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5	A microfluidic approach to fabricate sucrose decorated liposomes with increased
6	uptake in breast cancer cells
7	
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14 Abstract

15 Nanocarriers are known to control the non-selective activity of conventional chemotherapies that is still a major limitation in the fight against cancer. Developing a targeted drug delivery system is an urgent 16 need to decrease the side effects and increase the drug's efficiency. Most cancer cells show an increased 17 sugar consumption compared to healthy cells due to the deregulation of sugar transporters. 18 Consequently, liposomes, as a biocompatible nanocarrier, could be surface decorated by sugars to 19 enhance drug targeting into cancer cells. Our work outlines a new strategy to easily manufacture 20 sucrose decorated liposomes using sucrose stearate, a biocompatible and biodegradable non-ionic 21 surfactant, with a scalable microfluidic approach. Sucrose decorated liposomes were loaded with 22 23 berberine hydrochloride, a well-known phytochemical compound to investigate its effects on triple-24 negative breast cancer cells (MDA-MB-231). Using the microfluidic manufacturing system, we 25 prepared berberine-loaded liposomes using a mixture of phosphatidylcholine and cholesterol with and without sucrose stearate with a size up to 140 nm and narrow polydispersity. Stability was confirmed 26 for 90 days, and the *in vitro* release profile was evaluated. The formulations showed acceptable *in vitro* 27 28 biocompatibility and significantly higher anti-proliferative effect on MDA-MB-231 cell line. These 29 results have been confirmed by an increased uptake evaluated by flow cytometry and confocal 30 microscopy. Taken together, our findings represent an innovative, easy, and scalable approach to obtain

- sugar decorated liposomal formulations without any surface-chemistry reactions. They can be
 potentially used as an anticancer targeted drug delivery system.
- 33 Keywords: Additive manufacturing; 3D printed microfluidic chips; nanomedicine; sucrose esters;
- 34 Surface modification.

35 **1. Introduction**

36 Based on World Health Organization statistics, breast cancers are the most prevalent cancer and the main reason for death in women (1). About 10-20 % of the diagnosed breast cancers are sorted as triple-37 negative breast cancer (TNBC) which is the most problematic subtype that has no therapeutic target 38 39 due to estrogen (ER), progesterone (PR), and HER2 protein lack of expression (2). A major drawback 40 for cancer therapy is the non-selectivity of conventional therapies which cause adverse side effects and a significant decrease in drug efficiency (3,4). Therefore, there is an imperative need to investigate new 41 42 therapeutic agents and targeted drug delivery systems (DDS) to cure invasive breast cancer more 43 efficiently.

44 Among nano DDS, liposomes are small spherical structures (5) that could be modified with different 45 active targeting agents (6). As a drug delivery system, liposomes have proper bioavailability, size controlling measures, long half-lives, control release features, low risk to benefit ratio (5), self-46 assembly potential, and the ability to protect drugs to have long-lasting activation (7). Thin-film 47 hydration, ethanol injection, and proliposome-liposome method are the conventional ways to develop 48 liposomes (8). With all these methods developing a proper liposome encounter some challenges, such 49 as high dispersed particles, batch to batch variability, time-consuming preparation, and non-50 reproducible formulations (8,9). 51

Microfluidics is an innovative, scalable liposome manufacturing method that can overcome these issues 52 by controlling the flow conduction of the fluids through micrometer channels (10,11). These 53 microchannels could have different geometries (12) which can affect the flow and consequently the 54 self-assembly of the nanocarriers (11). In recent years, by the advent of microfluidic technology, which 55 56 is a simple and automized technique, the physical characteristics of the liposomes could be predictable, 57 tunable, and reproducible by the control of the main microfluidic parameters such as flow rate ratio (FRR) and total flow rate (TFR) (9,13). Subsequently, the microfluidic method allows the preparation 58 59 of liposomes in a more reproducible and time-saving manner (12) to deliver a precise amount of drugs 60 (14). Furthermore, liposomes have the ability to entrap both hydrophilic and hydrophobic drugs due to the amphiphilic nature (5,7). 61

Berberine, which has been chosen as a model drug in this work, is an isoquinoline alkaloid bioactive
which can be isolated mainly from Barberry (*Berberis vulgaris* L. belongs to the Berberidaceae family)
and attracts considerable interest due to its pharmacological effects. Investigations on berberine are

mostly on its anti-inflammatory (15-18), and anti-microbial properties (19-21). Moreover, there are 65 some in vitro and in vivo investigations on berberine's effect on different cancer cell lines, such as 66 67 prostatic (22,23), gastrointestinal (24-26), hepatic (27), dermal (28), and breast (27,29). It has been demonstrated that berberine can arrest the G0/G1 cell cycle and induce apoptosis in breast cancer cells 68 with estrogen receptors (27) and various cell lines (30). Despite all these characteristics and therapeutic 69 70 effects of berberine, low solubility (31) and low bioavailability (0.68%), which is probably related to 71 berberine structure as quaternary ammonium (32), are some of the limitations for berberine clinical 72 usage.

