1 Biophysical analysis of lipidic nanoparticles.

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7 Highlights

- 8 Biological nanoparticles play key roles in research and therapeutics.
- 9 A range of techniques are available to study such particles in terms of size and
 10 concentration.
- Each technique has associated advantages and disadvantages.
- Novel approaches allow the analysis of the full range of nanoparticles with a single
 instrument.

14 Abstract

15 Biological nanoparticles include liposomes, extracellular vesicle and lipid-based discoidal systems. 16 When studying such particles, there are several key parameters of interest, including particle size and 17 concentration. Measuring these characteristics can be of particular importance in the research 18 laboratory or when producing such particles as biotherapeutics. This article briefly describes the major 19 types of lipid-containing nanoparticles and the techniques that can be used to study them. Such 20 methodologies include electron microscopy, atomic force microscopy, dynamic light scattering, 21 nanoparticle tracking analysis, flow cytometry, tunable resistive pulse sensing and microfluidic 22 resistive pulse sensing. Whilst no technique is perfect for the analysis of all nanoparticles, this article 23 provides advantages and disadvantages of each, highlighting the latest developments in the field. 24 Finally, we demonstrate the use of microfluidic resistive pulse sensing for the analysis of biological 25 nanoparticles.

26 Keywords: Nanoparticle, MPRS, nCS1, Extracellular vesicles, Liposomes

27 List of abbreviations

ACdEV	Apoptotic Cell-Derived Extracellular Vesicle
AFM	Atomic Force Microscopy
An-V	Annexin V
API	Active Pharmaceutical Ingredient
BSA	Bovine Serum Albumin
CCD	Charged-Coupled Device
CSD	Concentration Size Distribution
DDS	Drug Delivery System
DLS	Dynamic Light Scattering
EM	Electron Microscopy
ET	Electron Tomography
EV	Extracellular Vesicle
FCM	Flow Cytometry
	Extracellular Vesicle Flow Cytometry
FSC	Forward Scatter
HRFCM	High-Resolution Flow Cytometry

HRTEM IC LET MRPS MSP MVB NP NTA PS PZT RI RPS SEM SFM	High-Resolution Transmission Electron Microscopy Intracellular Communication Liposomal Encapsulation Technology Microfluidic Resistive Pulse Sensing Membrane Scaffold Protein Multivesicular Body Nanoparticle Nanoparticle Tracking Analyser Phosphatidylserine Piezoelectric Refractive Index Repulsive Pulse Sensing Scanning Electron Microscopy Scanning Force Microscopy
	•
SIOS	Scanning Ion Occlusion Spectroscopy
SMALP	Styrene Maleic Acid Lipid Particle
SSC	Side Scatter
TDFM	Transverse Dynamic Force Microscopy
TEM	Transmission Electron Microscopy
TRPS	Tunable Resistive Pulse Sensing
UC	Ultra-Centrifugation
3-D	3-Dimensional

29 Introduction

30 Nanoparticles

31 Nanoparticle (NP) is a term which encompasses several subpopulations. Nanoparticles can be 32 naturally occurring, for example, extracellular vesicles (EVs) released by cells across the three main 33 groups of the phylogenetic tree: Archaea, Bacteria and Eukarya. They can also be engineered, as with 34 viruses used for vaccines; they can be biological (for example liposomes) or non-biological (such as 35 metal or metal-derived NPs) or a combination of both. They could also be accidental in their 36 production, such as exhaust fumes. In this review, we will use the term "nanoparticle" to refer to 37 extracellular vesicles, liposomes and nanodisc systems, e.g. styrene maleic acid lipid particles 38 (SMALPs).

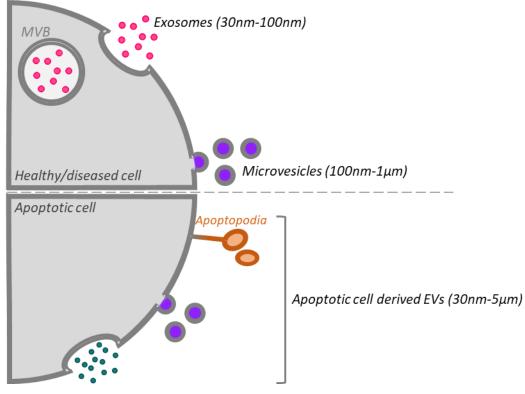
- 39 Whilst the international standards organisation (ISO)[1] defines nanoparticles as ranging in size from
- 1 to 100nm, it is generally accepted that biological nanoparticles may be larger than this within the
- 41 sub-micron range. Due to their size, NPs have a larger surface area to volume ratio when compared
- 42 to bulk materials, which give them interesting physical properties. Their size and composition can
- 43 make them challenging to study experimentally.

44 Extracellular vesicles

- Extracellular vesicles (EVs) are membrane-surrounded structures released by healthy, apoptotic and
 diseased cells. They were first described in 1967 by P. Wolf [2] and are found in most biological fluids.
 They are a highly heterogeneous population which vary in size, content, and mechanism of formation
 [3]. Their origin can also be used to classify them further. The field acknowledges three distinct
 subpopulations: apoptotic vesicles (30nm-5µm), microvesicles (100nm-1µm) and exosomes (30nm100nm) (Figure 1). Exosomes are initially formed inside multivesicular bodies (MVBs) and are further
 released through exocytosis into the extracellular environment. Alternatively, microvesicles and
- 52 apoptotic bodies are produced through budding of the plasma membrane.

Apoptotic cell-derived EVs (ACdEVs) encompass apoptotic bodies, apoptotic microvesicles and exosome-like vesicles. Evidence suggests that ACdEV biogenesis seems to be more complex as they can be formed through membrane protrusions termed apoptopodia [4], or through plasma membrane blebbing as mentioned previously. However, as little is reported in the literature regarding ACdEVs,

57 other novel mechanisms of biogenesis are worth investigating.



