density histogram distributions of vesicles diameter in resolving mode; comparison of the different m/z histogram distributions; measurement time series of the vesicle count rate in nonresolving mode; lipid vesicles number calculation; heatmaps for the charge vs mass for vesicles; DLS analysis of the blood serum sample; histogram distributions for serum samples in nonresolving mode; histogram distributions for serum samples in resolving mode (PDF)

AUTHOR INFORMATION

Corresponding Author

Anatolii Spesyvyi — J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; orcid.org/0000-0003-0220-0519; Email: anatolii.spesyvyi@jh-inst.cas.cz

Authors

Marek Cebecauer − J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; orcid.org/0000-0002-4606-1218

Ján Žabka – J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia

Agnieszka Olżyńska – J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; orcid.org/0000-0001-6533-5280

Michaela Malečková – J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; orcid.org/0000-0002-2992-9580

Zuzana Johanovská – J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; Faculty of Mathematics and Physics, Charles University, Prague 12116, Czechia

Miroslav Polášek – J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; © orcid.org/0000-0003-3753-4738

Ales Charvat – Institute of Chemical Technology, Leipzig University, Leipzig 04103, Germany; Leibniz Institute of Surface Engineering, Leipzig 04318, Germany

Bernd Abel – J. Heyrovský Institute of Physical Chemistry of the Czech Academy of Sciences, Prague 18223, Czechia; Institute of Chemical Technology, Leipzig University, Leipzig 04103, Germany; Leibniz Institute of Surface Engineering, Leipzig 04318, Germany; orcid.org/0000-0001-6032-1680

Complete contact information is available at: https://pubs.acs.org/10.1021/acs.analchem.4c05730

Author Contributions

Conceptualization: AS, MC, JZ, BA; software: AS; validation: AS, JZ, AO, AC, BA; formal analysis: AS, AO; investigation: AS, AO, MM, ZJ, MP, AC; resources: MC, AO, MP, AC; data curation: AS; writing—original draft: AS; writing—review and editing: AS, MC, AO, MP, AC, BA; visualization: AS; supervision: JZ, BA; project administration: AS, JZ; funding

acquisition: JZ, BA. All authors have read and agreed to the published version of the manuscript.

Notes

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

The authors acknowledge the financial support of the German Science Foundation (DFG) through grant AB 63/25-1 and by the Czech Science Foundation (GAČR) through grant No. 21-11931J. JZ and BA are grateful for funding within the European Union ERA-Chair Project 101186661 — SPACE. The authors thank Kateřina Paldusová for helping with the blood serum sample preparation. We are grateful to the reviewers for useful comments and helpful suggestions. Open access publishing facilitated by Ústav fyzikalní chemie J. Heyrovského, Akademie věd České republiky, as part of the ACS-CzechELib agreement.

