

Development of a microfluidic droplet platform with an antibody-free magnetic-bead-based strategy for high through-put and efficient EVs isolation

Marco Morani, Myriam Taverna, Zuzana Krupova, Lucile Alexandre, Pierre Defrenaix, Thanh Duc Mai

▶ To cite this version:

Marco Morani, Myriam Taverna, Zuzana Krupova, Lucile Alexandre, Pierre Defrenaix, et al.. Development of a microfluidic droplet platform with an antibody-free magnetic-bead-based strategy for high through-put and efficient EVs isolation. Talanta, 2022, 249, pp.123625. 10.1016/j.talanta.2022.123625. hal-03841027

HAL Id: hal-03841027 https://hal.science/hal-03841027

Submitted on 22 Jul2024

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.



Distributed under a Creative Commons Attribution - NonCommercial - NoDerivatives 4.0 International License

Version of Record: https://www.sciencedirect.com/science/article/pii/S0039914022004210 Manuscript_a6b50b13df23a10eea3c62677947f49e

- 1 -

1	Development of a microfluidic droplet platform with an antibody-free magnetic-bead-
2	based strategy for high through-put and efficient EVs isolation
3	
4	Marco Morani ¹ , Myriam Taverna ^{1,2} , Zuzana Krupova ³ , Lucile Alexandre ¹ , Pierre
5	Defrenaix ³ , and Thanh Duc Mai ^{1*}
6	
7	¹ Université Paris-Saclay, CNRS, Institut Galien Paris-Saclay, 92296, Châtenay-Malabry,
8	France.
9	² Institut Universitaire de France (IUF)
10	³ Excilone - 6, Rue Blaise Pascal - Parc Euclide - 78990 Elancourt - France
11	
12	Correspondence: E-mail: thanh-duc.mai@u-psud.fr
13	
14	
15	Keywords: extracellular vesicles; isolation; magnetic beads; polymer precipitation;
16	microfluidic droplets
17	
18	
19	
20	
21	
22	
23	
24	
25	

26 Abstract

27 In this study, we present a novel microfluidic droplet-based strategy for high performance 28 isolation of extracellular vesicles (EVs). For EVs capture and release, a magnetic bead-based 29 approach without having recourse to any antibody was optimized in batch and then adapted to 30 the microfluidic droplet system. This antibody-free capture approach relies on the presence of 31 a water-excluding polymer, polyethylene glycol (PEG), to precipitate EVs on the surface of 32 negatively charged magnetic beads. We significantly improved the reproducibility of EV 33 recovery and avoided positive false bias by including a washing step and optimizing the 34 protocol. Well-characterized EV standards derived from pre-purified bovine milk were used 35 for EVs isolation performance evaluation. An EVs recovery of up to 25 % estimated with 36 nanoparticle tracking analysis (NTA) was achieved for this batchwise PEG-based approach. 37 The confirmation of isolated EVs identity was also made with our recently developed method 38 using capillary electrophoresis (CE) coupled with laser-induced fluorescent (LIF) detection. 39 In parallel, a purpose-made droplet platform working with magnetic tweezers was developed 40 for translation of this PEG-based method into a droplet microfluidic protocol to further 41 improve the performance in terms of EVs capture efficiency and high throughput. The 42 droplet-based protocol offers a significant improvement of recovery rate (up to 50 %) while 43 reducing sample and reagent volumes (by more than 10 folds) and operation time (by 3 folds) 44 compared to the batch-wise mode.

- 45
- 46
- 47
- 48

49

51 **1. Introduction**

52 Extracellular vesicles (EVs) are phospholipid bilayer-delimited particles produced by most 53 cell types and present in many body fluids [1, 2]. EVs contain and carry diverse biomolecules 54 that are specific to the mother cells from which they are secreted, allowing them to transmit a 55 variety of essential signals under both normal and pathological conditions. Hence, the 56 potential of EVs as prognostic or diagnostic biomarkers has attracted significant attention in 57 recent years [3, 4]. Furthermore, due to their high specific targeting ability, EVs have gained 58 much interest as engineered drug delivery systems for clinical and pharmaceutical 59 applications [5, 6]. However, there are still technological hurdles to purify, analyze and 60 characterize such nanometric bio-entities. Many methods for isolating EVs have been developed so far, including ultracentrifugation (UC), gradient ultracentrifugation, 61 62 ultrafiltration (UF), polymer co-precipitation, size-exclusion liquid chromatography (SEC), 63 immuno-extraction [7]. Among these, ultracentrifugation is widely considered as the gold 64 standard in all EV applications. However, this technique presents many drawbacks, such as 65 time-consuming procedures, contamination of EV populations by protein aggregates and other 66 particles, damage to the EVs membrane structure and possible considerable loss of EVs (EVs 67 yield may drop to 2 %) [8, 9]. Thus, there is still an urgent need for emerging EVs isolation 68 approaches that can provide EV purity and integrity in a reproducible and high-throughput 69 manner. Many modern isolation methods have been developed in this direction, such as flow 70 field-flow fractionation, ion-exchange, electrokinetic approaches as well as the combination 71 of multiple techniques, like UC with UF or SEC [10-12]. In parallel, microfluidic 72 technologies have made significant progresses for such purpose, exploiting both physical and 73 biochemical properties of EVs at micro/nanoscale level for their capture and/or detection [13-74 17]. The majority of microfluidic approaches rely on immunoaffinity to selectively capture 75 Immunoaffinity bead-based kits allow highly selective isolation of EVs through EVs.

76 antibodies specific for target EVs surface proteins [18-20]. However, following isolation, 77 those commercial kits, which are rather used for subsequent EVs downstream lysis and analysis, do not provide any efficient elution possibility to recover intact EVs. Moreover, the 78 79 main disadvantage of those strategies is the absence of universal EVs markers to ensure total 80 capture of all EVs. Few recent works on capture and eventual elution of EVs on magnetic 81 beads have been reported, using either electrostatic interaction [21], polymer mediated 82 adsorption of EVs on magnetic beads [22], a DNA aptamer-based system [23] or DNA linker 83 spacers [24]. No EVs recovery efficiency was reported in these works that used cell culture 84 media and/or plasma samples as starting materials from which accurate EVs quantification is 85 not trivial. At the actual stage, these works had to be realized batchwise with multiple in-tube 86 steps without automation.