59 *Figure 1 Extracellular vesicles formation and different subpopulations.* Healthy and diseased cells 60 release two mains forms of EVs named exosomes and microvesicles. They are released through the

61 exocytosis of multivesicular bodies (MVBs) and via plasma membrane shedding. Apoptotic cells release

62 ACdEVs of various size through membrane protrusion, membrane blebbing and other mechanisms.

63 Green vesicles represent exosome-like ACdEVs.

64 It was first believed that EVs were cellular debris, but subsequent research has shown they contain a 65 specific subset of proteins, mRNAs, miRNAs, and lipids rather than random cellular components [5]. It 66 has been demonstrated that EVs play a major pathophysiological role in various biological process 67 (apoptotic cell clearance, infection, immune response, antigen presentation [6]). They function to 68 transport and deliver cargo molecules between cells [7]. This EV-mediated intercellular 69 communication (IC) exists in healthy tissues and when altered, can lead to a variety of diseases, namely 70 autoimmune diseases and cancer. In the past few decades, EVs have received a great deal of interest 71 for their diagnostic and therapeutic potential [8].

72 Liposomes

58

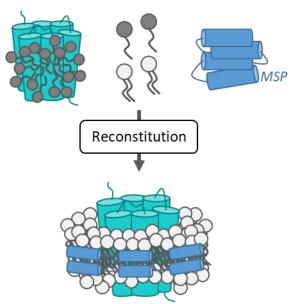
Liposomes were first discovered by Bangham and his co-workers in 1961 [9], and can be described as spherical artificial vesicles of various lamellarity: usually one or two phospholipid bilayers. Liposomes are formed spontaneously, but many different methods of preparation exist. These can be divided between passive or active loading techniques. The most common passive techniques are mechanical dispersion, solvent dispersion or detergent removal, such as sonication and extrusion [10]. Similar structures can also be formed from polymers (sometimes referred to as polymersomes) [11].

Conventional liposomes are commonly made of ester phospholipids such as phosphatidylcholine. A new generation of liposomes have modified formulations in order to try to overcome instability issues and enhance specificity [12]. For example, archeasomes contain one or more lipids found in archeobacterial membranes that offer higher stability because of di-ether linkages [13]. Additionally, virosomes are a combination of liposomes with virus-derived envelope proteins, allowing the fusion

- of the virosomes with the target cell [14]. Finally, novasomes are paucimellar vesicles formed of two
 to seven bilayer membranes commonly used in cosmetics [15].
- Liposomes can be loaded with different drugs and or molecules and used as drug carriers to deliver a range of active pharmaceutical ingredients (API) to a site of action for therapeutic and diagnostic purposes [16]. This liposomal encapsulation technology (LET) can involve drug molecules, gene therapies and bioactive agents. Hydrophobic API can be incorporated into the phospholipid bilayers, whereas hydrophilic API can be encapsulated inside the aqueous centre. LET has been shown to increase the stability and decrease the toxicity of the active molecules making it more efficient [17].
- 92 Different types of liposomal drug delivery system (DDS) exist [18], for example, by coupling the
- 93 liposomes to targeting ligands to increase the specificity. These carriers are most valuable because of
- 94 their biocompatibility and non-immunogenicity [19].
- 95 Another use for liposomes is their relevance as a biological model [20],[21]. They can be used to mimic
- 96 both extracellular vesicles and cells, offering a non-invasive alternative to study biological processes
- 97 and test drugs. Their composition, size and charge can be tuned to make a highly accurate model.

98 Disc systems

- 99 In membrane protein research, proteins often have to be isolated from the membrane in order to
- study them. This has often resulted in the use of detergents to solubilise the membrane, and therefore
- 101 the protein, into detergent micelles. Throughout the past decades, it has become more apparent the
- 102 native lipid environment surrounding the protein plays numerous roles in the structural stability and
- 103 functionality of the protein. Hence, alternatives to detergents are advantageous [22].
- 104 Membrane scaffold proteins (MSPs) are one of the alternatives to a purely detergent-based isolation
- system. Bayburt et al., 2002 [24] first described the formation of bilayer patches surrounded and
- stabilised by a ring of amphipathic helical protein, termed membrane scaffold proteins. The process
- 107 of forming these patches was described as a self-assembly mechanism whereby lipids, MSPs, and
- 108 detergent were added to protein-detergent micelles forming particles of around 10nm in size [25]
- 109 (Figure 2).



- **Figure 2 Schematic demonstrating the formation of MSP nanodiscs**. A protein is isolated from the
- 111 membrane via detergent solubilisation. A mixture of lipids and membrane scaffold proteins (MSPs) are
- added to the protein-detergent micelle solution, this results in the self-assembly of MPS encapsulated
- 113 protein-lipid nanodiscs.

- 114 A more recent approach is the use of styrene-maleic acid (SMA) copolymers and derivatives to isolate
- membrane proteins in a nanometer-sized disc, known as SMA lipid particles (SMALPs) (Figure 3). The
- technique was first described in 2009 by Knowles *et al.* [26] and has the distinct advantage that the
- 117 protein is extracted directly from the membrane without the need for detergent, and therefore
- 118 maintains a near-native lipid environment, which is beneficial for downstream research [26].

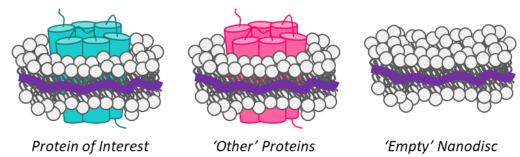


Figure 3 Schematic of SMALP encapsulated nanodiscs. SMA copolymers punch holes in the membrane
 encapsulating anything that is present, i.e. membrane proteins (protein of interest as well as other
 proteins) or 'empty' lipid only discs.

