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Possibilities and limitations of current technologies for quantification of biological extracellular vesicles and synthetic mimics 2

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1. Introduction

Extracellular vesicles (EVs) are lipid membrane-enclosed vesicles 43 released by cells and present in bodily fluids. EVs are heterogeneous in 44 composition and size, ranging from approximately 50 to 1000 nm, 45with the vast majority <200 nm in size [1,2]. EVs originate from their 4647donor cell as a result of outward budding of the plasma membrane. Alternatively, EVs form as a result of intracellular budding within late 48endosomes, from which vesicles are released upon fusion of these 49 50multivesicular bodies with the plasma membrane [3]. Regardless of their size and origin, EVs are the collective term adopted to designate 51 any type of cell-derived vesicle in the extracellular space. In recent 5253years, multiple reports have demonstrated EVs to play an important 54role in (patho)physiological processes, such as immune responses [4], 55blood coagulation [5], tissue repair [6] and tumor growth [7,8]. Current

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ABSTRACT

Nano-sized extracelullar vesicles (EVs) released by various cell types play important roles in a plethora of 24 (patho)physiological processes and are increasingly recognized as biomarkers for disease. In addition, 25 engineered EV and EV-inspired liposomes hold great potential as drug delivery systems. Major technologies 26 developed for high-throughput analysis of individual EV include nanoparticle tracking analysis (NTA), 27 tunable resistive pulse sensing (tRPS) and high-resolution flow cytometry (hFC). Currently, there is a 28 need for comparative studies on the available technologies to improve standardization of vesicle analysis in 29 diagnostic or therapeutic settings. 30

We investigated the possibilities, limitations and comparability of NTA, tRPS and hFC for analysis of tumor 31 cell-derived EVs and synthetic mimics (i.e. differently sized liposomes). NTA and tRPS instrument settings 32 were identified that significantly affected the quantification of these particles. Furthermore, we detailed the 33 differences in absolute quantification of EVs and liposomes using the three technologies. This study increases 34 our understanding of possibilities and pitfalls of NTA, tRPS and hFC, which will benefit standardized and 35 large-scale clinical application of (engineered) EVs and EV-mimics in the future.

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research focuses on obtaining improved insight into the formation and 56 function of EVs and on studying the potential of EVs for medical applica-57 tions. One of these applications is to use EVs present in body fluids as 58 biomarkers for diagnosis and monitoring of diseases [9,10]. In cancer, 59 tumor-derived EVs can serve as biomarkers since they contain proteins 60 and RNAs from their malignant donor cells [7,8]. Since tumor-derived 61 EVs are released in easily accessible bodily fluids, such as blood or 62 urine [7,11], analysis of these EVs for disease monitoring may circum- 63 vent biopsies [11], thereby reducing biopsy related morbidity and mor- 64 tality. A second important application of EV in the medical field is their 65 use as drug delivery systems. Although liposomes, which share the 66 bilayered membrane structure with EVs, have been employed as drug 67 delivery systems for many years, cross-pollination of knowledge in the 68 liposome and EV research fields now holds high promise for 69 improvement of current delivery systems. Various studies have indicated 70 that EVs can be exploited as carriers for delivery of exogenous therapeutic 71 cargoes, e.g. siRNAs, in vivo [12]. EV characteristics that facilitate efficient 72 delivery of biological drugs include their capacity to traverse intact 73 biological barriers (e.g. blood-brain barrier) and to deliver functional 74 RNA into cells, as well as their stability in blood (reviewed in [13]). 75

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Current research focuses on exploiting these features to either engineer natural EV for drug delivery to specific tissues, or to design EV mimics formulated as liposomes containing relevant EV components.

Even though EVs are increasingly recognized as important biological and therapeutical entities, standardized methods for their analysis are still lacking. Establishment of such methods is crucial for safe application of (engineered) EV in clinical practice, but EV quantification has proven technically difficult due to the small size of EVs and their heterogeneity in size and composition.

85 In recent years, several instruments have become available 86 that allow detection and characterization of individual EVs. These 87 techniques include nanoparticle tracking analysis (NTA) [18,19], tunable resistive pulse sensing (tRPS) [20] and high-resolution flow 88 89 cytometry (hFC) [21]. EV detection and quantification with these single-particle analysis techniques rely on distinct principles. NTA is 90 based on the illumination of particles in suspension with a laser beam, 91 followed by the recording of the scattered light by a light-microscope. 92 93 The Brownian motion of each particle is individually tracked to determine the mean square displacement of the individual particle. Since 94 temperature and viscosity of the suspension are known and controlled, 95 the Stokes-Einstein equation can be used to determine the hydrody-96 namic diameter of each individual particle. The total number of particles 97 98 is used for particle concentration estimation [18,22]. In tRPS, a nonconductive polyurethane membrane, punctured to contain a single 99 opening, separates two fluid cells [23]. By applying a voltage across 100 the membrane a flow of ions is induced. Once a particle moves through 101 the nanopore, the flow of ions is altered resulting in a brief "resistive 102103 pulse" which is recorded by the instrument [24]. The size-distribution [25] and concentration [26,27] of particles can be calculated by referring 104 the observed pulse height and rate to pulses induced by reference 105particles of known volume and concentration. Flow cytometric analysis 106 107of particles involves the sequential excitation of individual, fluorescently labeled particles in a liquid stream and detection of emitted light by 108diodes or photomultipliers [28]. In hFC, a high-end flow cytometer is 109optimized for the analysis of nano-particles. This optimization consists 110 of light scattering detection at customized angles, the usage of 111 high power lasers and high-performance photomultiplier tubes for 112 113 more sensitive light detection, and application of fluorescence-based thresholding to distinguish particles of interest from noise signals 114 [21]. In-depth description of the technical backgrounds of the tech-115 niques is beyond the scope of this manuscript and described elsewhere 116 for NTA [18,22,29,30], tRPS [24-26] and hFC [21,31]. 117