87

88 The goal of this study was to investigate bead-based strategies for isolating and recovering 89 intact EVs without the use of immunoaffinity recognition and to adapt them to droplet 90 microfluidics. For such purposes, well-characterized high-quality EVs isolated from bovine 91 milk were used as EVs standards rather than non-quantified EVs from cell culture and plasma 92 samples. Then, the batchwise approach with the superior performance was transferred into an 93 automated and high-throughput protocol relying on a microfluidic droplet train. Different 94 operations in microfluidic droplets were developed and optimized to overcome the challenges 95 of beads clustering and poor recirculation in droplets in the presence of viscous polymers, 96 allowing to realise EVs capture on beads, washing and elution with a droplet sequence. So far, 97 droplet microfluidics has been communicated only twice for immunoassay-based detection of 98 EVs [25, 26] and has never been exploited for high performance and high throughput isolation 99 of EVs.

101 **2. Materials and methods**

102 2.1. Chemicals and reagents

103 2-(cyclohexylamino)ethanesulfonic acid (CHES), phosphate buffered saline (PBS 10x), 104 sodium dodecyl sulfate (SDS, 98.5 % (GC)), tris(hydroxymethyl)aminomethane (Tris), 105 polyethylene glycol (PEG 8000), albumin from human serum, IgG from human serum and 106 human transferrin were all obtained from Sigma Aldrich (St. Louis, MO, USA). Sodium 107 hydroxide (1 M) and hydrochloric acid (1 M) were obtained from VWR (Fontenay-sous-Bois, 108 France). All solutions were prepared with deionized water purified with a Direct - Q3 UV 109 purification system (Millipore, Milford, MA, USA). Vybrant[™] CFDA SE Cell Tracer Kit 110 (dye 5-(and-6)-carboxyfluorescein diacetate succinimidyl ester, CFDA-SE) was purchased 111 from Thermo Fisher Scientific (Waltham, MA, USA). Fluorinert oil FC-40 (ZF-0002-1308-0) 112 was purchased from 3 M (USA). The surfactant 1H, 1H, 2H, 2H – perfluoro-1-decanol was 113 obtained from Sigma Aldrich. ExoCAS-2 kit containing poly-L-lysine-coated magnetic beads, 114 washing and elution buffer solutions was purchased from Microgentas (Seongbuk-gu, 115 Republic of Korea). Carboxylate functionalized magnetic beads (Dynabeads MyOne, 10 116 mg/mL, diameter of 1 µm) and silica-based magnetic beads (Dynabeads MyOne Silane, 40 117 mg/mL, diameter of 1 µm) were obtained from Thermo Fisher Scientific. EVs samples 118 isolated from bovine milk were provided by Excilone (Elancourt, France).

119

120 **2.2.** Apparatus and material

For macroscale protocols, all magnetic-bead-based assays in batch were carried out in protein LoBind 1.5 mL tubes purchased from Eppendorf (Hamburg, Germany). For retaining magnetic beads, a neodymium magnet purchased from Ademtech (Adem Mag MSV, Bessac, France) was used. Shaking of magnetic bead suspensions during the incubation and washing steps was realized with a mixer (Eppendorf ThermoMixer C).

126 For droplet microfluidic operations, a microfluidic droplet platform was constructed and was 127 inspired from our previous configuration [27]. It comprises a syringe pump (from Nemesys, 128 Cetoni GmbH) equipped with 1 mL glass syringes (purchased from SGE) and a motorized 129 pipettor arm (Rotaxys, Cetoni GmbH), used for droplets generation. A 96-well plate (Thermo 130 Scientific) was used for sample storage and reagent solutions. The plate was mounted on a 131 custom-made holder that can be moved in the X, Y, and Z directions. PTFE tubing with ID of 132 0.3 mm and OD of 0.6 mm (Z609692-1PAK, Sigma Aldrich) was used to conduct the droplet 133 trains. Fluorinated oil FC-40 mixed with the surfactant (1H, 1H, 2H, 2H - perfluoro-1-134 decanol, 2 % w/w) was used to separate aqueous droplets inside PTFE tubing. In the protocol, 135 a magnetic tweezer, prepared in-house was used. It is composed of a paramagnetic tip 136 activated by a magnetic coil. Macro Objective (MLH-10X) mounted on a low-cost CMOS 137 Cameras (acA1300-60 gm, Basler) and white LED back light illumination (Schott Lighting 138 and Imaging) were employed for droplet observation.

139

The analyses using CE coupled with laser induced fluorescent detection (CE-LIF) were performed with a PA 800 Plus system (Sciex Separation, Brea, CA) equipped with a solidstate LIF detector (excitation wavelength of 488 nm, emission wavelength of 520 nm) purchased from Integrated Optics (Art. No. 40A-48A-52A-64A-14-DM-PT, distributed by Acal BFi, Evry, France). Uncoated fused silica capillaries were purchased from CM Scientific (Silsden,UK). Data acquisition and instrument control were carried out using Karat 8.0 software (Sciex Separation, Brea, CA).

147

148 2.3. Methods

149 Isolation of bovine milk-derived EVs with sucrose gradient ultracentrifugation

150 Skimmed bovine milk samples were obtained by centrifugation of 50 mL whole milk at 3000 151 g for 30 min at 4 °C (Allegra X-15R, Beckman Coulter, France). The whey was obtained after 152 acid precipitation with 10 % (v/v) acetic acid, incubation at 37 °C for 10 min and 10 % (v/v) 1M sodium acetate for 10 min at RT followed by centrifugation at 1500 g, 4 °C for 15 min 153 154 and filtration using vacuum-driven filtration system Millipore Steritop, 0.22 µm. The whey 155 supernatants were concentrated using Amicon 100 kDa centrifugal filter units (Merck 156 Millipore) at 4000 g and 20 °C up to final volume of 6 mL. The obtained retentate was ultra-157 centrifuged for pelleting the EVs at 100000 g for 70 min at 4°C (Beckman Coulter, Optima 158 XPN-80, 50TI rotor). The pellets were solubilized in 500 µL of PBS then added to 11 mL of 159 pre-prepared sucrose gradient 5-40 % and ultra-centrifuged at 200000 g for 18 h at 4 °C 160 (Beckman Coulter, Optima XPN-80, SW41 rotor). Fractions of 1 mL were collected and the 161 selected ones containing targeted exosome population were diluted in 6 mL of PBS 1X and 162 finally centrifuged at 100000 g for 70 min at 4 °C (Beckman Coulter, Optima XPN-80, 50TI 163 rotor). The pellets were resuspended in 50 µL of PBS 1X and stored at -80 °C, until further 164 analyses.

