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# Mesoporous silica templated-albumin nanoparticles with high doxorubicin payload for drug delivery assessed with a 3-D tumor cell model

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## Abstract.

Human serum albumin (HSA) nanoparticles emerge as promising carriers for drug delivery. Among challenges, one important issue is the design of HSA nanoparticles with a low mean size of ca. 50 nm and having a high drug payload. The original strategy developed here is to use sacrificial mesoporous nanosilica templates having a diameter close to 30 nm to drive the protein nanocapsule formation. This new approach ensures first an efficient high drug loading (ca. 30%) of Doxorubicin (DOX) in the porous silica by functionalizing silica with an aminosiloxane layer and then allows the one-step adsorption and the physical cross-linking of HSA by modifying the silica surface with isobutyramide (IBAM) groups. After silica template removal, homogenous DOX-loaded HSA nanocapsules (30-60 nm size) with high drug loading capacity (ca. 88 %) are thus formed. Such nanocapsules are shown efficient in multicellular tumor spheroid models (MCTS) of human hepatocarcinoma cells by their significant growth inhibition with respect to controls. Such a new synthesis approach paves the way toward new protein based nanocarriers for drug delivery.

## 1. Introduction

Designing nanoparticles as controlled drug delivery systems for cancer therapy has attracted great attention to overcome the problems of chemotherapeutics side effects such as the well-known cardiotoxicity of anthracyclines.[1–5] Human serum albumin (HSA) derived nanoparticles show high potential for drug delivery.[6,7] HSA, the most abundant plasma protein (35-50 g/L human serum), has biocompatible, non-immunogenic and biodegradable properties.[8]

Furthermore, HSA shows high average half-life of 19 days which enhances protein tumor uptake.[9]

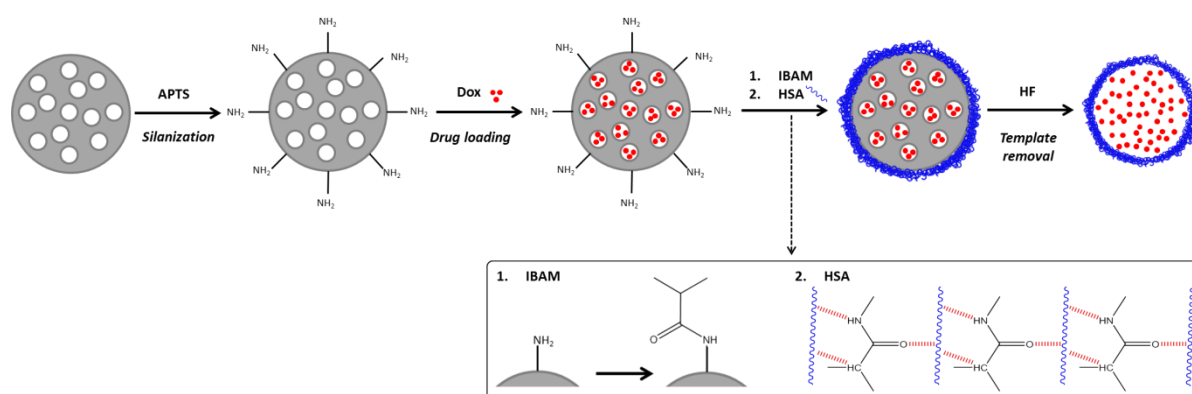
HSA nanoparticles can be prepared through a multitude of techniques, the main one being desolvation,[10–12] in which water soluble drugs are loaded either by incubation in the formed nanoparticles, or by incorporation in the albumin solution before particle formation and crosslinking.[13] For example Dreis *et al.*[14] prepared HSA nanoparticles and investigated the loading of the anti-cancer drug doxorubicin (Dox), by incubation (adsorption on particle surface) compared to incorporation. For both loading strategies, high loading efficiencies (70–95 wt%) were obtained and Dox loaded HSA nanoparticles showed improved anti-cancer effects on *in-vitro* cell cultures compared to free Dox molecules. However, one of the main important drawback generally encountered in such synthesis strategies is the use of toxic cross-linkers such as glutaraldehyde or formaldehyde. The potential release of such cross-linkers can induce some toxicity as glutaraldehyde is known to be genotoxic, mutagenic and carcinogenic.[15–17] Furthermore, there is a need to design drug delivery systems with mean sizes below 100 and even 50 nm to ensure their biodistribution and accumulation in diseases body parts.[18,19]

Considering the promising properties of HSA nanoparticles as drug delivery systems and to face the main challenges of high drug loading/delivery and mean sizes suitable to envision *in vivo* applications, an original strategy based on the use of silica templates has been innovatively adapted. Indeed, we and co-workers previously developed an approach to form tight and robust HSA micron and sub-micron sized particles assembled non-covalently *via isobutyramide* (IBAM) grafts and without the need of further cross-linking or addition of adjuvants.[20–22] In this strategy, a sacrificial template of silica is first modified with amine groups on which *isobutrylchloride* is grafted to form IBAM-functionalized surface. Then, HSA is coated on IBAM-modified silica by simple adsorption in aqueous solution. The driving force to immobilize HSA to IBAM groups was shown to be of a non-covalent nature and was assumed to result from strong hydrogen-bonding interactions.[20,21] Silica nanoparticles, which were extensively studied in the past two decades, appear as particularly relevant and original sacrificial templates. Indeed, they own outstanding intrinsic features like simple and well-referred synthesis processes leading to nanoparticles with controlled size and shape, tunable morphologies, tunable pore structure and easy surface modification.[23–25] Furthermore using plain or mesoporous silica nanoparticles influences the design of the final HSA nanoparticles obtained after complete dissolution of silica template with hydrofluoric acid (HF): full silica particles lead to the formation of HSA hollow capsules [21] whereas mesoporous silica nanoparticles with large porosities, allowing the diffusion of HSA into the pores, will form replicate HSA nanoparticles.[22] However the decrease of the mean size of such protein and polymer capsules, in order to reach the sub-micron scale is currently a great challenge, and to the best of our knowledge the use of template below 100 nm is scarcely investigated.[26,27]