73 In this work, liposomes and sucrose decorated liposomes were prepared using the microfluidic method 74 throughout innovative 3D printed chips with two different microchannel geometries that have been 75 developed previously by our group (33). Liposomes were prepared using soybean phosphatidylcholine 76 (PC), and cholesterol, meanwhile for sucrose decorated liposomes, sucrose stearate was added as a renewable, and functional non-ionic surfactant that allows the presence of a sugar moiety on the surface 77 78 of the nanocarrier (34). While there is deregulation in cancer cells sugar transporter, they take and 79 consume more sugar than the normal cell lines (35). Pathophysiological and specific molecular 80 characteristics of cancer cells could be exploited to reach targeted DDS (6,36). Different targeting agents such as peptides, folic acid (3,4), antibodies (4), and carbohydrates (3,4,36–39) have been used 81 82 to facilitate targeted drug delivery in cancer therapies. In spite of the fact that the targeting moieties are usually attached by chemical reactions (4,6) our work insert the sugar moieties on the surface of 83 84 liposomes with the aim of reaching an active targeting on cancer cells with no surface chemistry steps 85 needed before or after the formulation of the nanocarrier. Modifying nanocarriers with carbohydrate 86 targeting agents could enhance the drug uptake by the cancer cells (3,4) by active and passive 87 mechanisms (6). The integration of sucrose ester could enhance the drug efficacy on cancer cells growth inhibition by increasing the nanovesicles uptake by cancer cells. The colloidal systems were 88 89 characterized by their average particle size, polydispersity index (PDI), and encapsulation efficiency (EE%). The characteristics of optimized liposomes were evaluated by Fourier-transformed infrared 90 91 spectroscopy (FTIR), and thermogravimetric analysis (TGA). Cytocompatibility of the loaded and 92 unloaded liposomes was performed on AC16 cardiomyocyte cells by Sulforhodamine B (SRB) assay. Moreover, the antiproliferative effects of the mentioned formulations were tested to confirm the effect 93 of liposome to increase berberine's bioavailability and the sucrose ester effects on enhancing 94

95 liposome's antiproliferative effectivity. Cell uptake of the different formulations prepared was96 evaluated by flow cytometry coupled with confocal microscopy.

97

98 2. Materials and methods

99 2.1. Materials and cell lines

Berberine hydrochloride (BBH) was purchased from A.C.E.F. (Italy), sucrose stearate (SS) was
obtained from Chem Service (USA), cholesterol (Chol) was kindly obtained from CRODA (UK).
soybean phosphatidylcholine (PC) (soybean lecithin, 94% of phosphatidylcholine) was kindly provided
by Lipoid (Germany). Polypropylene (PP) was kindly gifted from BASF (Germany). All the other
solvents used were analytical grade.

105

106 2.2. 3D printing of microfluidic chips

107 The 3D printed chips were designed and printed as described before (9). Briefly, the design of the microfluidic chip was optimized using a computer-aided design (CAD) to obtain an effective passive 108 109 micromixing with a "zigzag" bas-relief (Z-chip) and "split and recombine" channels (C-chip). Thereafter, the chips were printed using polypropylene with a fused deposition modeling (FDM) 3D 110 printer (Ultimaker 3, Ultimaker, The Netherlands) at a print speed of 25 mm/s and with a nozzle 111 temperature of 220 °C (0.25 mm nozzle). This allowed obtaining leak-free and semi-transparent 112 113 devices. Probe needles were used to connect the chip to syringe pumps (Aladdin, WPI Europe, Germany) through PTFE tubing. 114

115

116 *2.3. Preparations of the liposomes by the microfluidic method*

For the preparation of normal liposome (NL) by the microfluidic method, a total concentration of 8 mg/mL of PC, and Chol with the weight ratio of 3:1 was dissolved in the ethanolic phase. Thereafter, the 3D printed chips were connected through PTFE tubing to two syringes, containing water and ethanolic phase, that were settled on syringe pumps to control the flow. The ethanolic and water phases were mixed with FRR (water/ethanol) of 2:1 and TFR 8 mL/min. Finally, the flow-out liposomes were collected from the chip's outlet. The ethanol in the final formulation was discharged with one round of centrifuge at 17200 RCF for 60 min at 4 °C. Thereafter, the pellets were resuspended in water to reach 124 the primary concentration. In order to prepare sucrose decorated liposomes (SL), the total concentration

of 8 mg/mL of SS, PC, and Chol with the weight ratio of 3:1:1 and 3:1:2 was dissolved in the ethanolic

phase. The rest of the process is as for normal liposomes with FRR 2:1 and different TFR (8, 10, 12,

and 14 mL/min). To obtain BBH loaded liposomes BBH@NL and BBH@SL, 3 mg/mL of BBH were

previously dissolved in the water phase at 60 °C and then mixed through the microfluidic device with

- 129 the other excipients contained in the ethanolic phase.
- 130

131 *2.4. Characterization of colloidal systems*

Physical characteristics of the formulation were investigated by measuring the average particle size (Z-average) and polydispersity index (PDI) using Malvern Zetasizer Nano S instrument (Malvern Instrument Ltd, UK). Prior to the measurements, the formulations were diluted at 1:10 in distilled water.
For further characterization, attenuated total reflectance Fourier-transformed infrared spectroscopy (ATR-FTIR, Spectrum Two FT-IR spectrometer with ATR accessory, Perkin Elmer, MA, USA) were operated at 400–4000 cm⁻¹ to evaluate the interaction between BBH, SS, Chol, PC, SL, BBH@SL, NL, BBH@NL. The FTIR test was performed on freeze-dried liposomes.

Moreover, thermo-gravimetric analysis (TGA) was used to investigate the thermal behavior of the component based on their weight loss with increasing temperature. TGA was performed using a TGA 4000 (PerkinElmer, Norwalk, USA) equipped with an intercooler (Intracooler 2, PerkinElmer, Norwalk, USA) in an inert nitrogen atmosphere. Chol+PC (NL), Chol+PC+BBH (BBH@NL), Chol+PC+SS (SL), Chol+PC+SS+BBH (BBH@SL) were analyzed in the temperature range of 30 to 500 °C with the heating rate of 20 °C/min and 10 °C/min from 30 to 80° C and 80 to 500° C, respectively, under nitrogen atmosphere with the gas flow of 30 mL/min.