122 The greatest challenge of single NP based isolation and analysis is still the size of the NPs, which are

123 below the reach of conventional detection methods [27]. In this review, we will present the main

124 methods for NP analysis and their limitations, focusing on single particle-based quantification and size

125 profiling. Finally, an advancement in NP analysing technology will be introduced as an alternative to

126 study NPs with preliminary data presented.

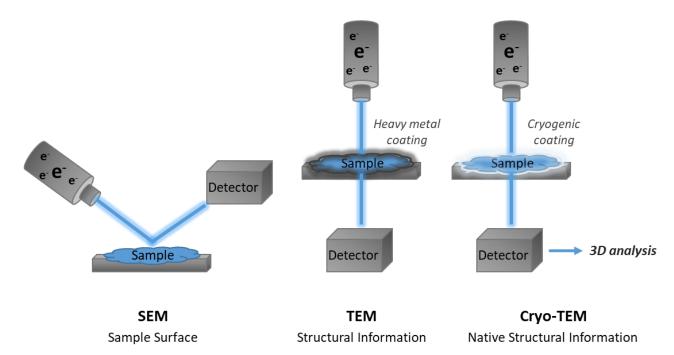
127 Conventional methods for NP analysis

Common isolation methods for NPs include; ultracentrifugation (UC) and precipitation alongside size-, immunoaffinity capture-, and microfluidic-based techniques [28]. Microfluidic isolation is a promising field, which offers miniaturisation of conventional techniques, quicker purification times and improved resolution while enabling a continuous separation of the NPs. Following isolation using any approach, it is often desirable to analyse the size of the NPs to ensure either efficient separation or to define the nature of the sample.

134 Electron microscopy (EM)

Conventional optical microscopes cannot be used due to NPs being below the diffraction limit of visible light (200nm). Traditionally, the only way to visualise NPs via microscopy is by using the shorter wavelength and higher frequency of an electron microscope. As the wavelength of an electron can be up to 100,000 times shorter than that of visible light photons, electron microscopes have a higher resolving power that can reveal the structure of smaller objects, i.e. in the nanoscale. The first electron microscope was built in 1931 by Ruska and Knoll. Nowadays, two types of electron microscopes exist

141 Scanning EM (SEM) and Transmission EM (TEM) (Figure 4).



142 Figure 4 Simple schematic of the most common EM techniques. Scanning electron microscopy

143 determines information about the sample surface. Transmission electron microscopy sends electrons

144 through the heavy metal coated sample, acquiring structural information. Cryo-transmission electron

145 microscopy utilises freezing of the sample to retain native structural information.

SEM is an electrical conductivity-based EM where the electron beam is focused onto the surface of the samples, scanning it line by line. A detector simultaneously counts the scattered secondary electrons. The images acquired by SEM only allow visualisation of the surface of the specimen, but SEM can also provide 3-dimensional (3D) images. However, this method offers a lower resolution than TEM [29].

TEM is an electron emission-based EM for thin samples (~100nm). Here, the beam of accelerated electrons passes through the samples. The detector is below the sample retrieving the electron to provide structural information. The entire electron path through the column must be under vacuum. High-resolution TEM (HRTEM) is now able to provide a resolution below 0.5 angstroms [30]. However, this resolution is limited to non-biological samples, regarding lipidic nanoparticles the resolution of ~1.5 angstroms is more accurate [31].TEM can have variants such as immune-EM, Cryo-EM (single-particle and tomography), 3D-EM and conventional TEM.

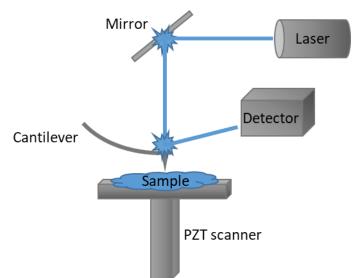
For conventional TEM, the thin sections are coated with heavy metals in order to obtain a better contrast and to visualise the lipid bilayer [32]. Depending on where the section was made, the diameter can differ. The morphology of the whole particle thus remains uncertain. Fixatives can be added to improve the sample retention. This method is a rapid way of confirming the presence of NPs in a sample.

The least invasive technique is cryo-TEM, where the sample is frozen using a cryogen (liquid ethane or propane) (Figure 4). Upon rapid freezing, a thin liquid film of the buffer is transformed into an amorphous solid, allowing the object beneath to be observed in the native frozen-hydrated state. The immediate freezing of the sample allows no time for crystal formation, removing the need to use fixative and heavy metals. The sample is then transferred in a cryo-electron microscope and observed at low temperature (-170 °C). The lipid bilayer appears as two thin dark lines because of the phosphorous atoms (the heaviest) scattering more electrons [33].

- 170 Phenotyping of EVs can be performed using cryo-TEM combined with immune-gold-labelling. Gold
- particles provide high contrast in EM thanks to their large electron scattering properties (i.e. plasma
- 172 EVs exposing phosphatidylserine (PS) labelled with annexin V (An-V) conjugated with gold particles)
- 173 [34]. Additionally, electron tomography (ET) can be used for 3D reconstruction based on 2-174 dimensional pictures taken at many different angles creating a tilt series. This can be combined with
- 175 cryo-TEM by creating electronic slices as thin as 1nm through the reconstructed 3-D volume [35]. The
- 176 details are much finer compared to cryo-EM alone and allow discrimination of vesicles spatially below
- 177 or inside another one. The angles of the image do not yet permit a fully 3D structure due to an artefact
- 178 that is referred to in the literature as "the missing wedge" [28, 29].
- 179 TEM has some potential disadvantages for particle analysis including possible radiation damage to the 180 sample such as re-conformation and de-crystallisation to the breaking of atom bonds, removal of side-181 chains and in general a loss of mass [38].