Considering the specifications of drug delivery systems with a high drug loading, mesoporous silica nanoparticles (MSNs) are well-known drug delivery systems since Vallet-Regi *et al.* reported in 2001 the loading of drug molecules within MCM-41 materials.[28] Indeed, MSNs exhibit large

surface area and pore volume providing great potential for drug adsorption and loading within the pore channels. In addition, the feature of tunable porosities enables better control of the drug loading and the type of drug loaded. Indeed, MSNs with large porosities ( $> 3$  nm) can load large drug molecules, siRNA, plasmid, proteins, quantum dots, nanoparticles[29] whereas MSNs with small porosities ( $\sim 3$  nm) are well known for the loading of small drug molecules such as anthracyclines, ibuprofen, cisplatin, aspirin etc[30,31] Therefore, besides their use as sacrificial templates for the formation of self-supported macromolecule assemblies, MSNs can serve in a first place as nanocontainers for drug loading and drive after their removal the final shape of the protein-based particle.

Herein, we report the design of MSNs templated self-supported HSA nanocapsules (size below 50 nm) with a high Dox content and demonstrate their potential as drug delivery system using a 3D tumor cell model. The MSNs addressed for this study exhibited a raspberry (RB) morphology with a narrow size distribution centered at about 30 nm and small porosities of 3 nm.[32] They are thus called RB NPs. This porosity has the benefits on one hand to be large enough to load chemotherapeutics into it by impregnation but, on another hand, to be small enough to provide an efficient HSA coating at the surface of the RB NPs leading thus to hollow-capsules formation after silica removal. As illustrated in **Scheme 1**, RB NPs were first synthesized and their surface functionalized with amine groups by using aminopropyltriethoxysilane (APTS) prior Dox loading. Indeed, very recently, we showed that an aminosilane layer condensed at very high silane concentration at the surface of carbon nanotubes@MS composites allowed to reach very high payloads of Dox ( $\geq 50\%$ ) depending on the feed weight ratio (fwr).[33] After such drug loading, the amino functions from APTS were modified with *isobutrylchloride* to form IBAM moieties. Then, HSA was absorbed on the IBAM-functionalized surface forming a tight protein shell onto the RB NPs through assumed non-covalent H-bonding with IBAM. Finally, after removal of mesoporous silica sacrificial templates dissolved in HF/ $\text{NH}_4\text{F}$  buffer ( $\text{pH} \sim 5$ ) and dialysis to remove fluoride ions, self-supported Dox loaded HSA nanocapsules were formed. To assess their efficiency as drug delivery systems, cytotoxicity and growth inhibition studies were carried out on human hepatocarcinoma spheroids.



**Scheme 1.** Schematic representation of the Dox loaded HSA nanocapsules synthesis strategy.

## 2. Experimental

### A. Materials

Tetraethyl orthosilicate (TEOS, Sigma-Aldrich), ammonium hydroxide solution ( $\text{NH}_{3\text{aq},25\%}$ ), triethanolamine (TEAH<sub>3</sub>, Sigma-Aldrich), cetyltrimethyl-ammonium bromide (CTAB, Roth), Human Serum Albumin (HSA, Sigma-Aldrich), isobutrylchloride (IBC, Sigma-Aldrich), Doxorubicin hydrochloride (Dox, 99%, OChem Inc, DesPlaines, IL), N,N-dimethylformamide (DMF, Sigma-Aldrich), fluorescein isothiocyanate (FITC), 3-aminopropyl-triethoxysilane (APTES, Sigma-Aldrich) and with poly(2-hydroxyethyl methacrylate) (poly-HEMA, Sigma-Aldrich) were used as received.

### B. Procedures

*Synthesis of MSNs with a raspberry morphology (RB NPs).* The RB NPs were synthesized according to a modified literature method.[32] Briefly, 960 mg of CTAB were mixed with 230 mg of TEAH<sub>3</sub> in 63 mL of Milli-Q water. This mixture was first placed for 1h under sonication then 1h at 80°C until the components were completely dissolved. Next, 4 mL of TEOS was quickly added to the above solution under stirring. The mixture was stirred at 80 °C for 2 h. The as-synthesized RB NPs were filtered, washed and dried in an oven at 100 °C for 24 h. The resulting powders were then calcined at 550 °C for 6 h to burn the surfactant molecules left in the porosities.

*Amino-functionalization of RB: RB-NH<sub>2</sub> NPs.* To functionalize the RB NPs with a layer of primary amine groups, the calcined RB NPs were re-dispersed in 40 mL of EtOH at a concentration of 1.5 mg.mL<sup>-1</sup> with 960 μL of NH<sub>3</sub> and 4 mL of APTES and the mixture was stirred for 2 hours at room temperature. The amino-functionalized MSNs (RB-NH<sub>2</sub> NPs) were then washed three times with EtOH by centrifugation.

*Dox loading.* Dox loading into RB-NH<sub>2</sub> porosities was performed in Milli-Q water by simple soaking. RB-NH<sub>2</sub> NPs dispersed at 0.5 mg.mL<sup>-1</sup> were mixed with Dox at a feed weight ratio (fwr) of 100%.

$$fwr\% = \frac{\text{weight of Dox fed}}{\text{weight of NPs}} \times 100$$

The mixture was stirred at room temperature in the dark for 24 h. Then the Dox loaded RB-NH<sub>2</sub> NPs were collected by centrifugation.