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147 2.5. Quantitative determination of berberine by HPLC

The amount of BBH in the experiments performed was evaluated by high-performance liquid chromatography (HPLC Agilent 1260 Infinity II, Agilent, USA). The mobile phase was a mixture of water and acetonitrile (65:35 v/v) with 0.05 % of Trifluoroacetic acid (TFA) at a flow rate of 1 mL/min. 151 The injection volume was 20 μ L in the column EC-C18, 100 × 4.6 mm, 2.7 μ m column (Agilent, USA) 152 monitored at 345 nm.

153

154 2.6. Encapsulation efficiency studies

Encapsulation efficiency (EE%) studies were assessed by direct method. Firstly, a calibration curve of BBH was performed with five concentrations in the range of 0.001 to 0.01 mg/mL with a coefficient of determination (R²) of 0.9986. The collected liposomes from the chip were centrifuged at 17200 RCF at 4° C for 60 min to settle the drug-containing liposomes. The supernatant was discharged to remove unloaded BBH and ethanol. Subsequently, 1 mL of methanol was added to dissolve the liposome structure and extract the encapsulated BBH. Then, the amount of loaded BBH was measured by HPLC.

161 Accordingly, the encapsulation efficiency (EE%) of the microfluidic liposome was calculated with the

162 following formula through the direct method.

163 $EE\% = (E_{drug}/T_{drug}) \times 100$

In the above formulation, E_{drug} is referred to the encapsulated amount of drug measured by the HPLC
 and T_{drug} is the total amount of used drug in the primary formulation.

166

167 *2.7. Stability test*

The stability of the optimized formulation was evaluated at 4 and 25 °C for up to 90 days in terms of size, PDI, and EE%. The measurements have been performed after 7, 14, 30, 60 and 90 days.

- 170
- 171 *2.8. Release studies*
- 172 2.8.1. In vitro release of berberine hydrochloride from liposomes

173 The release of BBH from NL and SL was investigated in phosphate-buffered saline (PBS pH 7.4) and

acetate buffer (pH 5.5) to simulate physiologic and cancerous cells pH (40), respectively. Briefly, 1 mL

of the BBH loaded liposomes (BBH@NL and BBH@SL) and free BBH at a concentration equal to the

176 BBH encapsulated into the liposome were placed in a dialysis bag (MW cut off 12k-14k Da,

177 Spectra/PorTM, Spectrum Labs, USA) and immersed into 50 mL of PBS and acetate buffer continuously

- 178 stirred at 100 rpm keeping the system at 37 °C. At each time point (0.25, 0.5, 1, 2, 3, 4, and 6 h), 1 mL
- of the medium was collected and replaced with the same amount of fresh release medium. The BBHconcentrations were measured with HPLC.
- 181 2.8.2. Mathematical Modeling of the Kinetics Release

The drug release data of the samples were modeled using the first-order, Higuchi, and Peppas-Salhin mathematical models using Origin Software (Origin Pro 2021, OriginLab, USA). The adjusted coefficient of determination (adjusted- R^2) was used to select the best-fitting model.

185

186 2.9. In vitro cytocompatibility on Human Cardiomyocyte Cell Line (AC16)

187 The human cardiomyocyte cell line AC16 was grown in Gibco Dulbecco's Modified Eagle Medium: Nutrient Mixture F-12 (DMEM/F-12) supplemented with 12.5 % FBS, 1 % penicillin/streptomycin 100 188 U/mL, and 2 mM L-glutamine. Cells were maintained in T25 flasks in a CO₂ incubator at 37 °C and 5 189 % CO₂. Cytocompatibility of different formulations with and without SS and BBH (BBH@SL, SL, 190 191 BBH@NL, NL) was studied by Sulforhodamine B (SRB) assays by evaluating the cell protein content. AC16 (4x10³ cells/well) were seeded in 96-well plates and treated with: a) NL 125, 250, 500 µg/mL; 192 193 b) SL 125, 250, 500 µg/mL; c) free BBH 3, 6, 12 µg/mL (corresponding to the amount of encapsulated BBH in the nanocarriers); d) BBH@NL 125, 250, 500 µg/mL; and e) BBH@SL 125, 250, 500 µg/ml. 194 195 After 24 h of incubation, test compounds were removed, cells were washed with PBS, and then fixed by adding trichloroacetic acid (TCA) 50 % (w/v). The plate was incubated for 1 h at 4 °C and then 196 197 rinsed with water 3 times. The plate was dried at room temperature and 0.4 % SRB was added to the wells and let the cells match the color for 30 min. Eventually, cells were rinsed with acetic acid 1 % as 198 many times to remove the unincorporated color. To solubilize the incorporated dye, 10mM tris was 199 added and the absorbance was measured at 570 nm in a microplate reader (Multiskan FC, Thermo 200 201 Scientific) (41). Data were expressed as a percentage (%) versus non-treated cells (controls).

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203

205 2.10. Antiproliferative studies on epithelial, human breast cancer cell line MDA-MB-231

206 The human triple-negative breast cancer cell line MDA-MB-231 was grown in DMEM medium supplemented 207 with 10 % FBS, 1 % penicillin/streptomycin 100 U/mL, 2 mM L-glutamine, and 1 % non-essential amino acid 208 solution. Cells were maintained in T25 flasks in a CO₂ incubator at 37 °C and 5 % CO₂. In order to investigate 209 the advantages of loaded drug compared to the free one, SS effect, and the microfluidic BBH-loaded liposomes on MDA-MB-231 breast cancer cells, different concentrations of NL, SL, BBH@NL, and 210 BBH@SL (125, 250, 500 µg/mL) were seeded in 96-well plates containing 5×10^{3} /well MDA-MB-211 212 231 cells and incubated for 24 h. Afterward, the cytotoxic effects of the microfluidic liposomes were 213 evaluated by SRB assays as explained before.