165

166 *Isolation of pony plasma and serum derived EVs with size exclusion chromatography (SEC)* 167 Preparation of plasma: Peripheral blood was collected into EDTA-coated vacutainer tubes. 168 After ten-time inversion, samples were processed within the 60 min of collection. Consecutive 169 centrifugation steps at 2500 g, 4 °C for 15 min and then at 15000 g for 10 min were performed 170 followed by filtration of the supernatant through 0.22 µm filters. Preparation of serum: Whole 171 blood was collected into anticoagulant-free tubes and allowed to clot at room temperature for 172 45 min. The clot was removed by centrifugation at 3200 g, 4 °C for 15 min, followed by 173 centrifugation at 15000 g, 4 °C for 10 min and filtration of the supernatant through 0.22 µm 174 filters. 500 µL of pre-treated plasma/serum was loaded onto a qEVoriginal SEC column (Izon Science, New Zealand) previously washed and equilibrated with PBS. Fraction collection (0.5 mL per fraction) was carried out immediately using PBS 1X as elution buffer. The selected elution fractions were pooled and subsequently concentrated using 100 kDa Amicon centrifugal filter units (Merck Millipore). Post-treatment processing with several washing steps with PBS was applied to obtain pure EV fractions that are highly enriched with exosomes.

181

182 EV isolation using commercial kits

183 ExoCAS-2 magnetic bead-based ion exchange kit was used for the study of EV yield 184 performance. The experimental procedure is described in a paper recently published [21]. 185 Briefly, PLL-coated beads were mixed with our standard EVs (bovine milk derived EVs), 186 followed by incubation of the mixture for 30 min at 4 °C in a rocking mixer. After incubation, 187 the EVs-bound beads were carefully washed with 2 mL of ExoCAS-2 washing solution and 188 then re-suspended in ExoCAS-2 elution solution by shaking for 5 min at 1000 rpm. Finally, a 189 magnet was employed to retain magnetic beads and the supernatant containing yielded EVs 190 was collected.

191

192 PEG-based EVs precipitation on magnetic beads in batch mode

For PEG-based EVs precipitation on magnetic beads in batch, a volume of 200 μ L of PEG (25 % m/v), 150 μ L of carboxylate functionalized magnetic beads (10 mg/mL) and 250 μ L of deionized water was incubated with 400 μ L of standard EVs or biological fluid (simulated human serum and pony plasma and serum, 5 times diluted with deionized water) on a mixer at 25 ° C for 1 h at 700 rpm. Beads were then carefully washed twice with PEG 5 % / NaCl 0.2 M. After removal of washing solution, the magnetic beads in tubes were re-dispersed in 400

µL of PBS 1X solution and then incubated on shaking for 5 min at 25 °C for EV elution. A
magnet was used to remove the magnetic beads, and EVs were recovered in the supernatant.

201

202 PEG-based EVs precipitation on magnetic beads using the microfluidic droplet platform

203 A robotic arm and a 500 µL syringe were utilized to pipette the droplets from separate 204 reservoirs into a PTFE tube in a fully automated manner. To complete one PEG-based EVs 205 precipitation protocol, a train of 5 droplets (each containing 6 µL) confined and separated by 206 oil was required. It includes: i) one droplet of carboxylate functionalized magnetic beads (1.5 207 mg/mL), ii) one droplet of EV sample in PEG 5 %, iii-iv) two droplets of washing solution 208 (PEG 5 % / NaCl 0.2 M) and v) one droplet of elution solution (PBS 1X). Beads were trapped 209 out of one droplet and released into another by electronic triggering of the magnetic tweezer. 210 The incubation was carried out with regular droplet back and forth movements at RT for 25 211 min. After incubation, two washing droplets are flushed over EVs-bound beads. The EVs-212 bound beads were then dispersed into the elution droplet with regular droplet back and forth 213 movements at RT for 5 min. The elution droplet containing released EVs was then collected 214 in a tube for downstream analysis.

215

216 Nanoparticle Tracking Analysis (NTA) of EVs

Particle concentration and size distribution were determined with a Nanosight NS300 instrument (Malvern, version NTA 3.2 Dev Build 3.2.16) equipped with a 405 nm laser, sCMOS camera type and the NTA software v3.1. The video acquisition was performed using a camera level of 14. 3 videos of 90 s with a frame rate of 30 frames/s were captured for each sample at 25 °C and subsequently analyzed with a threshold set up at 5. The results were validated with at least 2000 valid tracks for each triplicate. All experiments were carried out with samples pre-diluted in PBS according to input sample concentrations. The working particle concentrations were kept within the range of $10^6 - 10^9$ particles per mL for optimal analysis.

226

The Zetaview system (Particle Metrix, Germany) was equipped with a 488 nm laser. EVs samples were diluted to allow the measurements at 50–200 particles/frame. Each experiment was performed in duplicate on 11 different positions within the sample cell. The specifications used for all measurements were cell temperature of 25 °C, sensitivity of 70, shutter of 100, Max Area of 1000, Min Area of 10, Min Brightness of 25. The results were validated with at least 1000 valid tracks for each run. For data capture and analysis, the Nanoparticle Tracking Analysis Software (ZNTA) vs 8.05.04 was used.

234

235 CE-LIF of fluorescently labelled EVs

236 Details on the CE-LIF method for EVs analysis can be seen in our recent work [28]. Briefly, 237 fluorescently labelled EVs were prepared using the 5-(and-6)-carboxyfluorescein diacetate 238 succinimidyl ester (CFDA-SE). After removal of residual CFDA-SE via filtration with 239 commercial Exosome Spin Columns (MW 3000, Thermo Fisher Scientific Waltham, MA 240 USA), labelled EVs were analyzed with CE-LIF using a fused silica capillary having I.D. of 241 50 μ m, effective length (L_{eff}) of 50.2 cm and total length (L_{tot}) of 60.2 cm. The capillary was 242 pre-conditioned with water for 10 min, 1 M NaOH for 10 min, 1 M HCl for 10 min and then 243 water for 10 min. The rinsing between two analyses was carried out with 50 mM SDS for 5 244 min, 1 M NaOH for 5 min, deionized water for 5 min, and finally the running BGE composed 245 of Tris / CHES (IS 90 mM, pH 8.4) for 5 min using a pressure of 30 psi. EVs samples were 246 injected hydrodynamically from the inlet end by applying a pressure of 0.5 psi for 2 min. The 247 separation was carried out under 25 kV (normal polarity) at 25 °C and the samples were kept 248 at 5 °C with the sample storage module of the PA 800 Plus equipment.