For accurate EV quantification and characterization, it is impor-118 tant to know to what extent instrument-specific variables influence 119 particle characterization. For NTA, studies on how instrument 120settings affect the analysis of heterogeneous EV populations are lim-121 122ited [22,30,19], and the effects of specific variables on EV quantification and size-profiling by tRPS are largely unknown. For hFC, detailed 123reports on optimizing the instrument configuration and settings for 124accurate analysis of EVs and other nano-sized particles have recently 125been published [21,31]. In a few studies, two or three of the above 126127described techniques have been compared. However, these studies 128either focused on size-profiling of synthetic beads [34,35], or did not address effects of instrument settings on EV characterization 129and quantification [36,37]. 130

Here, we report a comprehensive comparative study on NTA, tRPS 131 132and hFC for analysis of populations of heterogeneous nano-sized EVs and synthetic mimics (i.e. polystyrene beads and calcein-loaded 133 liposomes). We identified different NTA- and tRPS-variables that 134significantly influenced the quantification of these particles. Further-135more, we assessed the comparability of NTA, tRPS and hFC in absolute 136quantification of liposomes and EVs. Based on these data, we stress 137 the importance of technical knowledge of the instruments, awareness 138 of analytical variables, and recognition of how instrument settings affect 139measurements when analyzing EV populations with unknown concen-140 141 tration and size heterogeneity.

2. Materials and methods 142

2.1. Polystyrene beads

115 and 203 nm polystyrene beads (Izon Science, Christchurch, New 144 Zealand) were analyzed using tRPS and NTA. For hFC, fluorescent 100 145 and 200 nm polystyrene beads (yellow-green-fluorescent FluoSpheres, 146 Invitrogen) were used. 147

2.2. Liposome preparation and characterization

Egg phosphatidylcholine (EPC), egg phosphatidylglycerol (EPG) (Li- 149 poid GmbH, Ludwigshafen, Germany) and cholesterol (Sigma-Aldrich 150 Chemie B.V., Zwijndrecht, The Netherlands) were dissolved in chloro- 151 form/methanol (1:1, v/v) in a round-bottom flask in a molar ratio of 152 2:0.06:1, respectively. A lipid film was prepared by rotary evaporation 153 (Rotavapor R3, Büchi Labortechnik AG, Flawil, Switzerland), followed 154 by drying under a stream of nitrogen. The lipid film was hydrated 155 with 10 mM calcein for 105 nm liposomes or 250 µM calcein for 156 "L146" and "L212" liposomes in HEPES buffered saline (HBS, 10 mM 157 HEPES, 137 mM NaCl, pH 7.4). Liposomes were sized by multiple 158 extrusion under nitrogen pressure using polycarbonate membranes 159 (Nuclepore, Pleasanton, CA, USA) with pore sizes of 200 nm and 160 100 nm in a Lipex high pressure extruder (Lipex, Northern Lipids, Van- 161 couver, Canada) or a Liposofast Extruder (Avestin, Inc, Ottawa, Canada). 162 Non-entrapped calcein was removed with dialysis against HBS for at 163 least 3 days using Slide-A-Lyzer dialysis cassettes with a cut off of 164 10 kD (Thermo Scientific, Bremen, Germany). The mean particle size 165 of the liposomes and the polydispersity index (PDI) was determined 166 by means of dynamic light scattering (DLS) using a Malvern ALV CGS-3 167 with a He-Ne laser source (Malvern Instruments, Malvern, UK). 168 Liposome sizes (L146 and L212) were 146 nm with a PDI of 0.03 and 169 212 nm with a PDI of 0.07. The zeta-potential of the liposomes (ζ poten-170 tial) was determined using a Malvern Zetasizer Nano-Z (Malvern Instru- 171 ments, Malvern, UK). The phosphate concentrations of the liposomes 172 were determined with a phosphate assay described by Rouser et al. 173 [38]. For final use, L146 and L212 liposomes were diluted with HBS till a 174 final total lipid (including cholesterol) concentration of 65 mM. 175

2.3. Cell culture and EV isolation

The human glioblastoma cell line U87-MG and the lymphoblastoma 177 cell line RN were cultured in medium containing FCS depleted from 178 bovine EVs as described previously [20,21]. After 24 h of incubation 179 the supernatant was isolated and centrifuged at 200 \times g for 10 min, 180 two times at 500 ×g for 10 min, followed by 10,000 g for 30 min. 181 100,000 \times g pelleted EVs were resuspended in phosphate buffered sa- 182 line (PBS) containing 0.2% BSA from an ultracentrifuged stock solution 183 [31]. EVs were fluorescently labeled with 7.5 µM PKH67 (Sigma-Al- 184 drich), mixed with 2.5 M sucrose, overlaid with a linear sucrose gradient 185 (2.0-0.4 M sucrose in PBS) in an SW60 tube (Beckman) and floated into 186 the gradient by centrifugation for 16 h at 192,000 \times g [31]. Gradient 187 fractions were collected, diluted in PBS and analyzed. Fraction densities 188 were determined by refractometry. 189

An LM14 Nanosight instrument (Nanosight Ltd, Salisbury, UK) 191 equipped with a CMOS camera (Hamamatsu Photonics, Hamamatsu, 192 Japan) and a 488 nm laser was used. Data acquisition and processing 193 were performed using NTA software 2.3 build 0025. Background extrac- 194 tion was applied, and automatic settings were applied for the minimum 195 expected particle size, minimum track length and blur settings. Since 196 samples were diluted at least 20 times in PBS, viscosity settings for 197 water were applied and automatically corrected for the temperature 198 used. Detection threshold and camera level settings varied as described 199

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in the Results section. Five movies of 60 s at 25 frames per second were
 recorded and designated as a single measurement. Only measurements
 with at least 1000 completed tracks were further analyzed [19]. For
 polystyrene bead dilutions, single measurements were performed for
 each dilution, whereas triplicates were recorded for liposome and EV
 samples.