*IBAM grafting and HSA coating.* Before the HSA adsorption, the RB-NH<sub>2</sub> NPs were functionalized with IBAM groups. In a typical procedure, 300 μL Et<sub>3</sub>N were mixed with 1.5 mL of RB-NH<sub>2</sub> at 10 mg.mL<sup>-1</sup> in DMF before the quick addition of IBC: DMF (405 μL: 3 mL) mixture forming a precipitate with RB-NH<sub>2</sub> NPs. After 1h of reaction under stirring, 15 mL of HSA at 2 mg.mL<sup>-1</sup> in Milli-Q water were added. The precipitate dissolved immediately after the addition of the aqueous solution and the mixture was put under stirring for additional 2h. The resulting RB-HSA NPs were collected and washed several time in Milli-Q water by centrifugation.

*Capsules formation by silica dissolution.* The RB sacrificial templates were dissolved with the use of ammonium fluoride buffered HF. Basically, 150  $\mu\text{L}$  of RB-HSA or RB-Dox-HSA NPs at 2  $\text{mg}\cdot\text{mL}^{-1}$  were mixed with 150  $\mu\text{L}$  of ammonium fluoride buffered HF (30 mM, pH 5) for 1 h before being dialyzed for 3 days against 3 mL of Milli-Q water (changed every day).

*Preparation of Huh7 multicellular tumor spheroid model (MCTS).* The Huh7-luc cells were cultured in a Dulbecco's Modified Eagle's Medium (DMEM) extended with 10 % of fetal calf serum (FCS), 5 % of PS and 1 % of non-essential amino acids. Prior to the culture, sterile 96-well plates were coated with poly-HEMA in order to avoid cell adhesion at the well-bottom. The coating was done by addition of a 0.5 % (w/v) poly-HEMA solution in ethanol (50  $\mu\text{L}$ ) into each well of a 96-well plate. For ensuring a total surface coating, the plate was put at 38°C for 24h until complete ethanol evaporation. The wells were then washed three times with PBS (100  $\mu\text{L}$ ). Adherent Huh-7 were treated with trypsin and suspended in complete medium at 5.3  $10^3$  cell/mL. The cell suspension (15  $\mu\text{L}$ ) was deposited as a drop on the inside face of the 96-well plate lid. The lid was then replaced on the PBS-filled plate and the cells in the hanging drop were left to grow in the incubator for 3 days. During these three days the cells present in the drop sedimented at the medium/air interface, aggregated and developed finally with subsequent proliferation and deposition of ECM into a multicellular tumor spheroid.[34] The PBS was then discarded and replaced with complete cell culture medium (100  $\mu\text{L}$ ). Each drop containing one single spheroid was then transferred from the lid into the well by centrifugation at 500 RPM for 1 min. MCTS were cultivated further at 37°C in 5 %  $\text{CO}_2$ .

*Evaluation of antitumoral effect on MCTS.* The experiments were carried out in octuplicate in 96-well plate using the spheroids as previously described. NPs were diluted at various concentrations in complete cell culture medium. NPs are immediately added to each spheroid by further dilution into the cell culture medium (100  $\mu\text{L}$ ). The spheroid growth was examined using a Nikon Eclipse TS100 microscope and a 10 X objective. Images were recorded at different days with a Nikon Color CCD DS-Fi2 Camera and a DS-L3 camera controller. The area (A) of each spheroid was measured using the NIH Image J software and reported in  $\mu\text{m}^2$ . The MCTS growth (G) was calculated relative to the initial area ( $A_0$ ; day 0) and express as relative growth versus control using the following equation:

$$G = \frac{P_t - P_0}{P_0} \times 100$$

Where

$$P_t = \frac{A_{tX}}{A_{t_0}}$$

And

$$P_c = \frac{A_{cX}}{A_{c0}}$$

with  $A_t$  the area of treated MCTS,  $A_c$  the area of control MCTS and X the number of days.

### C. Characterization Methods.

*TEM imaging.* The morphology and diameter of the RB MS NPs were determined by transmission electron microscopy (TEM, JEOL 2100) operating at an accelerating voltage of 200 kV. The average diameter of the NPs was calculated by measuring on ImageJ software at least 100 NPs from TEM images.

*SEM imaging.* The size and morphology of RB MS were characterized by scanning electron microscopy (Gemini500-ZEISS- 1nm resolution). One drop of the sample, in ethanol or water solution, was deposited on a silicon slide and then the solvent was evaporated.

*N<sub>2</sub> isotherm adsorption.* Nitrogen adsorption isotherms were obtained on a Micrometrics ASAP 2420 Accelerated Surface Area and Porosimetry System at 77K under continuous adsorption conditions. Brunauer-Emmett-Teller (BET) method was used to determine the surface area as well as the total internal pore volume at P/P<sub>0</sub> = 0.97. By applying the Barrett-Joyner-Halenda (BJH) method, the surface area, the cumulative pore volume, and the mesoporous pore size distribution (PSD) are determined during the adsorption and desorption steps. The microporosity pore size distribution (porosities < 1 nm) was calculated with the Horvath-Kawazoe model.

*Fourier Transform Infra-Red Spectroscopy (FTIR).* The surface composition of the NPs at different steps of synthesis was verified by Fourier Transform Infra-Red Spectroscopy (FTIR) (Perkin Elmer instrument).

*Thermogravimetric analysis (TGA).* TGA was carried out on a TA SDT 600 instrument in order to determine the Drug loading content (DLC) and the Coating content (CC):

$$DLC\% = \frac{\text{weight of Dox loaded}}{\text{weight of NPs}} \times 100$$

$$CC\% = \frac{\text{weight of HSA adsorbed}}{\text{weight of NPs}} \times 100$$

Before the analyses, the samples in suspensions were freeze dried. For the runs, the samples were heated from 25 to 100°C at a rate of 5°C / min.