214

215 *2.11. Cell uptake studies*

To evaluate NL and SL uptake by MDA-MB-231, BBH-loaded liposomes were labeled with Rhodamine (1% w/w of 18:1 Lyss Rhod PE, Avanti Polar Lipid, USA). The fluorescent lipid was solved in the ethanolic phase together with the other excipients and incorporated during liposome microfluidic assembly.

Cancer cells were seeded in 12-well plates $(1.5 \times 10^5 \text{ cells/well})$ and treated with rhodamine-labeled NL and SL 250 µg/mL for 4 and 24 h. After the incubation time, liposomes were removed, cells detached by trypsinization, and washed with PBS at 1200 rpm for 10 min.

To label and trace lysosomes, the acidotropic dye LysoTracker Deep Red (LTDR) (Thermo Fisher Scientific, Waltham, MA, USA) was used. The amount of fluorescence obtained from staining with LysoTracker is directly related to the volume of lysosome-related organelles in a cell (42). Cell pellets were resuspended in 300 μl PBS containing 100 nM of LysoTracker. After 30 min of incubation, red lysosomal fluorescence was detected by flow cytometry and confocal microscopy (43).

228 CD71 antibody is specific for the human transferrin receptor, which is essential for iron transport into 229 proliferating cells (44); indeed, the presence of iron within the cell is necessary for cell proliferation 230 and is responsible for many cell functions such as DNA synthesis, oxygen sensing, and transitioning 231 from the G1 to the S phase in the cell cycle (45). To evaluate CD71 surface expression, FITC-232 conjugated anti-CD71 monoclonal antibody (clone L01.1) (BD Biosciences, San Jose CA, USA) was 233 added to 70 μ L of cell pellet, at dilutions according to the manufacturer's instructions. After 20 min of 234 incubation at RT, samples were acquired by flow cytometry and confocal microscopy.

To investigate the involvement of sugar receptors in SL uptake, MDA-MB-231 were seeded in 12-well plates $(1.5 \times 10^5 \text{ cells/well})$ and pre-treated with sucrose 0.75 M for 20 min at 37°C to block sucrose receptors (46) before incubation with rhodamine-labeled SL 250 µg/mL for two different periods (20 min and 40 min).

239

240 2.12. Cytometric Investigations

Cytometric experiments were carried out with a FACSCanto II flow cytometer (BD, Franklin, Lakes,
NJ, USA) equipped with an argon laser (Blue, Excitation 488 nm), a helium-neon laser (Red, Excitation
633 nm), and a solid-state diode laser (Violet, Ex 405 nm). Analyses were performed with the
FACSDivaTM software (BD); 10,000 cell events were acquired for each sample.

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246 2.13. Confocal Microscopy

Confocal microscopy analyses were performed with a Leica TCS SP5 II confocal microscope (Leica
Microsystem, Germany) with 488, 543, and 633 nm illuminations and oil-immersed objectives. For
confocal live imaging, MatTek glass-bottom chambers (MatTek Corporation, Bratislava, Slovak
Republic) were used. The images were further processed and analyzed in ImageJ software (NIH,
Bethesda, MD, USA).

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253 2.14. Statistical Analysis

All data were expressed as the mean \pm standard deviation (SD) based on at least three tests. Statistical analysis was carried out using Origin software (Origin Pro 2021b). A *P* value <0.05 was considered statistically significant.

258 **3. Results and discussion**

259 3.1. Microfluidic assembly and characterization of liposomes

260 In this study, we prepared a new class of microfluidic assembled liposomes decorated with sucrose to 261 increase the uptake in MDA-MB-231 breast cancer cells. BBH was successfully loaded into this 262 innovative system to formulate liposomes. The sucrose decoration was obtained using sucrose stearate 263 that belongs to the class of sugar esters, non-ionic surfactants that can be effectively used alone or in 264 combination with other lipids to formulate colloidal drug delivery systems (34) The use of these 265 surfactants allows to insert the sugar moieties on the surface of liposomes with the aim of reaching an active targeting on cancer cells with no surface chemistry steps needed before or after the formulation 266 267 of the nanocarrier. To obtain the optimum ratio of SS, PC, and Chol we characterized the liposomes in terms of size and stability starting from different weight ratios (3:1:1, 3:1:2) and different microfluidic 268 269 parameters (TFR and FRR) using the 3D printed C-chip and Z-chip (Fig. 1). Our results (Fig. 2) showed that the 3:1:2 ratio possessed both lower size and higher stability due to increased amount of cholesterol, 270 which can arrange alkyl chains ordering of the lipid and increase membrane density (47), while the 271 liposomes with 3:1:1 ratio were not stable and they showed 40 nm increase in the size after one week. 272 273 Moreover, different trials with TFR (8, 10, 12, and 14 mL/min) and the two different chips (C and Zchip) showed smaller sizes using Z-chip and 14 mL/min (Fig. 2). Based on the liposome size and 274 stability, the finest TFR and FRR are 14 and 2:1, respectively. The highest concentration of BBH (3 275 mg/mL) without any precipitation was used and solved in water at 60 °C. 276

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278





Fig. 2. Sucrose decorated liposomes optimization considering different TFR (8, 10, 12, 14 mL/min) and T-shape chips (Cchip and Z-chip) with SS: PC: Chol ratio of a) 3:1:1 b) 3:1:2. Data are reported as the mean of three independent replication
of experiments ± SD.