We excluded data obtained at camera-level 15 (shutter: 1200, gain:
 500) as this camera-level resulted in the detection of substantial amounts
 of background detection, obscuring accurate data interpretation.

209 2.5. tRPS

For tRPS, the qNano instrument (Izon Science Ltd, Christchurch, New 210211Zealand) was used as described [39]. Data was recorded and analyzed using the Izon Control Suite Software version 2.2.2.111. The default 212 minimum blockade height (0.05 nA) for particle detection was used. 213 For sample calibration and serial dilution experiments, polystyrene 214 beads supplied by the gNano manufacturer were used. Both 115 and 215203 nm polystyrene bead dilutions were recorded using NP100 216 nanopores. Liposome dilutions were recorded using two different 217NP100 nanopores at 0.8 kPa and 1.2 kPa pressure settings. EV samples 218 were analyzed using both an NP200 (1.2 kPa pressure) and NP150 219 220nanopore (1.4 kPa). The buffers of EVs and calibration beads were 221 kept identical by diluting the calibration beads in the appropriate fraction of a (mock-loaded) sucrose-based density gradient. 222

223 2.6. hFC

High-resolution flow cytometric analysis of individual EVs was 224performed using the BD Influx flow cytometer (Becton Dickinson, San 225Jose, CA, USA) with an optimized configuration, as described in detail 226 227before [31]. Light scattering was measured with a collection angle of 22815–25° (reduced wide-angle FSC) and detection was performed in log 229mode. Samples were run at low pressure (5 PSI on the sheath fluid and 4.2 PSI on the sample) using a 140 µm nozzle. The calculated flow 230rate at these settings was 52.2 µl per minute, as determined by 231weighing the volume aspirated during 30 min. Fluorescent 100 nm 232233 and 200 nm polystyrene beads (yellow-green-fluorescent FluoSpheres, Invitrogen) were used for calibration of the fluorescence, reduced 234wide-angle FSC, and SSC settings. EVs in sucrose fractions were diluted 235in PBS at least 20 times and time-based quantitative measurements 236were performed as described before [31]. Data was acquired using 237Spigot software version 6.1 (Becton Dickinson). Data was further 238analyzed using FCS Express software (De Novo Software, Los Angeles, 239USA). 240

241 2.7. Statistical analysis

Data analysis was performed using Prism version 5.0 (GraphPad 242 Software, La Jolla, CA, USA) or Microsoft Excel 2010 (Microsoft, Seattle, 243WA, USA). Two-tailed independent t-tests were used to test for 244245significant differences in means. One-way ANOVA followed by Tukey's 246post-test was performed to test differences between multiple groups. Pearson's correlation coefficient was used to determine the correlation 247between dilution and the measured concentration. Significance was 248determined and indicated as (*) p-value \leq 0.05, (**) p-value \leq 0.01 249250and (***) p-value \leq 0.001. Error bars represent the standard deviation (s.d.) unless stated otherwise. 251

252 **3. Results**

253 3.1. NTA-based particle quantification

Detection of nano-sized particles with NTA is influenced by two parameters: the camera-level (shutter speed and camera gain), which is set prior to data acquisition, and the detection threshold, i.e. the scattering intensity threshold above which particles are traced (set at 257 data processing). Here, we tested how these parameters affected the 258 quantification of nano-sized particles that differ in refractive index, 259 size, and heterogeneity. 260

First, we determined the accuracy for quantification of homoge- 261 neous populations of 115 or 203 nm sized polystyrene beads, which 262 have a high refractive index (r.i.) and consequently cause extensive 263 light scattering. Within a 32-fold dilution range $(0.9-29.0 \times 10^8/\text{ml})$ 264 the measured concentrations approximated the expected concentrations for 115 and 203 nm beads (Fig. 1A) (R²: 0.969 and 0.998 266 respectively). 267

For the 115 nm beads we were also able to obtain valid measure- 268 ments (>1000 completed tracks) outside this range. However, the 269 resulting s-shaped curve (Fig. 1A) indicates an overestimation of parti- 270 cles below 0.9×10^8 /ml and an underestimation above 29.0×10^8 /ml, 271 resulting in decreased correlation accuracy (R²: 0.859 for all measure- 272 ments). Although the range of refractive indices that EVs can exhibit is 273 largely unknown, polystyrene beads most likely have an r.i. that is sub- 274 stantially higher than the r.i. of the majority of EVs [19]. Next, we tested 275 how camera level and detection threshold settings affect the quantifica- 276 tion of calcein-labeled liposomes, which, similar to EVs, are enclosed by 277 a lipid bilayer. Movies were recorded, at camera level 6 (shutter: 150, 278 gain: 250), camera level 9 (shutter: 450, gain: 250), and camera level 279 12 (shutter: 600, gain: 350), which represent preprogrammed NTA 280 settings. After data acquisition each movie was processed at detection 281 threshold 4, 6, 8 and 10 (standard software setting). 282