REFERENCES

- (1) Welsh, J. A.; Goberdhan, D. C.; O'Driscoll, L.; Buzas, E.; Blenkiron, C.; Bussolati, B.; Cai, H.; Di Vizio, D.; Driedonks, T. A. P.; Erdbrügger, U.; Falco-Perez, J. M.; Fu, Q. L.; Hill, A. F.; Lenassi, M.; Lim, S. K.; Mahoney, M. G.; Mohanty, S.; Möller, A.; Nieuwland, R.; Witwer, K. W. J. Extracell Vesicles 2024, 13 (2), e12404.
- (2) Meldolesi, J. Curr. Biol. 2018, 28 (8), R435-R444.
- (3) Ciferri, M. C.; Quarto, R.; Tasso, R. Biology-Basel 2021, 10 (5), 359.
- (4) Mathew, B.; Mansuri, M. S.; Williams, K. R.; Nairn, A. C. Brain Sci. 2021, 11 (2), 258.
- (5) Pocsfalvi, G.; Stanly, C.; Vilasi, A.; Fiume, I.; Capasso, G.; Turiák, L.; Buzas, E. I.; Vékey, K. *Mass Spectrom Rev.* **2016**, 35 (1), 3–21.
- (6) Askeland, A.; Borup, A.; Ostergaard, O.; Olsen, J. V.; Lund, S. M.; Christiansen, G.; Kristensen, S. R.; Heegaard, N. H. H.; Pedersen, S. Biomedicines 2020, 8 (8), 246.
- (7) van der Pol, E.; Coumans, F. A. W.; Grootemaat, A. E.; Gardiner, C.; Sargent, I. L.; Harrison, P.; Sturk, A.; van Leeuwen, T. G.; Nieuwland, R. J. Thromb Haemost 2014, 12 (7), 1182–1192.
- (8) van Niel, G.; D'Angelo, G.; Raposo, G. Nat. Rev. Mol. Cell Bio 2018, 19 (4), 213–228.
- (9) Piontek, M. C.; Lira, R. B.; Roos, W. H. Bba-Gen Subjects 2021, 1865 (4), 129486.
- (10) Nikoloff, J. M.; Saucedo-Espinosa, M. A.; Kling, A.; Dittrich, P. S. P Natl. Acad. Sci. USA 2021, 118 (38), e2106630118.
- (11) van der Koog, L.; Gandek, T. B.; Nagelkerke, A. Adv. Healthc Mater. 2022, 11 (5), 2100639.
- (12) Cebecauer, M.; Amaro, M.; Jurkiewicz, P.; Sarmento, M. J.; Sachl, R.; Cwiklik, L.; Hof, M. Chem. Rev. 2018, 118 (23), 11259–11297.
- (13) Weiss, V. U.; Wieland, K.; Schwaighofer, A.; Lendl, B.; Allmaier, G. Anal. Chem. **2019**, 91 (6), 3860–3868.
- (14) Brown, B. A.; Zeng, X. Y.; Todd, A. R.; Barnes, L. F.; Winstone, J. M. A.; Trinidad, J. C.; Novotny, M. V.; Jarrold, M. F.; Clemmer, D. E. *Anal. Chem.* **2020**, 92 (4), 3285–3292.
- (15) Brown, B. A.; Guda, P. R.; Zeng, X. Y.; Anthony, A.; Couse, A.; Barnes, L. F.; Sharon, E. M.; Trinidad, J. C.; Sen, C. K.; Jarrold, M. F.; Ghatak, S.; Clemmer, D. E. *Anal. Chem.* **2022**, *94* (25), 8909–8918. (16) Todd, A. R.; Jarrold, M. F. *Anal. Chem.* **2019**, *91* (21), 14002–
- (16) Todd, A. R.; Jarrold, M. F. Anal. Chem. 2019, 91 (21), 14002–14008.
 (17) Xiong, C. Q.; Liu, H. H.; Li, Y. Z.; Meng, L. W.; Wang, J. Y.;
- Nie, Z. X. Anal. Chem. **2022**, 94 (6), 2686–2692. (18) Lee, J.; Marino, M. A.; Koizumi, H.; Reilly, P. T. A. Int. J. Mass Spectrom. **2011**, 304 (1), 36–40.
- (19) Bhanot, J. S.; Fabijanczuk, K. C.; Abdillahi, A. M.; Chao, H. C.; Pizzala, N. J.; Londry, F. A.; Dziekonski, E. T.; Hager, J. W.; McLuckey, S. A. *Int. J. Mass Spectrom.* **2022**, 478, 116874.