182 Atomic force microscopy (AFM)

- 183 Atomic force microscopy (AFM) is a type of scanning force microscopy (SFM) developed in the 1980s
- by Binning, Quate, and Gerber [39]. Typically, a near-horizontal cantilever is used with a "nanofiber"
- tip facing towards the sample. Lasers are then reflected off the back of the cantilever to track the
- 186 change in its displacement, which depending on its mode of operation, provides different information
- 187 (Figure 5).
- 188



- **Figure 5 Schematic representation of Atomic Force Microscopy.** The cantilever is in contact with the sample and the sample stage (piezoelectric (PZT) scanner) moves the x, y, and z position. A laser monitors the position of tip with feedback via a photodiode to the piezoelectric scanner to move the stage z-position to build a topological image at atomic resolution.
- 193 There are three modes of AFM; the first mode is contact-based, where the sample stage is positioned
- 194 to provide a small upwards deflection of the tip when contact occurs. The sample is then moved along
- 194 to provide a small upwards deflection of the tip when contact occurs. The sample is then moved along
- the x- and y-axis. As the topology of the sample changes so does the level of cantilever deflection,
- some operate by simply tracking the height of the cantilever in absolute terms and generating the
- topology from this [40]. In others, the system will for each x and y co-ordinate aim to deliver a
- 198 consistent level of deflection, and the required changes in the samples z position will generate the
- 199 topological map.

- 200 The second mode is non-contact. Here instead of an upwards deflection, a downwards deflection is
- observed. This is due to atomic forces, e.g. Van der Waals on this scale interacting with the cantilever,
 pulling it towards the sample [41]. Again, the concept is to maintain a consistent deflection and
 generate a topological map through record stage positioning.

There is a third non-contact mode. This involves the cantilever being oscillated at very high frequencies. As the tip gets close to the sample, either atomic forces or physical interactions, often with a layer of moisture present, leads to a dampening of the oscillation frequency. A consistent level of damping is required to generate the topological map [42].

Some advantages are that AFM can be operated in ambient conditions and often requires no special preparation of the sample. The sample sizes are often small, e.g. <200nm squared. Advances in piezoelectrics and control systems allow for near real-time scanning to generate videos, particularly useful for studying biological samples [43],[44]. Improved methods are always being developed, e.g. Transverse dynamic force microscopy (TDFM) which uses a vertical cantilever [45].

213 Dynamic light scattering (DLS)

214 Dynamic light scattering (DLS) is one of the most common and simple techniques used in the analysis

of particle size distribution [46]. DLS is a measure of time-dependent fluctuations in the scattered light

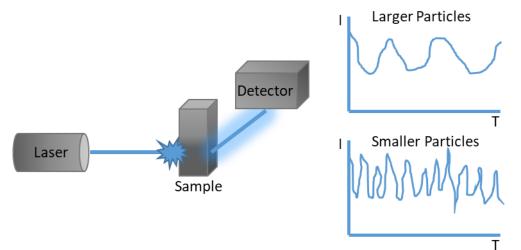
of particles undergoing Brownian motion (diffusion caused by random collisions with solvent

217 molecules). A particle's hydrodynamic diameter is calculated as a function of the diffusion coefficient.

218 Where the diffusion coefficient is inversely proportional to the particle size according to the Stokes-

219 Einstein equation (Figure 6). This calculation assumes that all particles are spherical, reporting an

220 equivalent particle diameter which can be disadvantageous for studying discoidal systems.



221 Figure 6 Schematic of Particle Size Analysis by Dynamic Light Scattering. A laser illuminates particles

222 undergoing Brownian motion in a sample. A photodetector measures the fluctuations in the intensity

of scattered light over a time period. Stokes-Einstein's equation is used to calculate the hydrodynamic

224 diameter of the particles.

225 DLS can analyse particles ranging from 0.5 to 10,000nm dependent on the instrument and can analyse 226 as little as 45μL of the sample, although common practice is to use a standard size cuvette which

requires around 500μL for analysis. DLS also uses Rayleigh's approximation, where the intensity, I, of

scattered light is proportional to the sixth power of the particle diameter, d: $I \propto d^6$ (i.e. a particle of

100nm will scatter light with a million times the intensity of a 10nm particle)[46],[47]. This means that

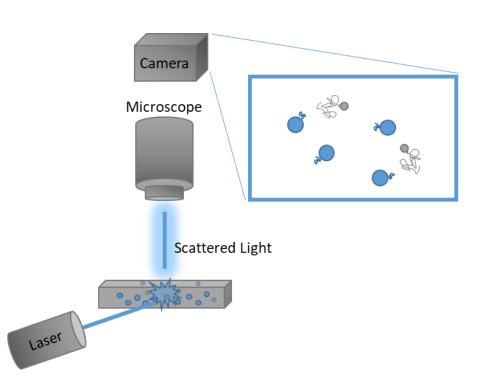
DLS has an inherent bias towards larger particles and therefore, a tendency to oversize particles in a
 polydisperse mixture and as such, it is not suitable for highly polydisperse samples.