As expected, at increased camera levels the particles appeared 283 brighter and increased detection was observed of weak-scattering 284 particles (Fig. 1B, top-panel). The number of detectable particles was 285 also increased by reducing the detection threshold (Fig. 1B, bottom- 286 panel) (A complete overview of screenshots at different camera levels 287 and detection thresholds is provided in Suppl. Fig. S1A). Numerical 288 analysis of these data revealed that the quantification of liposomes is 289 significantly influenced by the NTA settings, with measurement of 290 higher concentrations after increasing the camera level or decreasing 291 the detection threshold (Fig. 1C and Suppl. Fig. S1B). At both camera 292 levels 9 and 12, accurate linearity in measured concentration was 293 observed for multiple dilutions of liposomes (applying camera level 6 294 resulted in an inadequate number of completed tracks) (Fig. 1D). 295 Thus, relative comparison of liposome concentrations is feasible with 296 different NTA settings, but measurement of the exact concentration 297 strongly depends on the camera-level and detection threshold settings. 298 Of interest, the increased number of liposome detection after increasing 299 the camera level is not accompanied by increased detection of smaller 300 sized liposomes (Suppl. Fig. S1C). It has previously been suggested 301 that smaller particles may be over-scattered by larger particles, which 302 would especially occur after concentrating samples [19,40]. However, 303 this appears not to occur for liposomes (Suppl. Fig. S1D). 304

Next, we tested to what extent the camera-level and detection 305 threshold influence the quantification of EVs, which are more variable 306 in size and r.i. than liposomes. For these experiments, we used EVs 307 derived from the RN lymphoblastoma and U87-MG glioblastoma cell 308 lines that were purified from contaminating protein aggregates by 309 sucrose density gradient ultracentrifugation. As expected, increasing 310 the camera level resulted in an increased brightness of detected parti- 311 cles (Suppl. Fig. S2A). Similar to the liposome analysis, different EV 312 quantification data were obtained at different camera level settings 313 (Fig. 1E). A maximum fold change of 3 was observed (camera-level 6 314 versus 12 at detection threshold 10). The influence of detection thresh- 315 old on particle quantification was less prominent for EVs compared to 316 liposomes (Fig. 1E and Suppl. Fig. S2B). Similar data were obtained for 317 the U87-MG derived EVs (data not shown). Sample dilution did not 318 significantly influence measurement of the raw EV concentration (i.e. 319 the measured sample concentration multiplied by the dilution factor) 320 (Fig. 1F). This was corroborated by the observation that the size- 321 distributions and mode sizes were similar at the different dilutions 322

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Fig. 1. NTA-based quantification of beads, liposomes and EVs. (A) Quantification of 115 and 203 nm polystyrene beads. The measured concentration of the beads is plotted against the expected concentration based on the manufacturer's supplied stock concentration. Detection was performed at camera level 5 (shutter: 100, gain: 200) for the 115 nm beads and camera level 3 (shutter: 20, gain: 0) for the 203 nm beads. (B–D) Quantification of 115 nm-sized liposomes. The effect of camera level and detection threshold was assessed, demonstrating visual differences in particle imaging (screenshots in (B)) as well as differences in the calculation of raw concentrations (C). Dilution of liposomes showed linearity with the measured liposome concentration, at camera levels 9 and 12 (D). (E–F) Quantification of purified EVs. The effect of camera level and detection threshold on quantification of EVs (from RN cells) is shown in (E). The effect of sample dilution (1:20 and 1:100) on quantification is shown in (F), with EVs included from RN cells and U87-MG cells, and analysis at three different camera levels. Data are mean \pm s.d. (n = 3).

(Suppl. Fig. S2C and S2D). Thus, although EVs are more heterogeneous
in size than liposomes, the presence of infrequent EV that displayed a
higher level of scattering did not substantially affect detection of neighboring EVs.

In conclusion, camera-level and detection threshold variables were
 found to affect the NTA-based quantification of liposomes and EVs.
 The influences were more profound for the relatively homogeneous
 liposomes than for the heterogeneous EVs.

331 3.2. tRPS-based particle quantification

As an alternative to NTA, we tested tRPS for liposome and EV quantification, with a specific focus on establishing the most suitable measurement conditions. As tRPS-based quantification requires a linear correlation between the particle count rate (particles per minute) and the concentration of particles, we first measured a dilution range of polystyrene calibration beads. We observed linearity over a 64-fold dilution range for the 115 nm beads (R^2 : 0.979) and over a 32-fold 338 range for the 203 nm beads (R^2 : 0.994) (Fig. 2A). 339

Particle detection above threshold levels is dependent on the block- 340 ade height (resistive pulse) generated by a particle moving through a 341 nanopore. This blockade height is determined by the particle's volume 342 relative to the volume of the nanopore opening, the applied voltage, 343 and buffer used. These parameters together determine a 'tRPS setup' 344 and thus determine if particles surpass threshold levels (0.05 nA at 345 default software settings). High-sensitivity tRPS setups can be used to 346 detect the smallest particles. To obtain a high-sensitivity setup one 347 should apply a high voltage, low stretch (to establish a minimal opening 348 size of the nanopore) and a small nanopore (NP100/NP150) [36,39]. 349 Nanopore characteristics are known to differ between individual 350 nanopores, as well as over time [41]. To assess the effect of this on 351 liposome quantification, we compared three cases. First we compared 352 two new NP100 nanopores (setups #1 and #2). Subsequently, the 353 nanopore used for setup #1 was tested again after approximately 7 h 354