Based on the characterization, we choose the ratio of 3:1:2 of SS:PC:Chol, FRR 2:1 (water: ethanolic phase), and TFR (14 mL/min) for SL, and the ratio of 3:1 of PC:Chol has been chosen for NL. NL shows an average size and PDI of 130 ± 0.06 nm and 0.15 ± 0.004 , respectively; and physical characterization of SL by DLS shows a good particle distribution (PDI 0.06 ± 0.001) with an average size of 140 ± 2.4 nm. The EE% of the BBH@NL and BBH@SL investigated by HPLC was 3.5 and 7%, respectively.

320 *3.2. Stability test*

The stability of the formulation was evaluated at two different temperatures (4, 25 °C) for up to 90 days. The formulation resulted stable at 4 °C meanwhile, a slight increase in size and PDI was seen after 90 days in the 25 °C condition. The amount of encapsulated drug was more stable at 4 °C and shows 1.04%, 1.26%, and 1,33% decrease in 7, 30, and 90 days. While more significant decrease in EE% has been seen in 25 °C which shows 1.08 %, 1.6%, and 2.38% decrease in 7, 30, and 90 days, respectively.

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332								
	4 °C		25 °C					
	Time (Days)	Size (nm)	PDI PDI	इन्ड्रि	Size (nm)	PDI	EE%	
334	7	141.2±0.01	0.06±0.01	1.04% ↓	140.1±6.1	0.06±0.01	1.08%↓	
335	14	139.75±0.45	0.059 ± 0.004	-	141.6±0.2	0.053±0.01	-	
336	30	144±1.3	0.056±0.001	1.26% ↓	151.35±1.05	0.058±0.002	1.6 % ↓	
337	60	145.65±0.05	0.053±0.003	-	160.65±1.45	0.053±0.001	-	
338	90	145.65±0.63	0.048±0.006	1.33%↓	176.05±2.7	0.065 ± 0.004	2.38%↓	

Table 1. Characterization of microfluidic liposomes. The decrease is announced based on the EE% at day zero. Data are
 reported as the mean of three independent replication of experiments ± SD.

339

340 *3.3. Fourier-transform infrared spectroscopy (FTIR)*

FTIR analysis was carried out to evaluate the chemical structure of the microfluidic assembled 341 342 liposomes (Fig. 3). In the spectrum of BBH@NL and BBH@SL, peaks at 1505 and 1331 cm⁻¹ are assigned to the presence of BBH in the BBH loaded liposome's structure. In addition, the peak at 1728 343 344 cm⁻¹ is related to the existence of SS in both SL and BBH@SL. In the liposome's spectrum, the peak at around 3342 cm⁻¹ is related to cholesterol with a slight shift to the right and a remarkable increase in 345 intensity due to merging the characteristic peaks of cholesterol and the low-intensity peak of PC. The 346 slight shift to the right might be evidence of the hydrophobic interaction and hydrogen bonding in the 347 intermolecular structure of the liposomes (48). IR spectrum of BBH revealed an intense band at around 348 1003 cm⁻¹ and 1363 cm⁻¹, indicating the symmetric O-C-O stretch of the dioxolane ring and C-H group 349 wag vibration in C_6H_2 and C_5H_2 rings (49). In the BBH spectrum, the band 2845 cm⁻¹ indicates alkane 350 C-H stretching, and 1635 cm⁻¹ and 1331 cm⁻¹ refer to medium bending N-H and C-N in the amine 351 group, respectively. The band at 1568 cm⁻¹ and 1505 cm⁻¹ show medium stretching C=C in cyclic 352 alkene (50). In the spectrum of PC, the peaks at 2924 cm⁻¹ and 2854 cm⁻¹ are ascribed to C-H stretching 353 group in the structure. The peaks around 1735 cm⁻¹, 1249 cm⁻¹, and 1173 cm⁻¹ are due to the stretching 354 band of the ester carbonyl group, moderate stretching C-N amine band, and strong C-O stretching ester 355 band, respectively (48). In the cholesterol spectrum, the characteristic peaks can be seen at 3405 cm⁻¹, 356 2866 cm⁻¹, 1466 cm⁻¹, 1376 cm⁻¹, and 1056 cm⁻¹ which refers to stretching hydroxyl group, symmetric 357

358 C-H stretching vibration, asymmetric stretching vibration of methylene and methyl group, bending 359 vibration of methylene and methyl group and bending vibration of C-O group, respectively (34). The 360 peak at 1728 cm⁻¹ shows the characteristic peak of SS (51) which is referred to a strong C=O band in 361 an α , β -unsaturated ester.



Fig. 3. Physicochemical characterization by FTIR analysis of a) BBH, SS, SL, and BBH@SL and b) BBH, PC, Chol, NL,
 BBH@NL demonstrated that interactions between components of NL and SL were able to slightly change the location of
 bands, indicating liposome formation.