*AFM imaging in liquid phase.* AFM images and force spectroscopy measurements were performed using a Bioscope Resolve instrument (Bruker Nano, Palaiseau, France). Nanoparticles were deposited onto PEI coated silicon wafer during 10 minutes and then extensively rinsed with water prior to morphology and mechanical analyses in liquid phase by AFM. Topographical images of nanoparticles were performed in peakforce tapping™ mode and mechanical analysis derived from peakforce QNM mode. Silicon nitride cantilevers of conical shape purchased from Bruker (Peakforce HIRS-FA, Bruker, France) with spring constant of about 0.32-0.35 nN/nm were used for both imaging and nanomechanical measurements. All images were recorded with a resolution of 256 × 256 pixels and a scan rate of 1 Hz. Nanomechanical properties of nanoparticles under solvent condition were extracted from the analysis of at least 4 peakforce captured (PFC) images at different locations over the silicon wafer surface. The elastic modulus *E* was calculated by analyzing the force-indentation curves according to the Sneddon model.[35] In this model, the Young modulus is related to the applied force according to the equation given below:

$$F = \frac{2E \tan \alpha}{\pi(1-\nu^2)} \delta^2 f_{BECC} \quad (1)$$

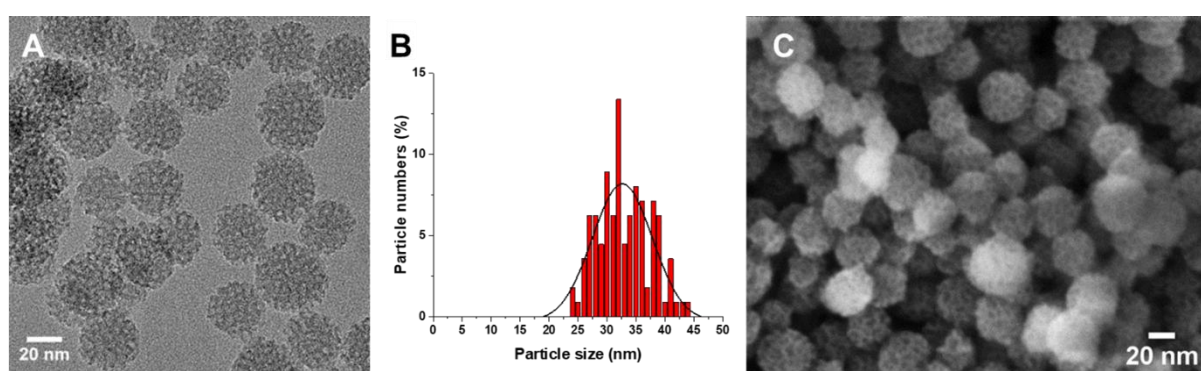
where  $\delta$  is the indentation depth,  $\nu$  the Poisson coefficient,  $\alpha$  the semi-top angle of the tip and  $f_{\text{BECC}}$  is the bottom effect cone correction function that take into account the presence of the substrate stiffness.[36] All the PFC were analyzed with an automatic Matlab algorithm described elsewhere.[37]

### 3. Results and discussion

#### A. Synthesis and characterization of RB NPs and surface modification with APTS

The morphology and structure of as-synthesized RB NPs before any functionalization were characterized by TEM and SEM imaging. As depicted in **Figure 1.A** the synthesis yielded uniform RB NPs of *ca.* 30 nm in size with apparent porosities throughout the entire particle. The RB NPs statistical average diameter was measured of *ca.*  $33 \pm 4$  nm and the resulting histogram is presented in **Figure 1.B**. The size distribution and morphology of the RB NPs were also confirmed by SEM images, which depict very nicely the nanostructure and demonstrated that the pore network was open to outside (**Figure 1.C**).

The  $\text{N}_2$  adsorption-desorption isotherms, (**Figure S1.A**) exhibited according to IUPAC classification, a type IV isotherm which is characteristic for mesoporous channel. The BET surface area and the total pore volume of the calcined RB NPs were measured at *ca.*  $708 \text{ m}^2\cdot\text{g}^{-1}$  and  $1.62 \text{ cm}^3\cdot\text{g}^{-1}$  respectively. These values are consistent to those describe in literature for RB NPs (BET surface area:  $675 \text{ m}^2\cdot\text{g}^{-1}$ ; total pore volume:  $1.27 \text{ cm}^3\cdot\text{g}^{-1}$  [32]). The corresponding pore size distribution (PSD) (**Figure S1.B**) calculated for pore sizes above 2 nm using the BJH method exhibited a single peak centered on 2.8 nm. With the Horvath-Kawazoe method (used for porosities  $< 1$  nm) a single peak at 0.7 nm was recorded. Via those two PSD calculation methods, we found out that these RB NPs showed a bimodal porosity centered at 7 and 28 Å.



**Figure 1.** A) TEM image of RB NPs. B) Particle size distribution of RB NPs from TEM pictures (>100 particles). C) SEM image of RB NPs.

RB NPs were then functionalized with amine groups with a concentration ensuring the formation an aminosilane layer providing high drug loading.[33] FTIR-ATR analyses were achieved to confirm the RB surface modification by APTS. **Figure S2** compares the IR spectra of RB NPs with RB-NH<sub>2</sub> NPs. The intense and broad absorption peak centered at  $1065 \text{ cm}^{-1}$  is assigned to Si-O-Si asymmetric stretching vibration, whereas the peak around  $800 \text{ cm}^{-1}$  is assigned to Si-O-Si symmetric stretching vibrations. The peak at  $970$