- (20) Patil, A. A.; Liu, Z.-X.; Chiu, Y.-P.; Lai, T. K. L.; Chou, S.-W.; Cheng, C.-Y.; Su, W.-M.; Liao, H.-T.; Agcaoili, J. B. A.; Peng, W.-P. *Talanta* **2023**, 259, 124555.
- (21) Schlunegger, U. P.; Stoeckli, M.; Caprioli, R. M. Rapid Commun. Mass Sp 1999, 13 (18), 1792–1796.
- (22) Peng, W. P.; Lin, H. C.; Chu, M. L.; Chang, H. C.; Lin, N. H.; Yu, A. L.; Chen, C. H. Anal. Chem. 2008, 80 (7), 2524–2530.
- (23) Zhu, Z. Q.; Xiong, C. Q.; Xu, G. P.; Liu, H.; Zhou, X. Y.; Chen, R.; Peng, W. P.; Nie, Z. X. *Analyst* **2011**, *136* (7), 1305–1309.
- (24) Nie, Z.; Cui, F.; Tzeng, Y. K.; Chang, H. C.; Chu, M.; Lin, H. C.; Chen, C. H.; Lin, H. H.; Yu, A. L. *Anal. Chem.* **2007**, *79* (19), 7401–7407.
- (25) Douglas, D. J. Mass Spectrom Rev. 2009, 28 (6), 937-960.
- (26) Marmet, P.; Proulx, M. Int. J. Mass Spectrom. 1982, 42 (1-2), 3-10.
- (27) Shinholt, D. L.; Anthony, S. N.; Alexander, A. W.; Draper, B. E.; Jarrold, M. F. *Rev. Sci. Instrum.* **2014**, *85* (11), 113109.
- (28) Opacic, B.; Hoffman, N. M.; Clowers, B. H.; Reilly, P. T. A. *Int. J. Mass Spectrom.* **2018**, 434, 1–6.
- (29) Simke, F.; Fischer, P.; Marx, G.; Schweikhard, L. Int. J. Mass Spectrom. 2022, 473, 116779.
- (30) Hu, R.; Gundlach-Graham, A. Rapid Commun. Mass Sp 2024, 38 (12), e9753.
- (31) Remes, P. M.; Syka, J. E. P.; Kovtoun, V. V.; Schwartz, J. C. Int. J. Mass Spectrom. **2015**, 377, 368–384.
- (32) Londry, F. A.; Hager, J. W. J. Am. Soc. Mass Spectrom. 2003, 14 (10), 1130-1147.
- (33) Spesyvyi, A.; Zabka, J.; Polásek, M.; Charvat, A.; Schmidt, J.; Postberg, F.; Abel, B. *J. Am. Soc. Mass Spectrom.* **2023**, 34 (5), 878–892.
- (34) Spesyvyi, A.; Zabka, J.; Polasek, M.; Maleckova, M.; Khawaja, N.; Schmidt, J.; Kempf, S.; Postberg, F.; Charvat, A.; Abel, B. *Philos. T R Soc. A* **2024**, 382 (2273), 20230208.
- (35) Keifer, D. Z.; Pierson, E. E.; Jarrold, M. F. Analyst 2017, 142 (10), 1654-1671.
- (36) Peng, W. P.; Lee, Y. T.; Ting, J. W.; Chang, H. C. Rev. Sci. Instrum. 2005, 76 (2), 023108.
- (37) Harris, C. R.; Millman, K. J.; van der Walt, S. J.; Gommers, R.; Virtanen, P.; Cournapeau, D.; Wieser, E.; Taylor, J.; Berg, S.; Smith, N. J.; Kern, R.; Picus, M.; Hoyer, S.; van Kerkwijk, M. H.; Brett, M.; Haldane, A.; del Río, J. F.; Wiebe, M.; Peterson, P.; Gérard-Marchant, P.; Sheppard, K.; Reddy, T.; Weckesser, W.; Abbasi, H.; Gohlke, C.; Oliphant, T. E. *Nature* **2020**, *585* (7825), 357–362.
- (38) Virtanen, P.; Gommers, R.; Oliphant, T. E.; Haberland, M.; Reddy, T.; Cournapeau, D.; Burovski, E.; Peterson, P.; Weckesser, W.; Bright, J.; van der Walt, S. J.; Brett, M.; Wilson, J.; Millman, K. J.; Mayorov, N.; Nelson, A. R. J.; Jones, E.; Kern, R.; Larson, E.; Carey, C. J.; Polat, I.; Feng, Y.; Moore, E. W.; VanderPlas, J.; Laxalde, D.; Perktold, J.; Cimrman, R.; Henriksen, I.; Quintero, E. A.; Harris, C. R.; Archibald, A. M.; Ribeiro, A. N. H.; Pedregosa, F.; van Mulbregt, P.; et al. *Nat. Methods* **2020**, *17* (3), 261–272.
- (39) Hunter, J. D. Comput. Sci. Eng. 2007, 9 (3), 90-95.
- (40) Lutomski, C. A.; Gordon, S. M.; Remaley, A. T.; Jarrold, M. F. *Anal. Chem.* **2018**, 90 (11), 6353–6356.
- (41) Liang, S. Y.; Estayan, M. I. C.; Hsieh, L. W.; Pan, M. C.; Li, K. X.; Chang, H. C.; Peng, W. P. Anal. Chem. **2024**, 96 (18), 7179–7186.
- (42) Hogan, J. A.; Jarrold, M. F. J. Am. Soc. Mass Spectrom. 2018, 29 (10), 2086–2095.
- (43) Shu, A.; Collette, A.; Drake, K.; Grün, E.; Horányi, M.; Kempf, S.; Mocker, A.; Munsat, T.; Northway, P.; Srama, R.; Sternovsky, Z.; Thomas, E. Rev. Sci. Instrum. 2012, 83 (7), 075108.
- (44) Hellwig, N.; Martin, J.; Morgner, N. Biochem Soc. T **2022**, 50 (3), 1057–1067.
- (45) Gerlich, D. Adv. Chem. Phys. 1992, 82, 1-176.