249

3. Results and discussion

3.1. Batchwise EVs isolation development

252 With the aim to establish a high-performance and high-throughput microfluidic droplet 253 system for EVs isolation, we evaluated two recently communicated batchwise EV-enrichment 254 strategies that are alternative approaches to the immunoaffinity-based ones. They hold high 255 potential for subsequent translation into a microfluidic format in terms of minimal forefront 256 preparations, non-laborious operations, as well as ease of manipulation and step transition 257 thanks to the use of magnetic beads as cargos. The first one is a magnetic bead-mediated selective adsorption strategy (MagExo), exploiting the presence of a water-excluding 258 259 polymer, polyethylene glycol (PEG), to lock up a significant number of water molecules, 260 forcing thereby the EVs to aggregate and precipitate on the surface of magnetic beads [22]. 261 The second strategy called ExoCAS-2 relies on an ionic exchange mechanism, using magnetic 262 beads coated with a polycationic polymer, poly-L-lysine (PLL), to quickly trap negatively 263 charged EVs via electrostatic interaction [21]. For performance comparison, EVs standards 264 derived from bovine milk having narrow size distribution, with well-defined concentrations 265 and exhaustive characterizations by NTA, DLS, LC-MS/MS and TEM, were used as the 266 starting sample. TEM images revealed the absence of contaminating protein residues in these 267 EVs standards (Fig. S1A in the supporting information ESI). Major protein contaminants (e.g. 268 α -s1 casein, β -casein, α -Lactalbumin and serum albumin) were not found according to the 269 LC-MS/MS analysis (Fig. S1B). The use of EV standards of high purity allowed to evaluate 270 the EVs recovery more accurately. In our case, quantification of EVs isolation yields obtained 271 with MagExo and ExoCAS-2 methods was possible by comparing the EVs concentrations 272 before and after isolation processes (see Fig. 1). Unsatisfactory EVs recovery (less than 10 %) was obtained with commercial PLL-coated beads, compared to that achieved when using PEG 273

274 and silica magnetic beads (34 %). NTA data also showed a size shift towards smaller particles 275 when EVs were eluted from PLL beads. This could be a consequence of PLL release from the 276 beads during the elution process. This was confirmed by the CE-LIF analysis (discussion 277 below). This led to a critical PLL concentration in the eluent that is high enough to cause EVs 278 lysis. Indeed, several studies have shown that PLL can penetrate through vesicles via 279 interaction with the lipid membrane [29-31], provoking vesicle lysis from a certain threshold 280 concentration of PLL [32]. In parallel to NTA measurements for EVs recovery evaluation, we 281 used our recently developed CE-LIF approach [28] to validate the identity of EVs collected 282 with MagExo and ExoCAS-2 methods (see Fig. 2). The peak profile of EVs collected with PEG-based protocol corresponds well to the fingerprint of bovine milk-derived EVs, 283 284 confirming the presence of intact EVs in the eluent (Fig. 2A). On the contrary, with the ion 285 exchange method, multiple tagged species were detected. To understand the origin of these 286 peaks, solutions containing different PLL concentrations without the presence of EVs were 287 analyzed. As can be seen in Fig. 2B, many peaks were still detected, whose intensities were 288 related to the PLL concentrations. These results confirmed the hypotheses that the peaks 289 observed come from PLL leakage from the magnetic beads, which could not be visualized 290 with conventional NTA. The released PLL which can be labelled by the residual CFDA-SE 291 dye through its amino groups lead to unwanted products in the eluent. The ExoCAS-2 method 292 was therefore not further considered; and the PEG-based method (MagExo) was chosen for 293 further optimisation.

294

Besides the preservation of intact EVs after purification, protocol reproducibility is another important point to consider to guarantee consistency of isolated EVs population and quality. Indeed, by reproducing the MagExo procedure described in reference [22], we observed significant batch-to-batch variation in vesicle concentration (RSD of 30 %). The lack of EVs 299 washing between the capture and elution steps in the original protocol led to misleading data. 300 To minimize this cross contamination and to recover intact EVs for further characterization, 301 we developed a washing protocol after the EVs capture step. By adding two consecutive washing steps with PEG 5 % w/v to remove residual unbound EVs while maintaining 302 303 captured EVs on magnetic beads, we significantly improved the repeatability (RSD) to 8 %. 304 This came with some penalty, as the EV isolation yield fell to 17 %. To further improve the 305 performance of EVs capture and thus isolation yield, we carried out different optimizations on 306 magnetic bead concentrations (0.5 - 2 mg/mL), bead chemical surface (with carboxylic or silane groups), incubation temperature for EVs capture (4 - 25 °C) as well as PEG 307 308 concentrations (5-15 % w/v). Note again that the performance of EVs capture relies on the 309 presence of PEG to force the EVs to precipitate (by locking up a significant number of water 310 molecules) on the surface of magnetic beads. As can been seen in Fig. S2 in the ESI, much 311 better EVs capture performance was achieved with beads with carboxylic groups (EVs yield 312 of 61 %) than with silane groups (less than 5 %). The incubation temperature was found not to 313 significantly influence the EVs capture performance, with no remarquable difference 314 observed at 4 °C (yield of 17.2 %) and 25 °C (yield of 20 %). The concentrations of magnetic beads and PEG played important roles in the on-bead retaining of EVs. A compromise of their 315 316 concentrations had to be made to allow facile PEG-induced precipitation of EVs on sufficient 317 quantity of carboxylate beads, while avoiding i) poor recirculation or even clustering of beads 318 at too high concentrations and ii) hinderance of the arrival of EVs on to the beads surface due 319 to too elevated medium's viscosity at too high PEG concentrations. Overall, the highest EVs 320 recovery rate (after 2 washing steps) of 25 % \pm 8 % was achieved with the optimized 321 conditions using PEG concentration of 5 % and carboxylic magnetic beads' concentration of 322 1.5 mg/mL with 1 h incubation at 25 °C. A higher bead concentration of 2 mg/mL was tested 323 but no satisfactory results were obtained due to clustering and poor circulation of beads. PEG