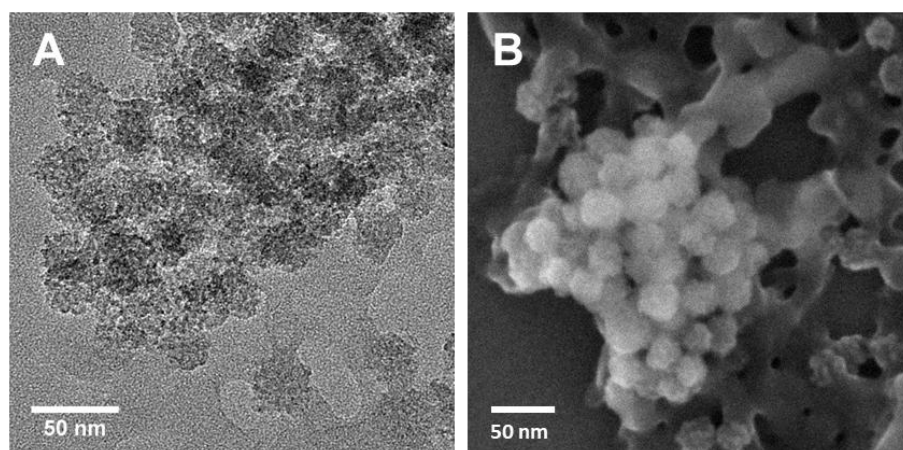


$\text{cm}^{-1}$  is attributed to Si-OH bond. Finally, the IR band at  $450 \text{ cm}^{-1}$  is due to O-Si-O bending vibrations. The appearance of new peaks for the RB-NH<sub>2</sub> sample at  $1356$  and  $1408 \text{ cm}^{-1}$ , specific for the stretching bands of C-N groups, and at  $1573$  and  $1618 \text{ cm}^{-1}$ , assigned to the bending vibration of N-H groups, proved the effective amino-functionalization of the RB surface.

## B. Synthesis and characterization of RB-Dox-HSA NPs

Then, RB-Dox-HSA and the corresponding Dox free RB-HSA NPs were obtained according to the procedure described in the Experimental section. Dox loading into RB-NH<sub>2</sub> porosities was performed in Milli-Q water by simple soaking (24h, room T) at a fwr of 100%. Then, the RB-NH<sub>2</sub>-Dox NPs and the corresponding Dox free RB-NH<sub>2</sub> NPs were functionalized with IBAM groups by reaction between amine groups and IBC reagents in the presence of Et<sub>3</sub>N. After 1h of reaction under stirring, HSA was adsorbed at  $2 \text{ mg}\cdot\text{mL}^{-1}$  in Milli-Q water and the resulting RB-DOX-HSA NPs or RB-HSA NPs were washed several times in water. The protein coating and the Dox loading were first characterized by ATR-IR (**Figure S3**). For both the RB-HSA and RB-Dox-HSA NPs samples in comparison to raw RB NPs, multiple peaks in the ranges of  $1300$ - $1800 \text{ cm}^{-1}$  and  $2800$ - $3000 \text{ cm}^{-1}$  evidenced the presence of various organic groups usually encountered with polypeptides.

The RB-Dox-HSA NPs were then visualized by TEM (**Figure 2.A**) and SEM (**Figure 2.B**) analyses. Both imaging showed RB NPs tangled in a dense HSA network. Thus, the RB NPs seemed to form large aggregates that were trapped into HSA. These particles being collected after centrifugation, we expect that any free albumin was separated from our sample. However, the drying of the TEM grids and then the vacuum conditions seem to have triggered the leaking of a part of the HSA coating and the partial formation of an HSA matrix on the TEM grid/SEM substrate.

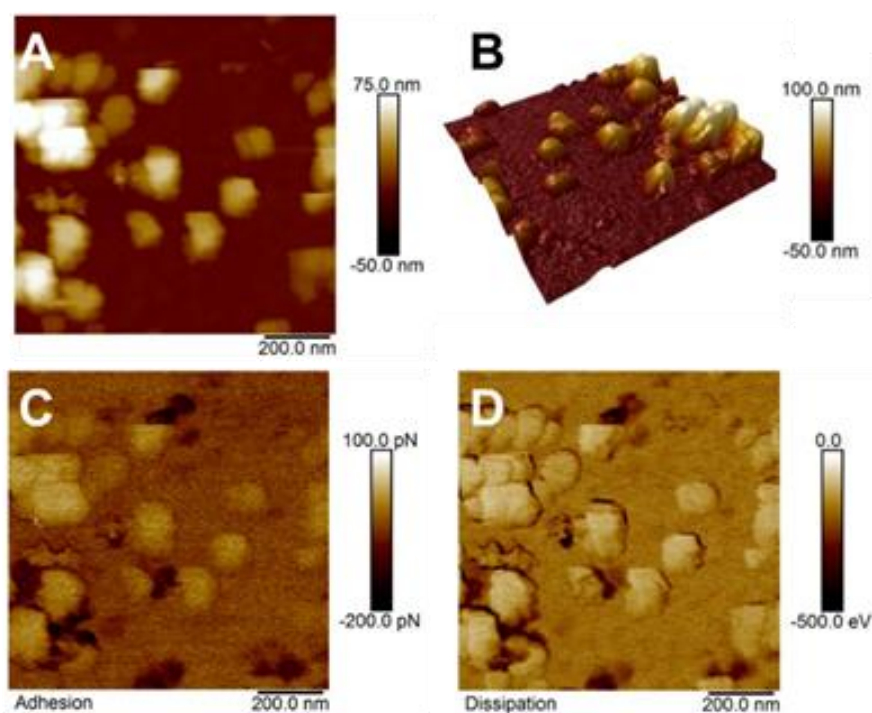


**Figure 2.** (A) TEM and (B) SEM images of RB-Dox-HSA-NPs in dried conditions from low to high magnifications respectively from the left to the right.