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366 *3.4. Thermal gravimetric analysis (TGA) characterization*

Principal data on the composition of the liposomes, their water content, thermal stability, and their mass 367 368 loss in specific temperatures (0- 500 °C) were obtained from TGA characterization. TGA curves showed the rates of water evaporation, water binding capacity, and stability of BBH, Chol, SS, PC, NL, 369 BBH@NL, SL, BBH@SL (Fig. 4 a, b) (52). BBH presented thermal stability up to 350 °C showing a 370 four-step degradation in the TGA curve from 350 to 500 °C confirming data reported in the literature 371 372 (53). Sucrose ester's thermal behavior showed a broad step of mass loss decomposition, starting in the 373 100-250 °C range, which is ascribed to a random molecular decomposition (54). PC thermal degradation was found at approximately 97 °C and the TGA analyses of Cholesterol showed that the 374 weight loss starts at 220-280 °C (55). The total weight loss of BBH, SS, PC, and Chol in thermal 375 degradation was 44.8%, 96.4%, 94.9%, and 97.3%, respectively. The TGA graph of the NL and 376 BBH@NL shows a thermal degradation refers to water loss starting at 100 °C and 5% of thermal 377 degradation at 150 °C; on the contrary, no water loss occurred in SL and BBH@SL. The onset of initial 378

weight loss is around 256, 263, 218, 214 °C for NL, BBH@NL, SL, BBH@SL, respectively. The
occurrence of the maximum degradation could be demonstrated from derivative thermogravimetry
(DTG) analysis (Fig. 4c). The maximum degradation happens at <300 °C for the SL (253.4 °C and
296.3 °C for SL without and with BBH) and >300 °C for normal liposomes (341.65 °C and 332.9 °C
for NL and BBH@NL, respectively). The residues in the NL and SL formulations are 5.15% and 6.14%
of the initial weight, while the residue for the formulations containing the BBH is 10.17% and 17% for
BBH@NL and BBH@SL, which refers to undegraded BBH in the formulation.



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Fig. 4. a, b) TGA (upper thermograms) c) and DTG (lower blue thermograms). The DTG thermograms are plotted for a
better understanding of the rate of material weight changes upon heating up to 500 °C.

- 390 *3.5. Release studies*
- 391 *3.5.1.* In vitro release of berberine hydrochloride from liposomes
- 392 The release profiles of BBH from BBH@SL and BBH@NL were studied at two different pH values of
- 393 7.4 and 5.5 to simulate physiologic and cancerous cells pH, respectively (40) (Fig. 5 a, b). The release

results in both mediums (acetate buffer and PBS) show that loading BBH into liposomes could decrease the release rate of the drug and provide a sustained release profile compared to free BBH solution. Moreover, BBH@SL showed faster drug release compared to BBH@NL due to the presence of SS, a permeability enhancer (56), which increases the liposome membrane permeability and drug release compared to the liposome without SS (0.23± 0.003 mg/mL, 0.2±0.007 mg/mL in BBH@SL and 0.14±0.004 mg/mL, 0.13±0.001 mg/mL in BBH@NL in acetate buffer and PBS, respectively).



Fig. 5. BBH release from BBH@NL, BBH@SL and free BBH solution in a) acetate buffer (pH 5.5) b) PBS (pH 7.4) at 37
°C. Data are reported as the mean of three independent replication of experiments ± SD.

402

403 *3.5.2. Mathematical Models of Releasing Kinetics*

The results obtained from the modeling of drug release profiles for each system evaluated at different environmental pH's, as well as their respective kinetic constants and adjusted-R² are shown in Table 2, where $k_1(h^{-1})$ is the first-order rate constant, $k_H(h^{-\frac{1}{2}})$ is the dissolution constant, $k_d(h^{-0,43})$ is the Fickian diffusional contribution and $k_s(h^{-0,43})$ the matrix swelling contribution (57).

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Sample	Mathematical Model	Kinetic constants	R^2
NL-	First-order: ${}^{M_t}/{M_0} = e^{-k_1 t}$	$k_1 = 0.03 \pm 0.01(h^{-1})$	0.33
buffer	Higuchi: ${}^{M_t} / {}_{M_0} = k_H \sqrt[2]{t}$	$k_H = 0.30 \pm 0.08 \ (h^{-1})$	0.64
	Peppas-Salhin:	<i>m</i> = 0.43	0.98
	$\frac{M_t}{M_0} = k_d \times t^m + k_s \times t^{2m}$	$k_d = 1.06 \pm 0.06 \ (h^{-0.43})$	
		$k_s = -0.30 \pm 0.03 \ (h^{-0.43})$	
NL-PBS	First order: ${}^{M_t}/{}_{M_0} = e^{-k_1 t}$	$k_1 = 0.03 \pm 0.01(h^{-1})$	0.49
	Higuchi: ${}^{M_t}/{M_0} = k_H \sqrt[2]{t}$	$k_H = 0.3 \pm 0.1 \ (h^{-1})$	0.66
	Peppas-Salhin:	m = 0.43	0.97
	$M_t / M_0 = k_d \times t^m + k_s \times t^{2m}$	$k_d = 0.99 \pm 0.01 \ (h^{-0.43})$	
	, ,	$k_s = -0.3 \pm 0.3 \ (h^{-0.43})$	
SL-	First-order: ${}^{M_t}/{}_{M_0} = e^{-k_1 t}$	$k_1 = 0.03 \pm 0.01(h^{-1})$	0.45
buffer	Higuchi: ${}^{M_t}/{M_0} = k_H \sqrt[2]{t}$	$k_H = 0.3 \pm 0.1 \ (h^{-1})$	0.70
	Peppas-Salhin:	m = 0.43	0.99
	$M_t / M_0 = k_d \times t^m + k_s \times t^{2m}$	$k_d = 1.10 \pm 0.04 \ (h^{-0.43})$	
	, ,	$k_s = -0.31 \pm 0.02 \ (h^{-0.43})$	
SL-PBS	First order: ${}^{M_t}\!/_{M_0} = e^{-k_1 t}$	$k_1 = 0.03 \pm 0.01(h^{-1})$	0.42
	Higuchi: ${}^{M_t}/{M_0} = k_H \sqrt[2]{t}$	$k_H = 0.3 \pm 0.1 \ (h^{-1})$	0.66
	Peppas-Salhin:	m = 0.43	0.98
	$M_t / M_0 = k_d \times t^m + k_s \times t^{2m}$	$k_d = 1.06 \pm 0.04 \ (h^{-0.43})$	
	, 0	$k_s = -0.31 \pm 0.02 \ (h^{-0.43})$	

412 **Table 2.** Kinetic constants are derived from the application of mathematical models to the drug release profiles.