324 concentrations lower than 5 % w/v were not considered in our work, based on previous optimization already reported [22]. Conveniently, the working temperature of 25 °C is well 325 326 adapted for subsequent translation of batchwise protocol into a microfluidic format where 327 cooling function is not readily available. To evaluate the reusability of magnetic beads 328 functionalized with carboxylic groups for repeated isolation of EVs, the same beads employed 329 for the first round of capture and elution of EVs were recovered in PBS and then subjected to 330 the second and third rounds. As shown in Fig. S3 in the ESI, the concentration range of 331 recovered EVs in the second round is similar to that obtained from the first round. The 332 variation of recovered EVs concentrations is however significantly higher (RSD of 24 % vs. 9 333 %, respectively). While the surface charge of the beads is expected not to be hammered by the 334 elution media (i.e., PBS), the possible presence of residual PEG on the surface of beads after 335 the first elution step could explain such an elevated variation in the second round. When using the same beads for the third cycle of EVs capture - elution, no satisfactory results were 336 337 obtained, with the recovered EVs concentration almost 5-fold lower than those obtained in 338 previous cycles (Fig. S3 in ESI). While the reusability of magnetic beads for EVs capture and 339 elution is possible for 2 cycles with careful consideration of PEG presence, fresh beads 340 without recycling are nevertheless required for further experiments in order to minimize the 341 risk of EVs isolation uncertainty.

342

343 **3.2. Droplet microfluidics for EVs isolation: proof of concept**

The optimised batchwise PEG-based method was subsequently converted into a microfluidic droplet protocol in order to provide a high level of automation and integration, significant reduction in sample/reagent amounts, and a higher performance in terms of isolation efficiency. The instrumental setup of the purpose-made microfluidic platform is shown in Fig. 3. It is composed of a syringe pump, a motorized pipettor arm for droplet production, a 96349 well plate for sample and reagent storage and a magnetic tweezer for manipulation of 350 magnetic beads. With this system, we used a train of 6 µL droplets containing in a defined 351 sequence (i) the magnetic bead suspension, (ii) the EVs sample, (iii) the washing and (iv) 352 elution solutions to replace different tube-based steps. When working with the droplet-wise 353 multi-step protocol, one hurdle encountered is the difficulty to efficiently transfer the target 354 species (EVs in this case) from one droplet to another without any risk of cross 355 contamination. This was expected to be overcome with our setup through the use of magnetic 356 beads as the controllable carrier of target analytes between droplets. A purpose-made 357 magnetic tweezer, composed of a paramagnetic tip activated by an electrical coil [27, 33], was 358 employed to manipulate magnetic beads between droplets via application of an external 359 electrical field. Table 1 provides an overview of the operation sequence. A train of droplets 360 containing different solutions and sample in a defined order was delivered through the 361 magnetic tweezer where the beads were extracted from the first droplet and transported into 362 the sample one containing EVs. The incubation was subsequently performed by pushing the 363 droplet train back and forth inside the tubing. The magnetic beads that retain EVs on their 364 surface were then trapped by the magnetic tweezer and the supernatant droplet was washed 365 away, followed by flushing of the trapped magnetic beads with two washing droplets. The 366 washed beads were finally released into an elution droplet that was finally collected in an oil-367 containing tube for further analyses.

368

369 Several problems were however encountered that led to failure of such droplet protocol in our 370 first experiments, notably beads clustering and poor recirculation in droplets in the presence 371 of PEG. Indeed, the beads often stayed at the rear of the moving droplet, forming undesired 372 bead clusters when the droplet moved inside the tubing over an extended distance. This was 373 ascribed to come from the presence of PEG which significantly increased droplet viscosity

[34], which in turn impaired the effective recirculation of the beads (see Fig. S4 in the ESI). 374 375 To enhance recirculation of the beads within a droplet, we put our efforts on optimizing the 376 droplet movement pattern and travel volume (i.e., the oil volume required to push and pull the 377 droplet) during the back-and-forth movement. Chaotic trajectories are known to enhance 378 mixing efficiency [35, 36]. Consequently, different droplet movement patterns were also 379 tested, including the straight, the U-shaped patterns (i.e., the tubing was shaped over a 380 metallic guideline) and the spiral-formed one (i.e., the tubing was coiled around a cylindrical 381 support) (see Fig. S5 in ESI). With the straight pattern, bead aggregation was always observed 382 in the presence of PEG, regardless of the travel volume (Fig. S4A). For other movement 383 patterns, bead clustering was alleviated but still visible when working with a large travel 384 volume of 10 µL (Fig. S4 B and C). This undesired phenomenon was finally avoided when 385 employing the spiral or U-shaped patterns with a short travel volume of $6 \mu L$ (equivalent to a 386 droplet volume). By keeping these optimized setups and conditions, we further investigated 387 the in-droplet incubation duration (from 5 to 45 min) during the EVs capture step (Fig. S6 in 388 the ESI). The EVs recovery increased from 10 % for 5 min incubation to 27 % for 25 min 389 incubation, and remained stable even when the incubation time was increased up to 45 min. 390 For the spiral-formed pattern, the best EVs recovery of 27 $\% \pm 4 \%$ was thus found for 25 min 391 incubation Under the same incubation time, the U-shaped pattern offered the EVs recovery rate of 39 % \pm 3 %, indicating better interaction between EVs and magnetic beads, and thus 392 393 higher EVs capture efficiency for the U-shaped pattern. Among the tested droplet movement 394 ones, the straight one gave the least EVs recovery $(14 \% \pm 4 \%)$ and therefore was not further 395 considered. To obtain more precise information on the quality of the isolated EVs, the 396 samples were analyzed with Particle Metrix' ZetaView [37]. As revealed by Zetaview data 397 (Fig. 4B), 80 % of the recovered EVs fell within the range of the initial size distribution 398 (diameter of 171 nm, accounting for 79 % of the whole population), while the remaining 20 %

were represented by larger aggregates (16.7 % for the diameter of 315 nm, and 4.3 % for 472
nm). These aggregates are presumably formed due to the PEG capacity to wrap and condense
together two or three EVs (corresponding to the sizes of 317 and 472 nm, respectively),
making them hard to resuspend in the absence of PEG during the elution step.