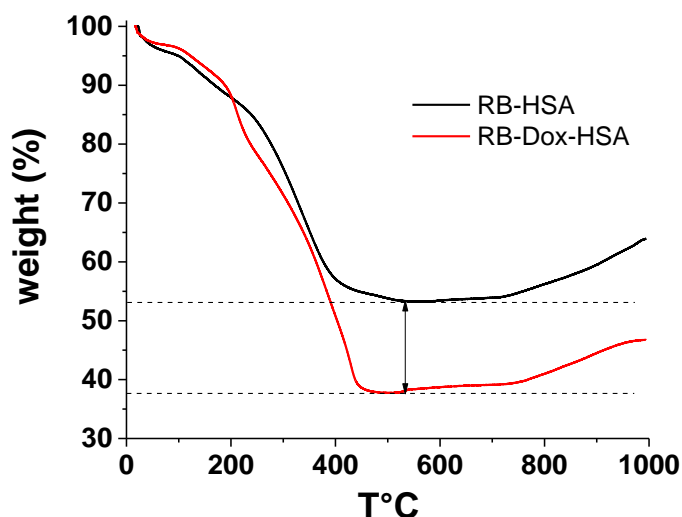
To ascertain this behaviour, AFM imaging in liquid phase was achieved on RB-DOX-HSA NPs in water (**Figure 3**). Careful inspection of images revealed aggregated and isolated particles uniformly distributed over the PEI-coated silicon wafer. Isolated nanoparticles were characterized by pseudo-spherical shape with a diameter in the range of  $30$ - $150 \text{ nm}$ . The average thickness of the isolated nanoparticles is about

67±23 nm. Concerning the aggregated nanoparticles, they are randomly distributed over the silicon wafer surface and their size never exceeds 300-400 nm for a thickness of about 60-150 nm. Moreover, the mappings of adhesion, energy dissipation and stiffness indicate homogeneity of both isolated and aggregated nanoparticles over the sample with average values of 285±49 pN, 0.18±0.09 keV and 2.28±0.17 MPa, respectively. The latter value is very low as compared to values previously reported for mesoporous silica.[38] Indeed, Young modulus of mesoporous silica is rather around 4-5 GPa and in the range of 10-100 MPa for nanocomposite gels blended with silica nanoparticles.[39,40] In our analysis, we found ca. 2 MPa and this low value is attributed to the layer of HSA proteins that embedded the hard silica nanoparticles.

Regarding the drug payload, as some Dox was released during the IBAM and HSA steps, the main difficulty in such drug loading studies was to determine the final DLC in the RB-Dox-HSA NPs. Indeed, the common analysis of supernatants by fluorescence or absorbance spectroscopy may induce some errors taking into account the different synthesis steps. Therefore, TGA analyses were run on RB-Dox-HSA and RB-HSA NPs samples to evaluate the Dox loading by comparing the weight loss of both samples. **Figure 4** shows the resulting TGA curves. The difference in weight loss between both samples suggested an efficient Dox loading. The first TGA curve on RB-HSA NPs indicated 53.3 wt.% of RB and 46.7 wt. % of HSA which corresponds to a protein coating content (CC) of 87.6 %. The second TGA curve obtained on RB-Dox-HSA NPs indicated 37.8 wt.% of RB and 62.2 wt.% of Dox-HSA. Considering the hypothesis that protein coating occurred in a similar way for both samples (with and without Dox) a CC of 87.6 % would give a weight-percentage of ~33 wt.% of HSA. Thus, the DLC of Dox is estimated at ~29 wt.% in the RB nanosilica.



**Figure 3.** AFM images of RB-Dox-HSA-NPs in liquid conditions (water).



**Figure 4.** TGA curves of RB-Dox-HSA and RB-HSA samples.

### C. Silica removal and formation of DOX-HSA NPs

With the aim to form silica free self-supported protein nanoparticles or capsules, RB sacrificial template was dissolved in HF/NH<sub>4</sub>F buffer. The resulting sample, Dox-HSA, was re-dispersed in water and formed a stable colloidal red suspension as pictured in **Figure 5.A** (photograph of the sample). Dynamic light scattering analysis (DLS) (intensity) performed on this sample, gave an average hydrodynamic size of 222±76 nm (**Figure 5. B**) indicating some effect of capsule aggregation which nevertheless did not lead to sedimentation with time.

Considering the DLC to be the percentage of the ratio between the mass of Dox and the mass of HSA capsules after complete dissolution of silica template, a high DLC value of 88 % was estimated. This value was calculated from the previous TGA analysis on RB-Dox-HSA sample considering that the entire RB sacrificial template was dissolved. **Table 1** compares DLC and DLE values of Dox loaded in HSA NPs reported in the literature[14,41–45] and the values obtained for our sample. Thus, compared to the various reported Dox encapsulation methods into HSA NPs, such as adsorption of Dox to preformed HSA NPs, incorporation of Dox to HSA matrix before desolvation or Dox conjugation to HSA molecules, our method of Dox soaking into sacrificial silica template led to high DLC's which is highly suitable for biological studies. That confirmed again that the formation of an aminosilane condensed at very high silane concentration is an efficient strategy to load high drug amount.[33]

**Table 1.** DLC and DLE values of Dox loading into HSA NPs from our sample compared to literature values

Ref.	Type of Dox loading	Size (nm)	DLC (%)	DLE (%)
<b>Our work</b>	Soaking into sacrificial silica template	43	88	-
Bae <i>et al.</i> <sup>39</sup>	Adsorption to preformed HSA NPs	200	4.8	94.5
Xu <i>et al.</i> <sup>40</sup>	Conjugation via disulfide bonds (via RPA) + physical adsorption	30	56.2	-

