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Enumeration of extracellular vesicles by a new improved flow cytometric method is comparable to fluorescence mode nanoparticle tracking analysis

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12 Abstract

13 Extracellular vesicles (EVs) play a role in a variety of physiological and pathological processes. However, use of EVs as biomarkers has been hampered by limitations of current detection and enumeration methods. We compared fluorescence-threshold flow cytometry (FT-FC) 14 to nanoparticle tracking analysis (NTA) for enumeration of cell culture-derived EVs. FT-FC and NTA utilising fluorescence mode (F-NTA) 15enumerated similar numbers of EVs stained with a membrane dye PKH67. Both methods were sufficiently sensitive to detect cell-derived 16 EVs above the background of culture medium. NTA in light scatter mode (LS-NTA) detected $10-100 \times$ more particles than either 17 fluorescence-based method but demonstrated poor specificity. F-NTA appeared to have better sensitivity for <100 nm vesicles, however, the 18 FT-FC method combined direct enumeration of EVs with high sensitivity and specificity in the >100 nm range. Due to wider availability and 19higher degree of automation and standardisation, FT-FC is a reasonable surrogate to emerging F-NTA for quantification of EVs. 20

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22 Key words: Extracellular vesicles; Flow cytometry; Nanoparticle tracking analysis; PKH67; Enumeration

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Introduction

The study of extracellular vesicles (EVs) is a rapidly growing 25 field in biology and medicine. EVs are small (<1 μ m in diameter) 26 bilipid membrane bound vesicles including exosomes, ecto- 27 somes, microvesicles, microparticles and apoptotic bodies among 28 other vesicle fractions, which can be released into blood and other 29 body fluids potentially by all cell types during cell activation, 30 intercellular interaction and cell death.¹ Attempts to classify these 31 heterogeneous vesicles by size or biogenesis have not been 32 universally accepted and many questions concerning biogenesis, 33 transport and other processes remain unanswered. Similarly, 34 there is no consensus on markers that identify the origin of 35 vesicles once outside their cell of origin. For example, exosomes 36 have traditionally been defined as originating from multivesicular 37 bodies and being smaller than 100-nm in size, whilst micro- 38 vesicles are considered to be larger than 100-nm and formed by 39 direct budding from plasma membranes. However, accumulating 40

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L. Pasalic et al / Nanomedicine: Nanotechnology, Biology, and Medicine xx (2016) xxx-xxx

experimental evidence suggests that particles less than 100-nm in 41 diameter can bud from plasma membranes, and that EVs with 42 exosome markers can be larger than 100-nm.² Despite these 43uncertainties, scientific interest in EVs has greatly expanded 44 45recently as EVs are recognised as central players in intercellular 46 communication. Furthermore, EVs are increasingly reported to have roles in many physiological and pathological processes 47 including intercellular communication, coagulation, angiogene-48 sis, cell survival, inflammation and immune response 49 modulation.^{1,3-5} Thus, EVs have significant potential as 50biomarkers for disease diagnosis, prognosis and monitoring. 51One particular area of interest is the potential use of EVs as 52diagnostic and prognostic biomarkers in cancer and cancer-53associated thrombosis.⁶⁻¹¹ As of February 2015, there were 53 54clinical trials registered with ClinicalTrials.gov (search terms: 55 biomarker AND (microparticles OR exosomes OR microvesi-56cles)) exploring use of EVs as biomarkers or therapeutic agents.¹² 57

A major obstacle to translation of EVs as biomarkers for 58 clinical use has been difficulty establishing standardised methods 5960 to measure EVs, primarily due to their small size, most being smaller than 300-nm in diameter.¹³⁻¹⁵ This is well documented¹⁵ 61 and includes problems with sensitivity in measuring sub-micron 62 particles as well as specificity when size is used as main 63 64 identification criteria. Several optical and non-optical techniques have been modified or developed for quantitative and qualitative 65 assessment of EVs including flow cytometry, dynamic light 66 scattering (DLS), nanoparticle tracking analysis (NTA), electron 67 microscopy (EM) and atomic force microscopy (AFM).¹⁶ 68

Flow cytometry remains the most commonly available and 69 widely used approach for enumeration and qualitative analysis of 70single vesicles. Benefits include availability, high throughput 71 and opportunities to explore multiple parameters using combi-72nations of dyes and fluorophores. Traditionally, forward scatter 73 signal in response to a 488 nm laser is used to define EVs by 74 size, with direct limitations including inability to resolve structures below 200 nm in diameter,¹⁷ false positive signal from detection of protein aggregates,¹⁸ and "swarm" detection, 7576 77 referring to registration of multiple vesicles simultaneously 78 identified as single event signals.^{19,20} 79