232 Nanoparticle tracking analyser (NTA)

233 Nanoparticle tracking analysis (NTA) can be used for various applications, including but not limited to 234 vesicles, exosomes, and proteins [48]. NTA uses the same underlying principle as DLS as it measures 235 time-dependent fluctuations in scattered laser light of particles undergoing Brownian motion to determine their hydrodynamic radius. Where NTA differs is how it detects the scattered light. NTA 236 237 uses a laser beam to illuminate particles, and the scattered light is easily visualised with a conventional microscope equipped with a 20x objective lens. Particle movement, i.e. scattered light, is then 238 239 recorded with a light-sensitive charged-coupled device (CCD) or CMOS camera, arranged at a 90° angle 240 to the irradiation plane [49]. The camera operates at 30 frames per second (fps), capturing a video file 241 of the particles moving under Brownian motion. The software tracks many particles individually and 242 using the Stokes-Einstein equation calculates their hydrodynamic diameters (Figure 7) [50]. As NTA is 243 capable of tracking individual particles, the concentration of particles within a sample can be calculated. However, this requires calibration with standards of a known size and concentration. 244

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Figure 7 Schematic Representation of Particle Size Analysis by Nanoparticle Tracking Analysis. A
 laser illuminates particles undergoing Brownian motion in a sample. Particle movement i.e. scattered
 light is then recorded with a light-sensitive charged-coupled device (CCD) or CMOS camera. Software
 tracks individual particles and uses the Stokes-Einstein equation calculates their hydrodynamic
 diameters.

- NTA is reported to size particles between 10 and 1000nm at a concentration range of 1E+06 to 1E+09
 particles.mL⁻¹; however, this is sample and system configuration dependent [50]. For samples (such as
 EVs), where the refractive index (RI) of the particle is close to the RI of the electrolyte it is suspended
- 256 in, the detection resolution is compromised, resulting in a restricted size analysis range and unreliable

concentration determination [51]. Like DLS, NTA also has restricted resolution for polydispersesamples due to Rayleigh's approximation, making it difficult to detect and track small particles [46].

259 Conventional Flow Cytometry (FCM)

Flow cytometry (FCM) is a laser-based technology developed for high throughput, multiparametric analysis of individual particles (e.g. cells). It allows the analysis of thousands of cells per second. However, in order to be detected, particles need to scatter light and/or induce fluorescence. NPs are much smaller than the cells for which flow cytometry was designed. Conventional FCM has a lower detection limit for polystyrene beads between 200 and 500nm [52]; consequently, the detection of NPs remains a challenge. Recently, dedicated small particle high-resolution flow cytometry (HRFCM) can offer a better resolution down to 40nm [53].

- Cells and NPs are guided through a laser beam. The first detector is parallel to this beam and measures
 forward scattered light (FSC). The other detector is perpendicular to the beam and measures the side
- scattered light (SSC). FSC is used to look at the specimen size, whereas SSC is used to look at its
- 270 granularity (Figure 8). The smaller the particle, the less the light scatters and the more background.
- FSC depends on the radius, the illumination wavelength, the refractive index and the light collection
- angle [54]. Thus, for an identical size, particles can have various refractive indices, resulting in a
- 273 different scattering (Figure 9A). For the analysis of EVs, it is better to use silica beads for calibration,
- as EVs refractive index is closer to that of silica beads than polystyrene beads.

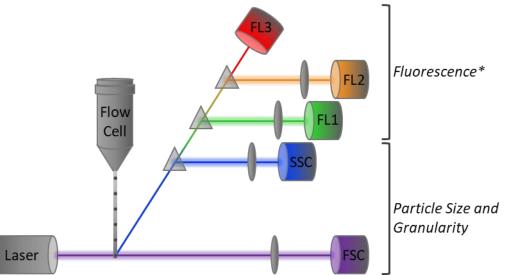


Figure 8 Schematic Representation of Particle Analysis by Flow Cytometry. A particle suspension
 enters the flow cell where the particles are aligned into a single stream and pass through a focused
 laser-beam. The resultant signal provides information on their size/granularity based on their forward
 and side scatter and the fluorescent intensities (*if fluorescently labelled) of each particle.

Fluorescent detection of surface proteins can also be achieved as long as the labelling of the NP is bright enough to discriminate NPs from the optical and electronic noise. To do so, the light scatter

threshold can be modified, or the excess of label can be removed using a density gradient. Focusing

on EVs, the fluorescence trigger can be used as long as the antigen is abundant enough to allow

detection. This can be an issue for smaller particles with fewer copies of the antigen of interest [55].

284 Thus, sensitivity limits should be kept in mind when using fluorescently-labelled antibodies.

The concentration of the particles can have an important impact on both scatter and fluorescence [56]. If the concentration is too high, it will result in a swarm detection (Figure 9B). This occurs when two or more particles are in the measurement volume when the data acquisition is triggered, resulting in multiple NPs counted as a single event [57]. In order to have a better single NP detection, the samples need to be carefully diluted to an appropriate particle concentration. FCM offers a great possibility for single NP based analysis as long as there is a standardisation of the sample preparation as well as the appropriate instrument calibration.

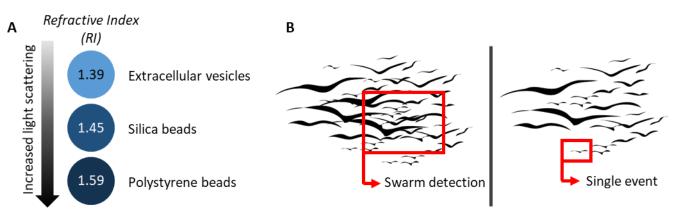


Figure 9 Principles of flow cytometry for NP analysis A. Effect of the refractive index (RI) of different
 nature of nanoparticles on the light scattering. B. Representation of particle flowing: swarm detection
 in opposition to a single event detection. Individual seagull representing an individual particle.

295 Tunable resistive pulse sensing (TRPS)

Tunable resistive pulse sensing (TRPS), previously known as scanning ion occlusion spectroscopy (SIOS)
[58], is suitable for biological and inorganic NP size, concentration and charge analysis. The equipment
most largely associated with TRPS is produced by Izon Science Ltd and can be referred to as qNano.