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Fig. 2. tRPS-based particle quantification. (A) Quantification of 115 and 203 nm polystyrene calibration beads. As tRPS quantification is based on the conversion of observed particle per minute counts to that of polystyrene calibration beads of known concentration, the read-out is displayed as "particles per minute". (B) Three tRPS setups displaying the observed blockade heights for the same 115 nm calibration beads (left-panel). The dashed line illustrates the detection threshold (both panels). Reconstruction of the recorded data for beads and liposomes at the three different setups (right-panel), illustrating that the lower detection limit is the highest for setup #3, followed by setups #1 and #2. Bin size 15 pA. (C) Representative liposome size-distributions obtained at the three different tRPS setups (left-panel). For each of the three setups the measured concentration was corrected for the dilution factor to obtain raw concentration estimations (n = 6) (right-panel). Bin size 5 nm. (D) Quantification of serially diluted liposomes at two different pressure levels (n = 3). (E) Representative size-distribution obtained for RN-derived EVs on an NP200 nanopore setup and an N150A nanopore. Bin size 5 nm. (F) Raw particle concentrations were determined for RN and U87-MG derived EVs at both the NP200 and NP150 nanopore setups (n = 3). Data are mean \pm s.d.

of usage (termed setup #3). The most optimal (i.e. high-sensitivity) 355 356 settings were applied in all three cases. Measurement of 115 nm poly-357styrene beads for tRPS calibration showed different mode blockade heights detected for setups #1, #2, and #3, with #2 > #1 > #3 (Fig. 2B, 358left panel). This indicated that the lower detection limit, as determined 359by the height of the calibration bead blockade relative to the threshold 360 level of 0.05 nA, was different at the different setups. This is illustrated 361 by reconstructing the mean blockade heights of 115 nm calibration 362 particles and liposomes for the three setups (Fig. 2B, right panel). 363 Using setup #3, the peaks of the 115 nm calibration particles are closer 364 to the detection threshold level. The lower detection limit in setup #3 is 365 therefore higher than in setups #1 and #2, implicating that the blockade 366 height induced by smaller liposomes may not surpass the detection 367 threshold. This could result in the detection of only larger-sized lipo-368 somes. Secondly, these observed differences indicate that characteris-369 370 tics of nanopores, such as resolution, may change over time.

The variation in detectable size range for setups 1–3 resulted in 371 substantial differences in absolute quantification of the liposomes 372 (difference setup #2 versus #3: 2.43 fold; Fig. 2C). Setup #3 allowed de- 373 tection of >80 nm liposomes only (Fig. 2C, left-panel), and consequently 374 yielded the lowest liposome quantification. Differences in particle 375 concentration (1.43 fold) were also observed for setups #1 and #2, 376 despite the comparable efficiency in detection of small liposomes at 377 these setups (Fig. 2C, left-panel). 378

Besides absolute quantification of liposomes, we also determined 379 how accurate a range of liposome dilutions could be quantified by 380 tRPS (Fig. 2D). An NP100 nanopore was used for this test, and we concomitantly investigated whether the pressure level influenced liposome 382 quantification. For both applied pressure levels we observed accurate 383 detection and linearity over a 4-fold dilution range. Surprisingly, changing the applied pressure led to significantly different liposome sizing 385 estimations (Suppl. Fig. 3A and B). 386

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EV measurements by tRPS indicated that the size distributions of RN 387 388 (Fig. 2E) and U87-MG derived EVs resembled those obtained using the NTA, with the majority of EVs being 100-200 nm in size. Similar to 389 390 what we observed for NTA, tRPS showed the presence of a small number of larger (200–600 nm) EVs. Due to the presence of large EVs, we tested 391 two larger nanopores (NP200 and NP150) for EV quantification, to 392 reduce clogging events. Even though frequent nanopore clogging was 393 394 observed, overall particle detection was stable and reproducible for 395 each triplicate of sample measurements (Suppl. Fig. S4A). Applying 396 the NP150 nanopore, which theoretically allows for detection of 100-120 nm particles, yielded significantly higher EV particle concen-397 trations as compared to the measurement with the NP200 pore 398 (Fig. 2F) (difference RN-derived EVs 1.45 fold, U87-derived EVs 1.50 399 fold). The ability to measure smaller sized EVs with the NP150 nanopore 400 (Suppl. Fig. S4B), led to significant differences in the calculated mean 401 and mode sizes of the EVs (e.g. mode sizes of 136.3 nm (NP200) and 402117.8 nm (NP150) for RN-derived EVs) (Suppl. Fig. S4C). 403

In conclusion, quantifications of liposomes and EVs can differ
 between (high sensitivity) nanopore setups and this is most likely relat ed to the lower detection limit. Since the required lower detection limit

may be unknown for liposomes and EVs, tRPS measurement may result 407 in underestimation of the concentration. 408

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3.3. hFC-based particle quantification

Reliable quantification of nano-sized particles using fluorescence- 410 based hFC requires that sufficient numbers of fluorophores are associat- 411 ed to the particle to be detected above the fluorescence threshold, and 412 that maximal sensitivity in fluorescence detection is obtained. The 413 optimal configuration and settings for quantitative and qualitative anal-414 yses of nano-sized particles using the BD Influx have been determined 415 previously [21,31]. Instrument settings that were found to affect EV 416 measurements included the nozzle size and the applied sample/sheath 417 fluid pressure. Using the optimal settings, 100 nm fluorescent 418 polystyrene beads were efficiently detected above background noise. 419 Furthermore, for a 16-fold dilution range, hFC accurately detected 420 sample dilutions for both 100 (R^2 : 1.00) and 200 nm (R^2 : 0.999) fluores-421 cent polystyrene beads (Fig. 3A). 422

Calcein labeled, 105 nm sized liposomes could also be detected 423 above the fluorescence threshold (Fig. 3B). As expected, light scattering 424