Recently, a protocol,²¹ based on original work from Nolte-'t Hoen and colleagues,²² was published for fluorescence-triggered 80 81 flow cytometry characterisation of fluorescent EVs using 82 83 membrane dye PKH67, sucrose density gradient fractionation, and a customised high-end cytometer (BD-Influx; BD Biosci-84 ences). This method overcame several problems associated with 85 conventional light scatter-based flow cytometry. Using mem-86 brane dye coupled with sucrose gradient fractionation, interfer-87 ence by protein aggregates or calcium-monophosphate 88 precipitates was eliminated and specificity was significantly 89 enhanced. Increased sensitivity for detection of small particles 90 was achieved by triggering event detection on fluorescence 91 signal rather than light-scatter, enabling easy detection of 92fluorescent beads down to 100-nm in diameter. 93

However, many of these gains were dependent on substantial customisation and around 3 h of flow cytometer preparation time. Customisation steps included configuration of pulseprocessing channels, changes to the FSC obscuration bar, replacement of standard tubing lines, removal of the polarisation unit and neutral density filter in front of the FSC-PMT and 99 installation of 488/10 nm band-pass filter in front of the 100 FSC-PMT. Preparation time is not insignificant and involves 101 laser calibration, with regular monitoring of sheath and sample 102 pressures using an external pressure meter to ensure stable flow 103 rate and reproducible enumeration. 104

These modifications made this approach impractical in our 105 busy multi-user facility. Therefore, we sought to adapt this 106 method to a commercially available flow cytometer, without 107 need for end-user modifications to create a flow cytometry 108 method for a multi-user environment whilst retaining sensitivity 109 and specificity of the original method. Importantly, we also 110 aimed to develop a simple automated and reproducible protocol 111 for enumeration of fluorescently labelled EVs isolated from 112 cell-conditioned supernatants. In this study, we chose a system 113 with low level EV production to ensure that the method was 114 robust at lower limits of detection. We compared results with 115 those obtained using NTA in light scatter and fluorescent modes. 116

Methods

Materials

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Extracellular vesicle isolation and labelling

For EV production, HEY human ovarian cancer cells were 120 plated in RPMI medium containing 5% of vesicle depleted FBS. 121 Culture medium not previously exposed to cells, but otherwise 122 treated identically to cell-conditioned medium, was used as 123 negative control. Cell-conditioned and control media (22 mL) 124 were collected after 72 h at ~80-90% confluency. 125

EVs were isolated from cell culture supernatant and control 126 medium by differential centrifugation at 4 °C, (300 g for 5 min, 127 3000 g for 10 min and 10,000 g for 20 min). EVs were pelleted 128 from the 10,000 g supernatant by ultracentrifugation at 129 100,000 g for 120 min in a 45Ti rotor (Beckman Coulter Inc, 130 Fullerton, USA). Pellets were washed in 1 mL of 0.1 μ m filtered 131 PBS and supernatant removed by centrifugation (TLA-110 rotor; 132 Beckman; 110,000 g, 60 min). Finally, washed pellets were 133 resuspended in 20 μ L of filtered PBS supplemented with 0.2% 134 vesicle depleted bovine serum albumin (BSA).

Freshly isolated EVs were fluorescently labelled and 136 fractionated as described by Nolte-'t Hoen and colleagues.^{21,22} 137 Briefly, EVs stained with 7.5 μ M PKH67 (Sigma Aldrich) 138 membrane dye were mixed with 2.5 M sucrose and overlaid with 139 linear sucrose gradient (2.0-0.5 M sucrose in PBS) in a SW40 140 tube (Beckman). Following isopycnic sucrose density gradient 141 centrifugation (192,000 g, 16 h) twelve 1-mL fractions were 142 collected starting from the bottom of each gradient. Fraction 143 density was measured using a digital refractometer (HI96801, 144 Hanna Instruments, Australia). Multiple aliquots of each gradient 145 fraction were stored at -80 °C before analysis. 146

Fluorescence-triggered flow cytometry (FT-FC)

The BD LSRFortessa flow cytometer (BD Biosciences, 148 Australia), equipped with a 100 mW blue 488 nm laser and a 149 photo multiplier tube (PMT) for detection of forward scatter 150 (FSC) (FSC-PMT), was used to detect fluorescently labelled 151 EVs. A high throughput 96-well plate sampler with volumetric 152 Download English Version:

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