299 TRPS requires a nanosized stretchable pore that separates two conductive fluid compartments. After 300 applying a voltage across the nanopore, a current is established which is disrupted by the movement 301 of particles through the nanopore. When a particle moves through the opening, it decreases the flow 302 of ions through the nanopore, causing a transient current decrease known as a blockade event [59]. 303 The amplitude of this event provides information for the size profiling of the particle. The number of 304 blockades throughout time gives information about the concentration (Figure 10). NP measurement 305 using TRPS requires an initial run with a calibration sample of known diameter and concentration, 306 usually polystyrene beads of a modal size appropriate to the range of interest [41,42]. Samples and 307 calibration should ideally be performed in the same buffer, which can cause issues with biological 308 samples. Spiking the sample directly with calibration beads offers an alternative calibration method.

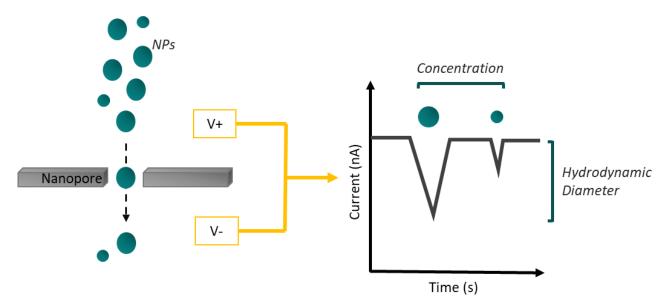


Figure 10 Principle of the TRPS technique to measure the diameter and concentration of
 nanoparticles. The magnitude of the blockade event gives information on the hydrodynamic diameter,
 whereas the number of blockades over time gives information on the concentration.

312 The size range of the nanopores can be tuned depending on the NPs of interest. Currently, the smallest

nanopore is the NP100 that has a size range of 70-200nm. However, the full range of manufactured

tunable pores extends from 70nm to 10μm. The measurement settings are flexible between each user.
 Different parameters can be adjusted in order to have the best measurement conditions, such as the

316 nanopore stretch, the applied voltage and the pressure. The same parameters must be used between

measurements to ensure the reproducibility of the data. A total of 500 particles or more is then

318 counted per measurement. The TRPS technique provides a particle-by-particle analysis of the size and

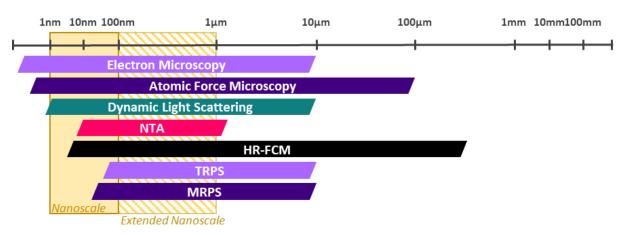
319 concentration using tunable nanopores. However, it can be challenging due to the size heterogeneity

320 of the samples that can often lead to the blockage of the nanopore; this can be especially true for

321 biological samples.

TECHNIQUE	ADVANTAGES	DISADVANTAGES
EM	High resolution Morphology 3D Contrast	Expensive equipment Specific training and expertise Radiation damage Sample preparation Invasive/destructive technique Highly time consuming
AFM	Atomic resolution Morphology/ surface roughness Sample preparation	Expensive equipment Specific training and expertise Highly time consuming
DLS	Quick Easy to use Relatively inexpensive High throughput	Low resolution in polydisperse samples Tendency to oversize particles due to Rayleigh's approximation of scattered light
ΝΤΑ	Quick Easy to use Relatively inexpensive High throughput	Data bias dependent on user settings Dependent on refractive index Low resolution in highly concentrated polydisperse samples due to Rayleigh's approximation. Moderately time consuming
FCM	Full NP diameter range High throughput Multiparametric analysis	Low resolution Risk of swarm detection Low sensitivity no information about morphology Moderately time consuming
TRPS	Inexpensive Reusable pore Single particle-based analysis High throughput	Easily clogged nanopores Calibration required Moderately time consuming Required standardisation

- 323 Table 1 Comparison of the conventional techniques used to characterize the size and concentration
- 324 of NPs.



325

326 Figure 11 Size ranges for particle analysis of discussed techniques.

Each technique for NP characterisation has different limitations and advantages (Table 1 and Figure 11) [62]. Thus, it is often recommended to combine several techniques instead of choosing one method, as there is no standard procedure for the characterisation of NPs to date.

331 Recent Advances in Nanoparticle Analysis

332 Microfluidic resistive pulse sensing

Microfluidic resistive pulse sensing (MRPS) is a promising advancement in the resistive pulse sensing techniques originally described in the Coulter Principle over half a century ago [63]. The last decade has seen advancement in the form of TRPS (see the previous section). Although TRPS has pushed the limits of size detection of the resistive pulse sensing (RPS) technique, it still has limitations which need to be addressed for research to progress. In recent years there has been rapid development in the manufacturing of microfluidic chips and devices. An American company, Spectradyne, has utilised this research to develop an RPS instrument with patented microfluidic technology, the nCS1 [64].

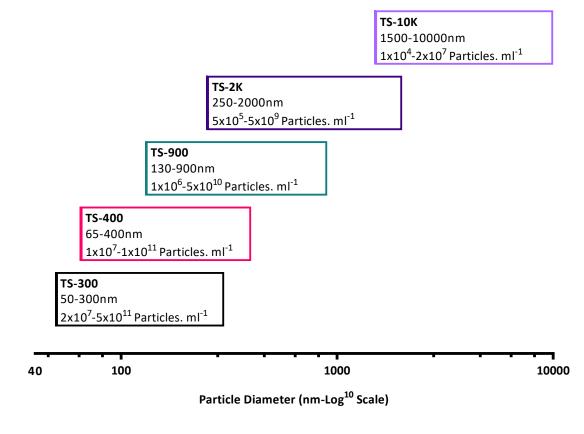
The microfluidic technology in question is a series of pre-calibrated, single-use, low volume microfluidic cartridges. Each cartridge has a distinct size and concentration range (Figure 12). This has

the benefits of making the technique an absolute method, i.e. no need for calibration material, the

disposable microfluidic cartridges also dramatically reduce the risk of sample cross-contamination,

with the added advantage of only requiring 3µL of sample to gain reliable, reproducible, statistically

345 significant results.