Fig. 3. hFC-based particle quantification. (A) Quantification of serially diluted 100 and 200 nm fluorescent polystyrene beads. Indicated are the mean number of beads detected in a fixed time window of 30 s. (B) Dotplots indicating that calcein-loaded liposomes can be detected above the fluorescence threshold (solid horizontal line) that excludes detection of non-fluorescent noise events (left-panel) and that light scattering levels induced by these liposomes are low (right-panel). (C) Quantification of serially diluted calcein-loaded liposomes. Indicated are the mean number of liposomes detected in a fixed time window of 30 s. (D) Dotplots indicating that PKH-67 labeled RN-derived EVs can be detected above the fluorescence detection threshold (left panel) and that the FSC and SSC signals induced by these heterogeneous are highly variable. (E) EVs were measured over an 8-fold range, and corrected for the dilution used to determine the raw concentration estimations. No statistical different raw concentration estimations between the dilutions were observed. Data are mean \pm s.d. (n = 3).

(FSC and SSC) levels generated by low r.i. liposomes were low and could
not be discriminated from those generated by noise, indicating the need
for fluorescence-based analysis (data not shown). Within the 16-fold
dilution range tested here, liposomes could be quantified with accurate
linearity (R²: 1.00) (Fig. 3C).

hFC-mediated detection of RN and U87-MG derived EVs relied on 430 fluorescent labeling of EV and efficient removal of unbound dye by 431 sucrose-gradient ultra-centrifugation [31] (Fig. 3D and Suppl. Fig. 5). 432 433 Although hFC does not allow for absolute size measurement of EV, variation in size and composition of EV are reflected in the light scattering 434 (FSC and SSC) and fluorescence signals observed. Similar to what was 435observed in the NTA and tRPS measurements of the RN and U87-MG 436derived EVs, hFC-based analysis also indicated substantial heterogene-437ity within these EV populations based on light scattering and PKH67 438 fluorescence levels (Fig. 3D and Suppl. Fig. 5). Quantification by hFC 439 indicated no significant differences in the estimation of EV concentra-440 tions over an 8-fold dilution range for both the RN and U87-derived 441 EVs (Fig. 3E). 442

In conclusion, once sufficient numbers of fluorophores are associated
 to liposomes or EVs to allow their detection above the fluorescent thresh old, hFC can be used for accurate quantitative analysis of fluorescently
 labeled liposomes and EVs in a range of sample dilutions.

447 4. Comparison of liposome and EV quantification using NTA, tRPS, 448 and hFC

For clinical application and research purposes, it is of utmost importance to reliably determine the concentrations of (engineered) EVs or synthetic mimics. Ideally, measurements of identical samples with 451 different technologies should yield comparable quantitative data. We 452 therefore compared quantification data obtained by NTA, tRPS, and 453 fluorescence-based hFC. Based on the previous experiments, a single 454 setup was selected for each instrument. We performed measurements 455 on relatively homogeneous populations of calcein-loaded liposomes 456 with (DLS-based) sizes of 146 and 212 nm (referred to as L146 and 457 L212 respectively), and a more heterogeneous population of purified 458 and PKH67 labeled EVs. NTA camera-levels were selected based on 459 the visually brightest detection of particles, without the occurrence of 460 abundant over-scattering events. The tRPS settings were selected to 461 allow for the highest-sensitivity measurement. More specifically, L146 462 measurements were performed with NTA camera-level 12/detection- 463 threshold 4 and nanopore NP100. For L212, camera-level 9/detection- 464 threshold 4 and nanopore NP150 were selected. The RN-EVs were 465 analyzed using NTA camera-level 12/detection-threshold 10 and an 466 NP150 nanopore. Optimized settings [21] were used for hFC, and hFC 467 settings were identical for measurements of both liposome populations 468 and EVs. On the three instruments L146 liposomes were quantified 469 within a 12.5 fold difference (Fig. 4A, left-panel). The highest concentra- 470 tions were measured with NTA (1.86×10^{14} /ml), followed by tRPS 471 $(5.33 \times 10^{13}$ /ml), and hFC $(1.5 \times 10^{13}$ /ml). Also for the L212 liposomes, 472 NTA measurements yielded the highest concentrations $(7.73 \times 10^{13} / 473)$ ml), followed by tRPS $(3.27 \times 10^{13}/\text{ml})$ and hFC $(1.12 \times 10^{13}/\text{ml})$ 474 (Fig. 4A, right-panel). Overall, the measured L212 concentrations on 475 the three instruments were within a narrower absolute fold-range 476 (6.92). We compared these quantifications with liposome concentra- 477 tion measurements based on dynamic light scattering (DLS)-sizing, 478



Fig. 4. Comparison of liposome and EV quantification using NTA, tRPS and hFC. (A) Comparative quantitative analysis of L146 and L212 liposomes using the three instruments (n = 3). Liposomes were diluted to match the required sample concentrations for the different instruments after which measured concentrations were calculated to raw concentrations. Horizontal lines indicate liposome concentration calculations based on lipid composition, phosphate quantification, and dynamic light scattering (DLS) sizing (dotted line) or NTA/tRPS sizing (solid line). Size-distributions for L146 and L212 liposomes as obtained by NTA (B) and tRPS (C). Bin size 5 nm. (D) Comparative quantitative analysis of RN-derived EVs by NTA (n = 4), tRPS and hFC (n = 3) at instrument specific concentrations, converted to raw concentration estimations. Data are mean \pm s.e.m.