403

404 **3.3. Microfluidic droplet-based isolation of EVs from biofluids**

405 The developed microfluidic droplet instrument and protocol were then used to isolate EVs 406 from more complex matrices. First, we used PBS mixed with three matrix proteins, including 407 albumin, IgG and transferrin to imitate human extracellular fluids [38]. This simulated human serum was then spiked with standard EVs (i.e., bovine milk derived EVs at 1.02 x 10¹¹ 408 409 particles/mL) and passed through the microfluidic droplet system to evaluate the EVs isolation performance. As indicated in Fig. 5, the concentration of collected EVs was 3.69 x 410 10^{10} particles/mL, giving an EVs recovery rate of 36 % with spiked simulated human serum, 411 412 which was not far from that obtained with pure standard EVs (39 %). We then applied the 413 droplet protocol to pony plasma and serum, and compared the results obtained with the in-414 tube batchwise protocol (Fig. S7 in ESI). When dealing with such complex biofluid matrices, 415 the droplet approach gave an overall particle concentration much higher (up to eight folds) 416 than those obtained with the in-tube method. For instance, with the pony serum sample, the recovered EVs concentration was 8 x 10¹⁰ particles/mL when using the microfluidic droplet 417 platform, whereas a significantly lower one (1.05 x 10¹⁰ particles/mL) was found for the in-418 419 tube method. A similar observation was found for pony plasma, with 17.7 x 10^{10} particles/mL vs. 4.9 x 10¹⁰ particles/mL for microfluidic and in-tube setups, respectively. Moreover, the 420 421 size distributions of EVs purified with the microfluidic droplet setup are much narrower with 422 only one main peak for each tested sample (Fig. S7). This was, on the other hand, not the case for the in-tube protocol where many peaks were found in the range of 70 - 400 nm for both 423

424 pony serum and plasma samples. To confirm the identity of EVs isolated from plasma and 425 serum samples, the elution droplets were also analyzed with CE-LIF to reveal different EVs 426 subpopulations (Fig. 6). Based on the peak intensities in each electropherogram, the highest 427 concentration was found for the fraction with the shortest migration time (9-12 min) whereas 428 the signal of the second peak zone (12-15 min) was less intense and this is more remarked for 429 the serum samples. This kind of EVs fingerprints was found similar to those obtained in our 430 previous work with EVs from pony plasma and serum isolated by SEC [28]. Interestingly, the 431 third peak zone (15-20 min) appeared in the electropherograms obtained with both in-batch 432 mode and the droplet system, suggesting that a distinct subpopulation emerges when using the 433 PEG-based isolation methods. Indeed, different isolation methods may lead to the shift or 434 differences in size distributions of the collected EVs, as already evidenced for SEC and UC 435 isolation methods [39, 40]. In our case, the PEG-based method is expected to capture non-436 selectively all EVs subpopulations, leading to more EVs fractions being visualized with CE-437 LIF as seen in Fig. 6. NTA measurements for both pony serum and plasma revealed a shift 438 towards smaller size distributions when using the droplet approach (Fig. S7 in ESI). This may 439 correspond to an increased concentration of the small-sized subpopulations, which may 440 correlate to the more pronounced appearance of the third peak zone (15-20 min) in the CE-441 LIF electropherograms. Nevertheless, no further speculation was made to interpret the 442 presence of these three subpopulations observed by CE as no clear relationship between size 443 and charge properties of EVs and their electrophoretic mobilities can be stated at this stage.

444

445 **4. Conclusion remarks**

We successfully developed a new approach (instrumentation and methodology) for EVs isolation from both pre-purified standards and biofluid samples, for the first time in microfluidic droplet format. Using a train of micrometric droplets, containing magnetic beads 449 in the first droplet, the sample, washing and elution solutions in the following ones, we 450 allowed significant sample and reagent volume reduction (by 5 times), minimization of 451 manual operations, diminution of operation time (by twice) and improvement of EVs recovery 452 rate by almost 2 folds. Thanks to automatization and miniaturization that we achieved with 453 droplet microfluidics higher throughput can now be expected. Integration of this microfluidic 454 EVs isolation module as a forefront of downstream EVs analysis and characterization is now 455 envisioned. In the present proof-of-concept study that deals with both instrumental and 456 methodological developments at the same time, the univariate approach was chosen to better 457 understand the impact of the different experimental factors. Indeed, the study on EVs is still at 458 early and emerging stage, and all parameters that would have influence on the EVs stability 459 and behaviour (e.g., morphology modification, surface charge changes, risk of lysis etc.) have 460 not been all identified and mastered, which in turn do not allow simultaneous investigation of 461 multiple variables during the optimization of EVs capture and release. Further optimizations 462 with the multivariate approach could be envisaged in the next phase when all risks of EVs 463 modifications during experiments could be mastered and understood.

- 464
- 465
- 466
- 467
- 468

469 The authors have declared no conflict of interest.

470

471 Acknowledgement

472 This work has been financially supported by the Institut Universitaire de France (for M.

473 Taverna, senior member) and the Agence Nationale de la Recherche (ANR, France) with the

474 grant no. ANR-18-CE29-0005-01. The European fellowship for Lucile Alexandre was
475 supported by the project H2020-MSCA-IF-2019 (IMPED 896313). The doctoral scholarship
476 for Marco Morani was supported by the doctoral school 2MIB (Sciences Chimiques:
477 Molécules, Matériaux, Instrumentation et Biosystèmes) – University Paris Saclay.

480	Table 1: Operation s	sequence of the m	icrofluidic droplet	protocol using o	droplets of 6 u	L each
100	I able II operation :	equence of the m	ieronaiaie aropiet	protocor asing .		L vavn

		1		
Operation	Droplet composition	Flowrate	Incubation	Back-and-forth
		(µL/s)	time (min)	droplet travel
				distance (+/- μL)
Sample incubation	EVs sample + PEG 5 %	4	25	6
	(w/v) + magnetic beads (1.5			
	mg/mL			
Washing 1	PEG 5 % / NaCl 0.2 M	0.5	Flush	-
Washing 2	PEG 5 % / NaCl 0.2 M	0.5	Flush	-
Sample elution	PBS 1X	4	5	6





Fig. 1. NTA measurements of EVs isolated from bovine milk before purification (A) and after purification with (B) MagExo or (C) ExoCAS-2 method. NTA histograms represent the mean of three replicate measurements of the same sample, with the standard deviation (SD) in grey.