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346

The principle of the technique is essentially the same as that described by Wallace Coulter [63], whereby, a voltage is applied across an aperture (hole) submerged in an electrolyte, as a particle passes through the aperture (also termed a nano-constriction) the displacement of the electrolyte causes a voltage drop which is proportional to the size/volume of the particle, enabling its hydrodynamic diameter to be calculated [63],[65],[66]. This allows the user to obtain analytical data for individual particles making the technique highly precise.

The microfluidic cartridge of Spectradyne's nCS1 also significantly reduces one of the major issues of the RPS technique, which is aperture blockages by incorporating a sample pre-filter embedded within the microfluidic cartridge. The only drawback with the incorporation of this pre-filter is overcoming the surface tension in the smallest (TS-300/-400) size cartridges. Spectradyne recommends the addition of a surfactant (polysorbate-20/ Tween-20) at 1%(v/v) to the sample or 0.1%(w/v) bovine serum albumin (BSA) as a wetting agent where surfactant cannot be used (i.e. with lipidic samples).

The technology also overcomes some of the more technical aspects of the technique, such as distinguishing agglomerated particles from monodisperse particles. It does this in the form of filters applied to the data with parameters specific to each of the different sized cartridges, such as transit time, the time it takes the particle to pass through the aperture, and symmetry.

366 Analysis of nanoparticles using the nCS1

367 To test the capabilities of the instrument, a polydisperse sample of polystyrene size standards 368 (Nanospheres) were analysed (Figure 13A). A broad range of Nanospheres were used (see Figure 13B) 369 this was to test the accuracy of the instrument over multiple size cartridges. The Nanospheres were 370 used at an approximate concentration of 4.2E+8 particles.mL⁻¹, i.e. a total concentration of 2.52E+9 371 particles.mL⁻¹ in the polydisperse sample, this is in line with the acceptable concentration range for 372 both the TS-400 and TS-2K cartridges used in the analysis. Approximately 2500 particles were analysed 373 with default cartridge filters applied (Table S1); the subsequent data was analysed graphically by 374 determining size ranges for Gaussian analysis/fit (Table S2).

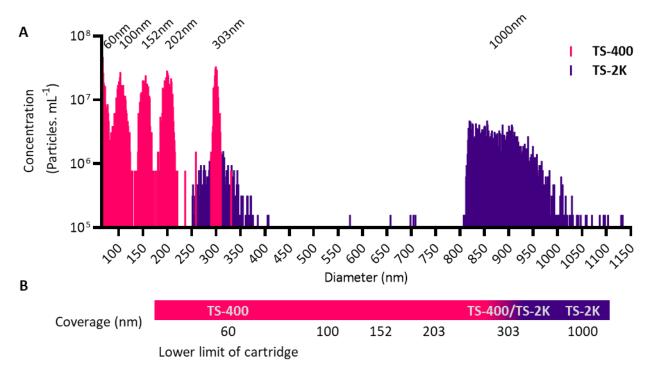


Figure 13 Analysis of a polydisperse polystyrene size standard sample using the nCS1. A. Size
distribution data demonstrating clear separation of peaks corresponding to the nominal size range of
the polystyrene standards, the TS-2K cartridge perfectly overlays with the TS-400 cartridge at 303nm
however demonstrates a broader distribution. B. Schematic demonstrates the size standards used in
the polydisperse mix and the theoretical coverage of the cartridges.

380 The data demonstrates that the nCS1 is capable of discerning multiple peaks in a single polydisperse 381 sample. The TS-400 coverage range starts at 65nm; therefore, the resolution of the 60nm Nanosphere is hindered, resulting in a slight oversizing of the particle (Nanosphere[™] NIST traceable certified 382 383 diameter of 60±4nm), these data would be best captured using a TS-300 cartridge which has a lower 384 detection threshold of 50nm. The other peak that demonstrated variability was the 1000nm peak; this 385 showed a broad peak with a mean of 873.3nm (Figure 13, Table S1). The remaining discernible peaks 386 fall within their certified mean diameter (provided on the safety data sheet), demonstrating the 387 accuracy/resolution of the nCS1.

388 Analysis of Extracellular vesicles using the nCS1

389 Following analysis of the polystyrene size standards, we sought to test the instrument's capabilities at 390 detecting a biological sample. Extracellular vesicles derived from apoptotic Jurkat cells were analysed 391 for both size range and concentration. The apoptotic cell-derived EVs (ACdEVs) were isolated using 392 centrifugation at 2000xg, and therefore it was suspected that the EVs could vary greatly in size; as 393 such analysis using both a TS-300 and TS-900 was conducted. The analysis was conducted without the 394 recommended surfactant (polysorbate-20) over concern that it could cause damage/ change the 395 integrity of the EVs. Acquisition with the TS-900 demonstrated a continuous size distribution with a 396 higher concentration of EVs at the lower size limit; this is in line with what was expected (