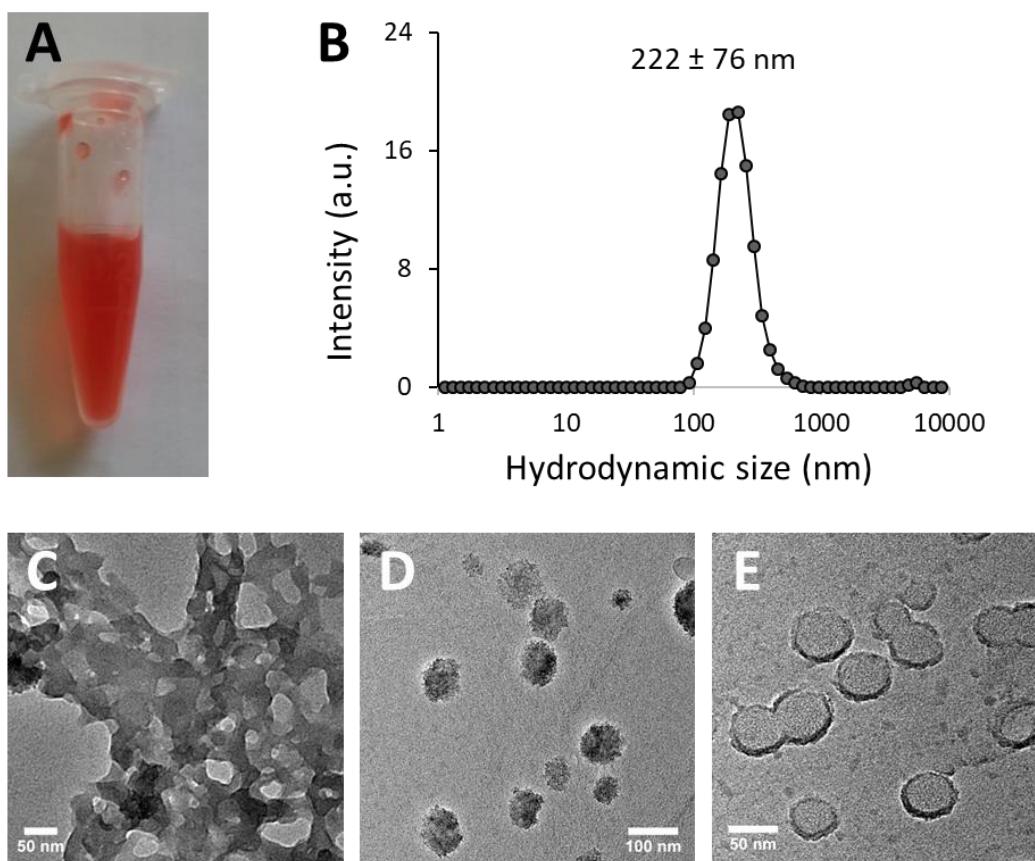
	Conjugation via disulfide bonds (via MPA) + physical adsorption		62.3	-
<i>Wagner et al.</i> <sup>41</sup>	Incubation to HSA before desolvation	400	5.7	-
<i>Dreis et al.</i> <sup>14</sup>	Adsorption to HSA in solution	-	14	75
	Adsorption to preformed HSA NPs	158	13.8	69
	Incorporation to HSA matrix before desolvation	404	4.82	96.3
<i>Lee et al.</i> <sup>42</sup>	Incorporation to HSA matrix before desolvation	183	5	-
<i>Li et al.</i> <sup>43</sup>	Adsorption to preformed BSA NPs	100	27.8	74

TEM analyses of the Dox-HSA sample were then performed. Three different kinds of morphologies were observed which may be due as described above to the TEM grids preparation and vacuum observation conditions. The first morphology, showed in **Figure 5.C**, was the major morphology observed for the Dox-HSA sample. This morphology was composed of a HSA network forming some kind of sticky like filaments that connected denser spherical HSA NPs together. These denser spherical HSA NPs exhibited sizes between 20 and 30 nm and thus, could correspond to HSA NPs formed from RB template dissolution as the RB NPs had sizes of 30 nm. Previous hypotheses could explain the formation of this filamentous network as it was also observed on the TEM and SEM images of RB-Dox-HSA (grapes stuck into a HSA matrix in **Figure 5**). A first assumption is that the HSA coating through IBAM grafts was not tight enough and that HSA tend to desorb from the RB surface creating this kind of network. A second assumption could be that this network simply results from an effect of leaking through sample drying on TEM grid before imaging.

The second morphology, as shown in TEM images in **Figure 5.D**, exhibited spherical HSA NPs of  $58 \pm 21$  nm. As this size was higher than the size of the original RB template, this could result for example from very thick HSA coatings on the RB NPs.

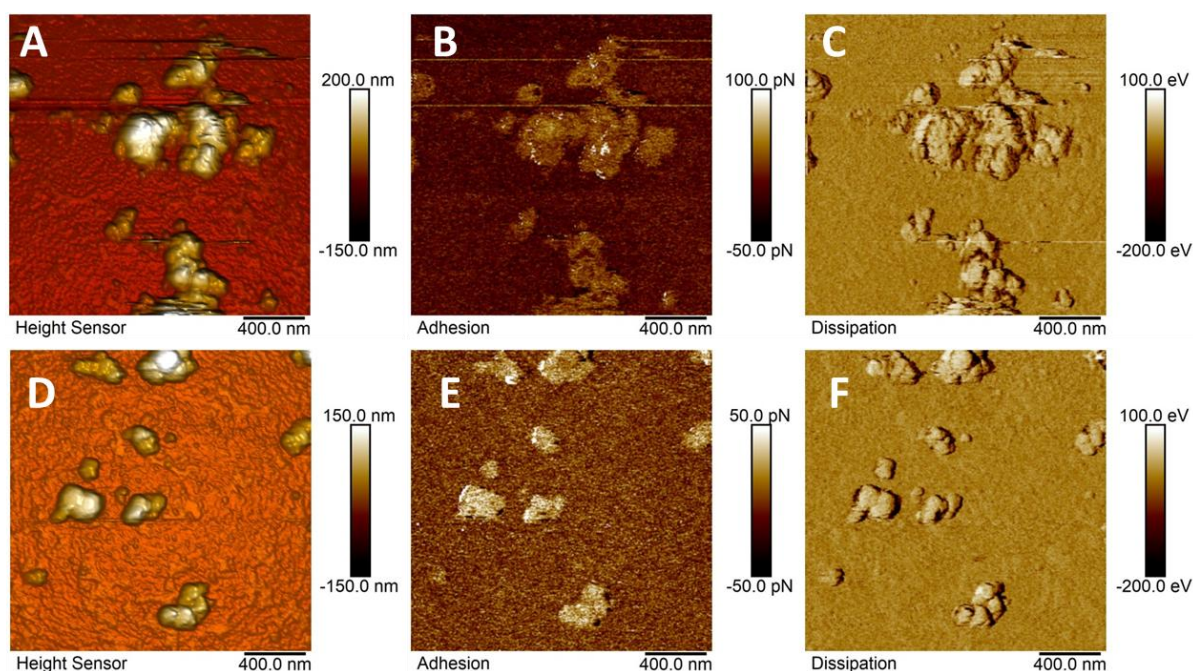
The third morphology, shown on TEM images in **Figures 5.E**, was typical for organic capsules. These capsules exhibited a size of  $43 \pm 10$  nm that corresponded to the RB sacrificial template size. This morphology should result from dried HSA NPs on the TEM grid.