504

- 505
- 506
- 507
- 508



Fig. 2. (A) CE-LIF measurements of EVs before purification (a), and after purification with (b) MagExo or (c) ExoCAS-2 method. (B) CE-LIF profiles of solutions of PLL at different concentrations, corresponding to PLL-coated beads suspension volumes of (a) 1 mL; (b) 0.5 mL; (c) 0.1 mL. CE-LIF conditions: fused silica capillary having I.D. of 50 μ m, effective length (L_{eff}) of 50.2 cm and total length (L_{tot}) of 60.2 cm; BGE composed of Tris / CHES (IS 90 mM, pH 8.4); applied voltage: 25 kV (normal polarity).



Fig. 3. Schematic drawing of microfluidic droplet system. B: droplet containing magnetic beads; S: sample droplet; W: washing droplet; E: droplet containing the elution solution. Droplets are separated by oil.



538 Metrix) for EVs standard before (a) and after (b) purification with the microfluidic 539 droplet platform using the U-shaped pattern. NTA histograms represent the mean of



Fig. 5. NTA measurements of EVs-spiked artificial serum before (A) and after (B) EVs isolation using the microfluidic droplet platform. The sample was 5-time diluted with PBS prior to microfluidic droplet operations. NTA histograms represent the mean of three replicate measurements of the same sample, with the standard deviation (SD) in grey.





Fig. 6. CE-LIF for EVs from pony plasma (A) and serum (B), isolated with PEG-based EVs precipitation method in batch mode (a); with the microfluidic droplet system using the U-shaped pattern (b) and isolated with SEC (c). CE conditions as in Fig. 2.

572 **References:**

- 573 [1] G. van Niel, G. D'Angelo, G. Raposo, Shedding light on the cell biology of extracellular
 574 vesicles, Nat. Rev. Mol. Cell Biol. 19(4) (2018) 213-228.
- 575 [2] L.M. Doyle, M.Z. Wang, Overview of extracellular vesicles, their origin, composition,
- 576 purpose, and methods for exosome isolation and analysis, Cells 8(7) (2019) 24.
- 577 [3] J. Howitt, A.F. Hill, Exosomes in the pathology of neurodegenerative diseases, J. Biol.
- 578 Chem. 291(52) (2016) 26589-26597.
- 579 [4] W. Guo, Y.B. Gao, N. Li, F. Shao, C.N. Wang, P. Wang, Z.L. Yang, R.D. Li, J. He,
- 580 Exosomes: New players in cancer, Oncol. Rep. 38(2) (2017) 665-675.
- 581 [5] P. Vader, E.A. Mol, G. Pasterkamp, R.M. Schiffelers, Extracellular vesicles for drug
 582 delivery, Adv. Drug Deliv. Rev. 106 (2016) 148-156.
- [6] H.C. Bu, D.G. He, X.X. He, K.M. Wang, Exosomes: Isolation, analysis, and applications
 in cancer detection and therapy, Chembiochem. 20(4) (2019) 451-461.
- 585 [7] Z. Zhao, H. Wijerathne, A.K. Godwin, S.A. Soper, Isolation and analysis methods of
- 586 extracellular vesicles (EVs), Extracellular Vesicles and Circulating Nucleic Acids 2(1)
 587 (2021) 80-103.
- 588 [8] C. Gardiner, D. Di Vizio, S. Sahoo, C. Thery, K.W. Witwer, M. Wauben, A.F. Hill,
- 589 Techniques used for the isolation and characterization of extracellular vesicles: Results
 590 of a worldwide survey, J. Extracell. Vesicles 5 (2016).
- 591 [9] M.Y. Konoshenko, E.A. Lekchnov, A.V. Vlassov, P.P. Laktionov, Isolation of
- 592 extracellular vesicles: General methodologies and latest trends, Biomed. Res. Int.593 (2018) 27.
- 594 [10] H. Yan, Y. Li, S. Cheng, Y. Zeng, Advances in analytical technologies for extracellular
 595 vesicles, Anal. Chem. 93(11) (2021) 4739-4774.

- 596 [11] T. Liangsupree, E. Multia, M.-L. Riekkola, Modern isolation and separation techniques
 597 for extracellular vesicles, J. Chromatogr. A 1636 (2021) 461773.
- 598 [12] M. Morani, T.D. Mai, Z. Krupova, G. van Niel, P. Defrenaix, M. Taverna, Recent
- 599 electrokinetic strategies for isolation, enrichment and separation of extracellular
- 600 vesicles, TrAC Trends Anal. Chem. (2021) 116179.
- [13] S. Hassanpour Tamrin, A. Sanati Nezhad, A. Sen, Label-free isolation of exosomes using
 microfluidic technologies, ACS Nano (2021).
- 603 [14] Y. Meng, M. Asghari, M.K. Aslan, A. Yilmaz, B. Mateescu, S. Stavrakis, A.J. deMello,
- 604 Microfluidics for extracellular vesicle separation and mimetic synthesis: Recent
- advances and future perspectives, Chem. Eng. J. 404 (2021) 126110.
- 606 [15] D. Yang, W. Zhang, H. Zhang, F. Zhang, L. Chen, L. Ma, L.M. Larcher, S. Chen, N. Liu,
- 607 Q. Zhao, P.H.L. Tran, C. Chen, R.N. Veedu, T. Wang, Progress, opportunity, and
- 608 perspective on exosome isolation Efforts for efficient exosome-based theranostics,
- 609 Theranostics 10(8) (2020) 3684-3707.
- 610 [16] W.T. Su, H.J. Li, W.W. Chen, J.H. Qin, Microfluidic strategies for label-free exosomes
- 611 isolation and analysis, TrAC -Trends Anal. Chem. 118 (2019) 686-698.
- 612 [17] S.J. Lin, Z.X. Yu, D. Chen, Z.G. Wang, J.M. Miao, Q.C. Li, D.Y. Zhang, J. Song, D.X.
- 613 Cui, Progress in microfluidics-based exosome separation and detection technologies for
 614 diagnostic applications, Small 16(9) (2020).
- 615 [18] M. Garcia-Contreras, S.H. Shah, A. Tamayo, P.D. Robbins, R.B. Golberg, A.J. Mendez,
- 616 C. Ricordi, Plasma-derived exosome characterization reveals a distinct microRNA
- 617 signature in long duration Type 1 diabetes, Sci. Rep. 7 (2017) 10.
- [19] M.N. Madison, J.L. Welch, C.M. Okeoma, Isolation of exosomes from semen for in vitro
 uptake and HIV-1 infection assays, Bio-protocol 7(7) (2017) 21.