The drug release profiles of all systems were better fitted by the Peppas-Salhin model as indicated by the R² values (**Table 2 and Fig. 6**). The Peppas-Salhin is a semiempirical model used to obtain information about the contributions of diffusion and matrix relaxation (or erosion) on drug release. The

417 coefficient m is the purely Fickian diffusion exponent related to a drug delivery system of any 418 geometrical shape. As can be seen in Table 2, for all the systems, the k_d kinetic constant is significantly 419 higher than the k_2 kinetic constant, which, in turn, displayed a not significative negative value. This 420 means that the drug release occurs by the Fickian diffusion mechanism without any contribution of the 421 matrix swelling or erosion (58,59).



Fig. 6. Drug release profile of Free BBH, BBH@NL, and BBH@SL in a) acetate buffer b) PBS fitted by the Peppas-Salhin
models.

424

425 *3.6. In vitro cytocompatibility on Human Cardiomyocyte Cell Line (AC16)*

426 Cytotoxicity of the liposomes toward normal cardiomyocyte cells (AC16 cells) was evaluated by SRB 427 assays (**Fig. 7**). No cytotoxic effects of NL and SL formulations were evident after 24 h treatment at 428 all the concentrations tested. Only in the cases of free BBH ($12 \mu g/mL$) and BBH@SL ($500 \mu g/mL$) a 429 statistically significant cell growth inhibition (up to 20 %) was observed by SRB test.

430



432

Fig. 7. AC16 cell growth (%) after 24 h treatment with NL, SL formulations (125, 250, 500 μg/mL) and free BBH (3, 6, 12 μg/mL) and BBH@NL and BBH@SL formulations 125, 250, 500 μg/mL (loaded with BBH 3, 6, 12 μg/mL). Cell viability
was investigated using SRB test. Data are expressed as the mean± SD (n=3). *p<0.05 vs CTR (Tukey's test).

437 *3.7. In vitro anti-proliferative test*

438 The anti-proliferative effect of the nanocarrier was determined by SRB assay (Fig. 8). Both SL and NL with and without BBH were studied to investigate if the presence of sucrose ester in the formulation 439 440 could affect the antiproliferative activity on MDA-MB-231 cancer cells. Free BBH at the same 441 concentration of the BBH loaded into liposomes was evaluated as a comparison. The SRB test demonstrates no cytotoxic effects of empty nanocarriers towards MDA-MB-231 after 24 h treatment. 442 443 Free BBH did not show inhibition of cell growth in the range of concentrations tested (3-12 μ g/mL 444 comparable to the concentration of encapsulated BBH), in accordance with previous findings which 445 demonstrated BBH cytotoxicity towards MDA-MB-231 cells only at very high doses (>20 µg/ml) (60). Significant cytotoxicity was observed in BBH@SL already at the concentration of 250 µg/mL 446 447 (corresponding to 6 µg/mL loaded BBH) meanwhile, at the same concentration, BBH@NL did not 448 show significant growth inhibition. Moreover, greater inhibition of cell growth was found by BBH@SL 500 µg/mL (-64 % vs. CTR) as compared to BBH@NL 500 µg/mL (-40 % vs, CTR). The difference 449 between two formulations was statistically significant proving that the presence of the sugar moiety on 450

451 the surface of BBH@SL gives a positive effect on the efficacy of the sugar decorated nanocarrier 452 compared to the normal ones.



453 Fig. 8. MDA-MB-231 cell growth (%) after 24 h treatment with NL, SL formulations (125, 250, 500 µg/mL) and free BBH 454 (3, 6, 12 µg/mL corresponding to BBH loaded in SL) and BBH@NL and BBH@SL formulations 125, 250, 500 µg/mL. 455 Cell viability was investigated using the SRB test. Data are expressed as the mean± SD (n=3). **p<0.01, ****p<0.0001 vs 456 CTR and free BB (Tukey's test).

457

458 3.8. Cell uptake studies

459 The uptake of NL and SL by MDA-MB-231 breast cancer cells was analyzed by flow cytometry and 460 confocal microscopy after 4h (Fig. 9) and 24h (Fig. 10) to evaluate differences between the two 461 nanocarriers. In the MDA-MB-231 cell line tested after 4h, a well-detectable, bright fluorescence was registered in SL samples (Fig. 9a, c, e) whereas in NL-treated cells only a negligible uptake was 462 463 detected, demonstrating that NL has a very slower uptake into the breast cancer cells compared to SL 464 which present the sugar moieties on the surface. The higher and, particularly, the faster liposome uptake in SL formulation could explain the more efficient anti-proliferative activity of SL on MDA-MB-231 465 466 evaluated by the SRB assay.

467 To further investigate the mechanisms that drive these strong differences in the internalization of NL

468 and SL, we detected the expression of the transferrin receptor (CD71, normally involved in the iron 469

cellular uptake and strongly expressed on cells with a high proliferation rate (Fig. 9b).