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lipid composition, and phosphate quantification, as a standard in the 479480 liposome field [42]. Using this method, the calculated liposome concentrations were 2.27×10^{13} /ml for L146 and 7.46×10^{12} /ml for 481 482 L212 liposomes (dotted horizontal lines in Fig. 4A). However, DLS is known to be heavily influenced by outliers [34], which may result in 483 overestimated size measurements. When replacing DLS size measure-484 ment with averaged liposome sizing data obtained by NTA and tRPS 485(124 and 156 nm for the L146 and L212 liposomes, respectively; 486 487 Fig. 4B and C), the calculated liposome concentrations were substantially higher (solid horizontal lines in Fig. 4A) and were most similar to the 488 489 concentrations obtained by tRPS.

The absolute concentration measurements of EV on the three 490 instruments were within a smaller fold-range difference compared to 491 492 the measurements of liposomes (4.44 versus 6.92 (L146) or 12.5 (L212); Fig. 4D). Interestingly, quantification of EVs by tRPS and hFC 493 vielded absolute particle concentrations in the same range (1.01×10^9) 494 ml and 1.40×10^9 /ml). However, similar to the liposome measurements, 495 NTA yielded substantially higher values for the EV concentrations 496 $(4.50 \times 10^{9} / \text{ml})$ (Fig. 4D). 497

In conclusion, the absolute quantifications as observed for both 498 homogeneous calcein-loaded liposomes and a purified population of 499 more heterogeneous, PKH67 labeled RN derived EVs differed signifi-500501cantly between the instruments. For liposomes, the difference in guantifications between the instruments decreased when measuring 502liposomes that were larger in size. The smallest difference in absolute 503concentration measurements between the instruments was found 504when measuring the more heterogeneous population of EVs, for 505506which tRPS and hFC yielded highly similar results.

507 5. Discussion

508Over the last decade, the interest in EVs has greatly intensified due to their proposed role in various biological processes and their potential as 509biomarkers for disease and as drug delivery systems. Approaches for 510accurate and standardized quantification of such nano-particles 511have not yet been established, but are crucial for safe application of EV 512513(-mimics) in clinical settings. Here, we compared quantification of different nano-sized particles, i.e. polystyrene beads, calcein-labeled fluo-514rescent liposomes and purified, PKH67-labeled EVs using three 515prominent single-EV analysis platforms; NTA, tRPS, and hFC. Moreover, 516517we identified variables that significantly influenced particle quantification using NTA and tRPS. 518

The particle concentration range at which accurate quantification 519520data could be obtained differed between the instruments. For NTA, the optimal concentration range was 9.0×10^7 /ml- 2.9×10^9 /ml, which is 521522a slightly larger dilution series than previously reported [18]. For tRPS, the required concentration for particle analysis increased as the particle 523volume decreased. Consequently, 203 nm beads were analyzed at 524 9.1×10^7 /ml- 2.9×10^9 /ml, whereas 115 nm particles were analyzed 525at 3.6 \times 10 $^8/ml-2.3$ \times 10 $^{10}/ml.$ hFC allows accurate quantification at 526527lower particle concentrations (a range of $4.6 \times 10^6/\text{ml}-7.3 \times 10^7/\text{ml}$ 528was analyzed in the current study). Our recent data indicate that concentrations up to 1.0×10^9 /ml can be reliably measured with hFC 529(manuscript in submission). 530

We identified the NTA camera level and detection threshold to be 531532significant factors in the quantification of liposomes (Fig. 1C). In contrast, the absolute differences induced by changing these variable 533settings were less prominent for quantification of EVs (Fig. 1D). This 534may be a result of the relatively higher light-scattering properties of 535EVs (due to the presence of surface/luminal proteins and/or m(i) 536RNAs), combined with increased heterogeneity in this population, 537which may make NTA-based detection of EVs less sensitive to differ-538ences in settings as compared to the detection of homogeneous lipo-539somes. Besides the empty liposomes used in this study, liposomes 540541engineered to contain proteins and/or nucleic acids show more structural resemblance with EV and quantification of such particles 542 may accordingly be less sensitive to NTA detection thresholding. 543

Our tRPS analyses showed inter-experimental variation in the 544 sensitivity of liposome and EV detection (Fig. 2C and E), which translated 545 into differences in concentration measurement. This sensitivity of tRPS-546 based measurements is determined by the size of the smallest detectable 547 particle. For quantification of homogeneous particle populations of a 548 known size, such as calibration beads, the most suitable nanopore setup 549 for detection of all particles can easily be selected. However, for samples 550 with an unknown size-distribution (e.g. EVs) this is more difficult and the 551 obtained size detection range may be insufficient for detection of all 552 particles. Besides this, we also noted slight differences in tRPS-based 553 concentration measurements (up to ~1.4 fold) between set-ups in 554 which the liposome size-distribution profiles and detections limits were 555 similar (Fig. 2C). We hypothesize that subtle differences in nanopore 556 size due to batch variations and nanopore longevity could have caused 557 these variations in particle quantification. The observed differences stress 558 the importance of comparing samples using exactly the same tRPS setup. 559