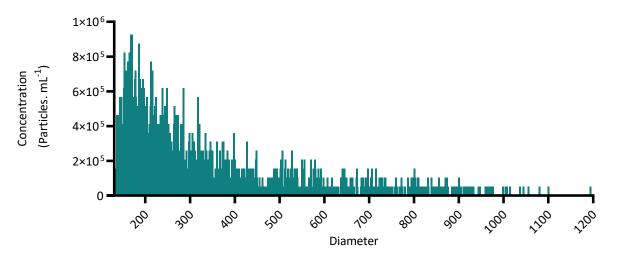
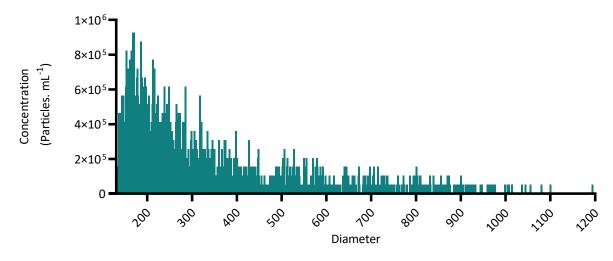




Figure **14**). Analysis across the entire TS-900 detection range (130-1000nm) reported a total of 2146 particles analysed in a time period of three minutes 30 seconds, demonstrating the high throughput capacity of the instrument. The concentration of particles across this range was calculated at 1.1E+08 particle.mL⁻¹, with a D10, D50, and D90 (corresponding to 10%, 50%, and 90% of particles in the concentration size distribution (CSD), see



404 Figure **14** for CSD) of 164.8, 269.2, and 669.4nm respectively.

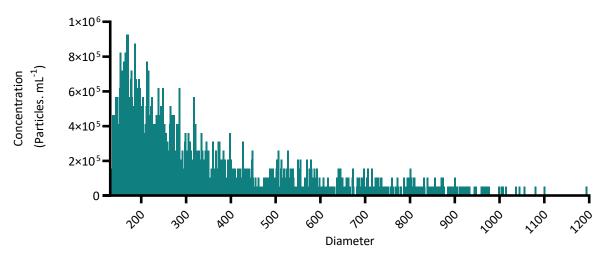


Figure 14 ACdEV analysis using the nCS1. 3μl of ACdEVs (Jurkat cells) in phenol-red free RPMI-1645
were analysed using a TS-900 cartridge on the nCS1. Approximately 2000 events/particles were
analysed and demonstrate a size distribution with a higher concentration of EVs at the lower detection
threshold.

411 ACdEV analysis using the TS-300 cartridge was relatively unsuccessful with only a few (<100) 412 discernible particles (data not shown). This did not acquire enough data for confident reporting of the 413 size distribution. The lack of apparent particulate is surmised to be due to one of two reasons; the lack 414 of surfactant in the sample and therefore a lack of wetting agent for the cartridge prefilter, or, the 415 sheer concentration of particles/EVs larger than the aperture being retained in the prefilter and 416 saturating it, and therefore severely reducing the flow of smaller particles. These hypotheses form 417 part of ongoing investigations to identify an alternative to detergent as a wetting agent and to 418 determine if pre-filtering for broad distributions is required. Nevertheless, MRPS shows great promise 419 for the analysis of lipidic nanoparticles.

420 Conclusions

There are a wide variety of techniques for studying the size and concentration of lipidic NPs. Each approach has advantages and disadvantages, so careful thought is required to determine the correct technique for a particular sample. This is particularly true when a sample is highly polydisperse and may even contain particles outside of the nano range. This article illustrates the benefits and drawbacks of the most commonly-used approaches for analysing NPs and highlights the promising new technique of MRPS. It is nearly always advantageous to apply multiple approaches to a particular sample to enable the most complete analysis to be performed.

428 Material and methods

429 Cell lines and culture

430 Jurkat T cell line, derived from an acute T cell leukaemia patient, were cultured at a density of 5E+05 431 cells.mL⁻¹ in RPMI 1640 medium (Sigma, Aldrich, UK) supplemented with 10% (v/v) FBS, 100 μ g.mL⁻¹ 432 penicillin/streptomycin, and 2mM glutamine. Cells were passaged every two to three days and 433 maintained at 37°C, 5% v/v CO₂.

434 EVs production

- 435 Jurkat cells were washed with PBS and resuspended in serum-free and phenol red-free RPMI-1640
- 436 (Sigma, Aldrich, UK) at a density of 4E+06 cells.mL⁻¹. Cells were then irradiated with a UV-B dose of 30

mJ.(cm²)⁻¹ using the UVP chromato-vue C71 cabinet. Analysis of apoptosis was performed using
annexin V-FITC / propidium iodide (PI) staining kit (BioLegend, San Diego) to confirm the cell death by
flow cytometry (CytoFLEX S, Beckman Coulter, USA).

440 Measurement on nCS1

• Nanosphere analysis

442 National Institute of Standards and Technology (NIST) traceable Nanosphere[™] size standards 443 (ThermoScientific) were suspended in Isoton-II solution with 1% Tween-20, filtered to 0.22 µm using 444 a PES syringe filter, to create a polydisperse mixture (60, 100, 202, 303, and 1000nm) with a final 445 concentration of ~2.52E+09 particles.mL⁻¹. 3µL of sample was loaded into two different cartridges, TS-446 400 and TS-2K, in order to visualize the different sized nanospheres.

447 • ACdEV analysis

3μL of ACdEVs suspended in phenol-red free RPMI (Sigma, Aldrich, UK) was loaded into TS-300 and
TS-900 cartridges and loaded into the machine for analysis.

The cartridge mould I.D. and date has to be inserted into the software, this provides the auto analysis engine with the parameters of the cartridge loaded, the machine then automatically primes the cartridge and collects the data. The machine will collect and analyse 10-second analysis windows until you either manually stop the runs or enter stop acquisition parameters, such as number of particles, in the auto-analysis run window.

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