621	M. Yáñez-Mó, M. Valés-Gómez, High sensitivity detection of extracellular vesicles
622	immune-captured from urine by conventional flow cytometry, Sci. Rep. 9(1) (2019) 1-
623	12.
624	[21] H. Kim, S. Shin, ExoCAS-2: Rapid and pure isolation of exosomes by anionic exchange
625	using magnetic beads, Biomedicines 9(1) (2021) 12.
626	[22] X. Fang, C. Chen, B. Liu, Z. Ma, F. Hu, H. Li, H. Gu, H. Xu, A magnetic bead-mediated
627	selective adsorption strategy for extracellular vesicle separation and purification, Acta
628	Biomater. (2021).
629	[23] K. Zhang, Y. Yue, S. Wu, W. Liu, J. Shi, Z. Zhang, Rapid capture and nondestructive
630	release of extracellular vesicles using aptamer-based magnetic isolation, ACS Sensors
631	4(5) (2019) 1245-1251.

- 632 [24] D. Brambilla, L. Sola, A.M. Ferretti, E. Chiodi, N. Zarovni, D. Fortunato, M. Criscuoli,
- V. Dolo, I. Giusti, V. Murdica, EV separation: Release of intact extracellular vesicles
 immunocaptured on magnetic particles, Anal. Chem. 93(13) (2021) 5476-5483.
- 635 [25] C. Liu, X. Xu, B. Li, B. Situ, W. Pan, Y. Hu, T. An, S. Yao, L. Zheng, Single-exosome-
- 636 counting immunoassays for cancer diagnostics, Nano Lett. 18(7) (2018) 4226-4232.
- 637 [26] J. Ko, Y. Wang, K. Sheng, D.A. Weitz, R. Weissleder, Sequencing-based protein
- 638 analysis of single extracellular vesicles, ACS Nano 15(3) (2021) 5631-5638.
- 639 [27] T.D. Mai, D. Ferraro, N. Aboud, R. Renault, M. Serra, N.T. Tran, J.-L. Viovy, C.
- 640 Smadja, S. Descroix, M. Taverna, Single-step immunoassays and microfluidic droplet
- 641 operation: Towards a versatile approach for detection of amyloid-beta peptide-based
- biomarkers of Alzheimer's disease, Sens. Actuators B 255 (2018) 2126-2135.

[20] C. Campos-Silva, H. Suárez, R. Jara-Acevedo, E. Linares-Espinós, L. Martinez-Piñeiro,

643	[28] M. Morani, T.D. Mai, Z. Krupova, P. Defrenaix, E. Multia, ML. Riekkola, M. Taverna,
644	Electrokinetic characterization of extracellular vesicles with capillary electrophoresis: a
645	new tool for their identification and quantification, Anal. Chim. Acta (2020).
646	[29] D. Volodkin, V. Ball, P. Schaaf, J.C. Voegel, H. Mohwald, Complexation of
647	phosphocholine liposomes with polylysine: Stabilization by surface coverage versus
648	aggregation, Biochim. Biophys. Acta-Biomembr. 1768(2) (2007) 280-290.
649	[30] D. Volodkin, H. Mohwald, J.C. Voegel, V. Ball, Coating of negatively charged
650	liposomes by polylysine: Drug release study, J. Control. Release 117(1) (2007) 111-
651	120.
652	[31] Y. Takechi, H. Tanaka, H. Kitayama, H. Yoshii, M. Tanaka, H. Saito, Comparative study
653	on the interaction of cell-penetrating polycationic polymers with lipid membranes,
654	Chem. Phys. Lipids 165(1) (2012) 51-58.
655	[32] A. Gorman, K.R. Hossain, F. Cornelius, R.J. Clarke, Penetration of phospholipid
656	membranes by poly-L-lysine depends on cholesterol and phospholipid composition,
657	Biochim. Biophys. Acta-Biomembr. 1862(2) (2020) 10.
658	[33] A. Ali-Cherif, S. Begolo, S. Descroix, JL. Viovy, L. Malaquin, Programmable magnetic
659	tweezers and droplet microfluidic device for high-throughput nanoliter multi-step
660	assays, Angew. Chem. Int. Ed. 51(43) (2012) 10765-10769.
661	[34] U. Gündüz, Viscosity prediction of polyethylene glycol-dextran-water solutions used in
662	aqueous two-phase systems, J. Chromatogr. B 743(1-2) (2000) 181-185.
663	[35] M.A. Stremler, F.R. Haselton, H. Aref, Designing for chaos: Applications of chaotic
664	advection at the microscale, Philos. Trans. R. Soc. A-Math. Phys. Eng. Sci. 362(1818)
665	(2004) 1019-1036.

666	[36] D. Bedding, C. Hidrovo, Asme, Investigation of mixing in colliding droplets generated in
667	flow-focusing configurations using laser induced fluorescence, Amer. Soc. Mechanical
668	Engineers, New York, 2016.
669	[37] D. Bachurski, M. Schuldner, P.H. Nguyen, A. Malz, K.S. Reiners, P.C. Grenzi, F.
670	Babatz, A.C. Schauss, H.P. Hansen, M. Hallek, E.P. von Strandmann, Extracellular
671	vesicle measurements with nanoparticle tracking analysis - An accuracy and
672	repeatability comparison between NanoSight NS300 and ZetaView, J. Extracell.
673	Vesicles 8(1) (2019).
674	[38] J.A. Jaros, P.C. Guest, S. Bahn, D. Martins-de-Souza, Affinity depletion of plasma and
675	serum for mass spectrometry-based proteome analysis, Proteomics for Biomarker
676	Discovery, Springer 2013, pp. 1-11.
677	[39] K. Takov, D.M. Yellon, S.M. Davidson, Comparison of small extracellular vesicles
678	isolated from plasma by ultracentrifugation or size-exclusion chromatography: yield,
679	purity and functional potential, J. Extracell. Vesicles 8(1) (2019) 1560809.
680	[40] K. Brennan, K. Martin, S. FitzGerald, J. O'Sullivan, Y. Wu, A. Blanco, C. Richardson,
681	M. Mc Gee, A comparison of methods for the isolation and separation of extracellular
682	vesicles from protein and lipid particles in human serum, Sci. Rep. 10(1) (2020) 1-13.
683	