Electro-kinetic forces were recently suggested [43] to influence the 560 movement of particles through smaller tRPS nanopore. In case particles 561 possess a different surface-charge compared to the polystyrene calibra- 562 tion particles, one of the two particle types may be more likely to pass 563 through the nanopore. Since this may cause inaccuracy in the calculated 564 particle/minute to concentration calculation, the manufacturer 565 suggested to perform quantifications at two or more pressure levels, 566 after which the tRPS software can determine a surface-charge corrected 567 concentration. Since we observed no difference in the measured 568 liposome concentration at two different pressure levels (Fig. 2D), we 569 conclude that electrokinetic forces at these settings do not significantly 570 influence the quantification of particles. The surface charge of the stud- 571 ied liposomes was -43.0 ± 0.87 mV, which is similar to the reported 572 surface charge characteristics of EVs [44-46]. Single-pressure tRPS 573 quantifications can therefore suffice for accurate EV quantifications. 574 The difference in blockade height when measuring the 115 nm calibra-575 tion particles at the two pressure settings (Suppl. Fig. S3A, left-panel) 576 was unexpected, because the applied pressure does not change the par- 577 ticle volume and nanopore diameter. Implications of this phenomenon 578 for particle characterization need to be further studied. When compar- 579 ing the liposome size-distributions obtained by tRPS and NTA (Fig. 2C 580 and Suppl. Fig. S2C), we conclude that both NTA and tRPS allowed 581 detection of liposomes as small as 55-60 nm in size, which for NTA is 582 the theoretical lower limit of liposome detection, limited by the r.i. of 583 the particle [18,22]. 584

In contrast to NTA and tRPS, for hFC the threshold for particle 585 detection is based on fluorescence intensity. Although the sensitivity 586 for detection is largely improved by the use of high power lasers and 587 by increasing the dwell time of the vesicles in the laser beam, particles 588 with low fluorescence intensity (e.g. due to low PKH67 labeling efficien-589 cy or because of small size) may not be detected using this technique. 590 Furthermore, the removal of unbound fluorescent dye by density gradi-591 ent ultracentrifugation can be seen as a time-consuming procedure. 592 However, the same procedure also allows separation of EVs from 593 protein aggregates that are abundantly present in culture media and 594 body fluids. This is essential, since such aggregates can mistakenly be recorded as vesicles by the technologies discussed here. 596

Comparability analysis of the three techniques indicated that 597 substantially larger differences in quantification were obtained for 598 liposomes, compared to EVs (Fig. 4A and D). In fact, no significant differ- 599 ence in raw concentration estimation was observed for quantification of 600 EVs by tRPS versus hFC. One potential explanation is that the EVs exhibit 601 higher fluorescence levels compare to the liposomes, either because EVs 602 are larger in size and incorporate more dye or because of differences in 603 labeling efficiency. Differences between the other instruments are 604 difficult to account for. For both liposome batches and EVs, higher raw 605 concentration estimations were obtained by NTA compared to tRPS. 606 We tested whether background particle detection (from the buffer in 607

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which particles were diluted) could explain the observed differences. However, measurement of PBS background particles at camera-level 9 and 12 revealed maximum concentrations of only 2.60×10^7 /ml and 3.3×10^7 /ml, respectively (data not shown), accounting for 3.29% (L146) and 2.33% (L212) of the measured concentrations of the liposomes.

So far, only one other study has directly compared NTA, tRPS and 614 flow cytometry (using a different high-end flow cytometer) by analyz-615 616 ing the size distributions of polystyrene beads and urine-derived EV [36]. Interestingly, comparable EV quantifications by NTA and tRPS 617 618 were reported, whereas the flow cytometry-based EV quantification 619 was 15 times lower. However, a direct comparison of these data to our 620 current study is difficult, because a crude preparation (1550 \times g centri-621 fugation followed by 0.2 µm filtration) of EVs from a different biological source (urine) was analyzed and because the flow cytometric measure-622 ments in that study were light scatter-based. However, it is interesting 623 to note that, in contrast to our findings, EV quantifications by NTA and 624 tRPS were found to be comparable. This could imply that the type of 625 EV and the degree of EV purification may also influence quantification 626 by the different instruments. 627

Several strategies have previously been suggested to calibrate 628 particle quantification in EV samples. For tRPS we spiked biological 629 630 fluids with polystyrene beads of known size and concentration to 631 improve EV quantification accuracy [20]. For NTA, on the contrary, this approach seems less suitable since the methodology does not 632 allow for accurate discrimination of particles of interest from beads 633 with a similar size [19,22,34]. Secondly, spiking a sample with large 634 635 (>500 nm) silica beads could lead to over-scattering of the EVs and skew characterization [22]. An alternative that has been proposed for 636 NTA calibration is application of a correction factor, based on the 637 measured concentration of silica beads compared to the expected con-638 639centration of these beads [19]. Although promising and potentially 640 valuable for measuring relatively homogenous populations of EVs, 641 such a calibration method is unsuitable for analysis of the heterogeneous EV preparations studied here. As NTA is less accurate in the detec-642 tion of size-based subpopulations [19,22,34], one would have to apply a 643 multitude of silica calibration beads, each covering a subpopulation of 644 645 EVs and subsequently aggregate analysis of these subpopulations. More research into the accuracy of such a calibration system will be 646 essential before it can be broadly applied. 647

In conclusion, we identified NTA and tRPS instrument settings that 648 649 affect particle quantification and showed that the impact of these parameters on quantification varies with the types of nano-sized 650 particles analyzed (i.e. polystyrene beads, liposomes and EVs). Our 651 data clearly indicate that absolute quantification of EVs and liposomes 652substantially differs using the three different technologies and that a 653 654golden standard for quantification of such particles is not available yet. Moreover, our data strongly underline the importance of technical 655 knowledge of the instruments for correct data interpretation, and 656 plead for awareness of the effects of instrument settings in case vesicle 657 populations with unknown concentration and size heterogeneity are 658 659 measured. Increased understanding of the possibilities and pitfalls of 660 these technologies will benefit standardized and large-scale clinical application of (engineered) EVs and EV mimics in the future. 661

Q4 6. Uncited references

663 [14,15,16,17,32,33,47,48]

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Appendix A. Supplementary data

Supplementary data to this article can be found online at http://dx. 673 doi.org/10.1016/i.jconrel.2014.12.041. 674

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