Indeed, CD71 is a marker of poor prognosis in breast cancer and can predict response to Tamoxifen
(61). We assessed the surface expression of CD71 on MDA-MB-231 cells, finding, in SL samples,
moderate but significantly lower levels than control cells (Fig. 9d).





Fig 9. a) Single confocal optical section to evaluate liposome uptake (red) of SL treated MDA-MB-231 cells, after 4 h. b) Single confocal optical sections of liposome uptake (red) and CD71 (green) on MDA-MB-231 cells after 4 h. Scale bars 10 μ m c) Statistical histogram of the liposome uptake after 4 h. Data are expressed as the mean± SD (n=3). ***p<0.0001 vs CTR and NL (Tukey's test). d) Statistical histogram of CD71 surface expression after 4 h. Data are expressed as the mean± SD (n=3). *p<0.01, vs CTR and NL (Tukey's test) e) Representative cytometric histogram of liposome internalization for Control, SL, and NL samples after 4 h.

481 CD71 also mediates the transport of IgA food complexes in epithelial cells (62) and, it may be involved 482 in a specific binding (63). Of note, the moderate decrease of CD71 fluorescence observed in SL samples 483 highlights a mild contribution of this surface molecule. We extended the analyses to 24 h after liposome 484 administration. Confocal, flow cytometric, and statistic evaluations are shown in (**Fig. 10**).

485 Data highlight the extensive and significant uptake of NL, slightly greater than SL uptake (Fig. 10 c,
486 f). In line with CD71 behavior observed for SL after 4 h, NL samples exhibit a mild decrease in CD71
487 expression (Fig. 10 d).



Fig. 10 a) Single confocal optical section to evaluate liposome uptake (red) of SL treated MDA-MB-231 cells, after 24 h.
b) Single confocal optical sections showing CD71 expression (green) and liposomes (red) on MDA-MB-231 cells after 24
h. Scale bars 10 μm. c) Statistical histogram of liposome uptake after 24 h. Data are expressed as the mean± SD (n=3).

491 **p<0.001, ***p<0.0001 vs CTR and NL (Tukey's test). d) Statistical histogram of CD71 after 24 h. e) Statistical histogram

- 492 of LTDR MFI after 24 h. Data are expressed as the mean \pm SD (n=3). ***p<0.0001 vs CTR and NL (Tukey's test). f)
- 493 Representative cytometric histogram of liposome internalization for Control, SL, and NL samples after 24 h.
- 494

Furthermore, to follow, at least in part, the intracellular journey of liposomes, we labeled breast cancer cells by the lysosomotropic probe LysoTracker Deep Red (LTG). Statistical and cytometric histograms are shown in (**Fig. 10 e**) and depict the increase of the number and functions of lysosomal structures and other acidic organelles (64). Furthermore, the confocal analysis highlights a co-localization of SL and acidic organelles (yellow spots) whereas no co-localized distribution is observable in NL treated cells.

These findings reveal profound differences also for the intracellular fate of the two different liposomal formulations, with a net SL tropism for lysosomal compartment, prospecting their possible use to target acidic vacuoles. Indeed, this peculiarity strengthens the evidence of the greater uptake of SL that, although degraded into lysosomes, reveal a similar NL fluorescence signal, after 24h, attesting a continuous, sustained SL uptake from the extracellular medium.

Finally, since cancer cells overexpress a wide variety of carbohydrate-binding receptors (3,4,36,65),
including members of the C-type lectin receptor, Siglec (sialic acid-binding immunoglobulin-type
lectin), galectin families (65), and the well-known CD206 receptor, highly expressed on MDA-MB231 cells (66), sucrose receptor blocking study was performed.

To elucidate the capability to massively involve a varied pool of carbohydrate-binding receptors, a sucrose-cell pre-incubation was performed, as explained in Materials and Methods. The cellular uptake and its eventual sucrose-derived inhibition of SL were investigated after 20 and 40 minutes (**Fig. 11**). NL was not considered in this study since they did not show any uptake after 4h. A partial uptake inhibition was noticed after 20 minutes (around 20%) and after 40 minutes (around 30%) from starting the test. These results confirm that a sucrose-dependent uptake is present, but it is not the only one used to internalized the nanosized formulation.



Fig 11. Statistical histogram of liposome uptake and its inhibition after 20' and 40' in normal-sized breast cancer cells,
excluding cell debris. Data are expressed as the mean± SD (n=3). *p<0.01, ***p<0.0001 vs CTR and NL (Tukey's test).

520 4. Conclusions

In this work, we developed an innovative sugar-decorated colloidal drug delivery system assembled by 521 522 3D printed microfluidics. 3D printed microfluidics open the possibility to researchers to have easy access to this technology with the chance to personalize it based on their specific needs. Moreover, by 523 employing the microfluidic method we have the potential to increase the liposomes preparation rate to 524 525 enhance the very low conventional lab-scale rates. Our approach allows us to easily obtain stable and reproducible surface decorated delivery systems with no chemical steps needed before or after the 526 production. The presence of the sugar moieties on the nanocarrier demonstrated an improvement of the 527 cancer cell uptake resulting in an increased antiproliferative effect of the berberine-loaded sugar 528 529 decorated liposomes. Taken together, this work demonstrates that, by microfluidic, it is possible to 530 formulate drug-loaded sugar decorated liposomes with active targeting properties and tunable characteristics with an easy and scalable approach. Moreover, the proposed microfluidic assembled 531 532 BBH@SL is a potential candidate as an antiproliferative carrier against triple-negative breast cancer cells. 533

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