Finally, the real morphology of the sample was rather difficult to determine through TEM and SEM analyses mostly because they involved drying this organic sample before imaging and imaging under vacuum conditions.



**Figure 5.** A photograph of the colloidal suspension of Dox-HSA NPs. B. DLS size distribution (intensity) of the red suspension. C-E. TEM images of the Dox-HSA NPs obtained after silica removal which showed three types of morphology : 1) HSA network like (C), 2) nanosphere like (D) and 3) nanocapsule like (E).

Thus to get additional insights in the morphology of the capsules in the hydrated state, we performed AFM in liquid conditions. AFM observations indicated that the HSA network (first morphology) was not anymore observed which tend to validate our previous hypothesis. AFM images showed that the Dox-HSA NPs in liquid conditions mostly appear as grapes of aggregated albumin nanospheres (**Figure 6**). The analysis of isolated nanoparticles evidenced pseudo-spherical shape with a diameter in the range of 50-150 nm. The average thickness of the isolated nanoparticles was found about 30-70 nm. Concerning the aggregated nanoparticles, they were randomly distributed over the silicon wafer surface and their size were measured in the range of 250-1000 nm for a thickness of about 150-250 nm. Moreover, the mappings of adhesion, energy dissipation and stiffness indicated homogeneity of both isolated and aggregated nanoparticles over the sample with average value of  $0.12 \pm 0.05$  MPa. The latter value is consistent with the silica dissolution and its absence within the nanocapsules as compared with the value of 2 MPa found in the presence of silica. Indeed, values between 0.01 to 0.5 MPa were already reported for polyelectrolytes and hydrogels nanocapsules.[46-48]



**Figure 6.** AFM imaging in liquid phase of the DOX-HSA capsules after silica removal forming aggregates of small capsules/vesicles mapped in two different zones (A and D). Images (B,C) on one hand and (E,F) on the other hand correspond respectively to adhesion and dissipation mapping associated respectively to the topographic images A and B.

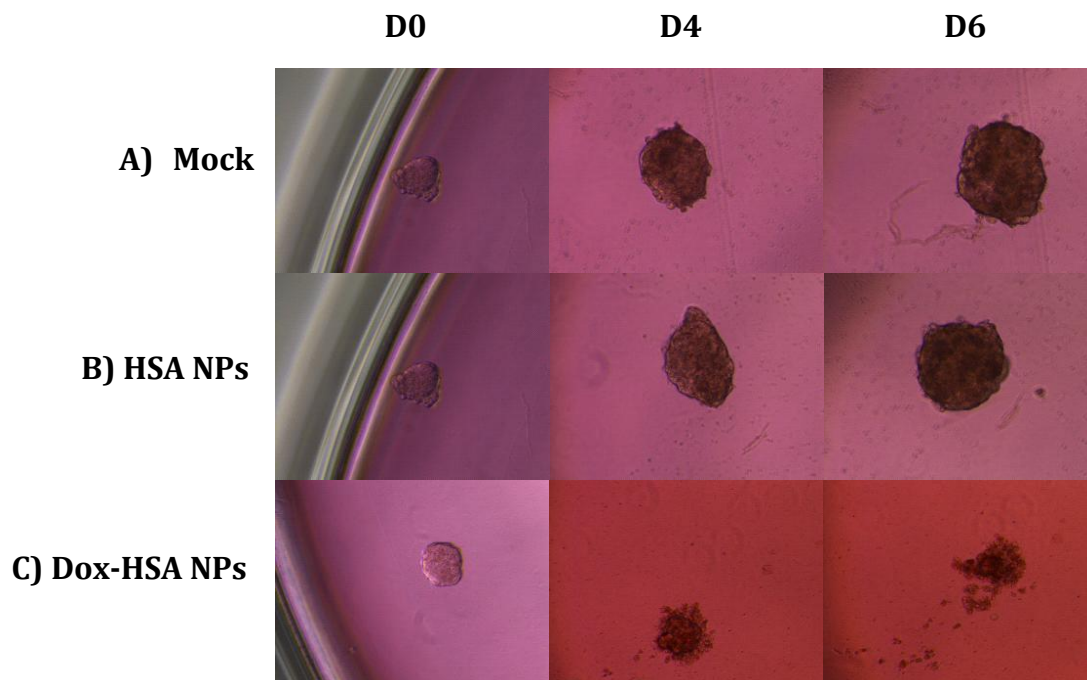
#### D. Drug delivery to multicellular spheroids and growth inhibition

Dox-HSA capsules were assessed directly to cancer cells assuming that the diversity of the cell environment would provide triggers for Dox delivery like proteolysis by secreted enzyme or lysosomal enzyme. For that, cytotoxicity assays were performed on a Huh7 multicellular tumor spheroid model (Huh7 MCTS). Such model reproduces more closely the *in vivo* behavior of tumor cells than 2D culture system. Indeed, even if two-dimensional (2D) cell monolayers have been conventionally used by the scientific community to evaluate drug efficacy, NPs cytotoxicity and their internalization, they poorly represent the complex tumor 3D microenvironment. Indeed, cells grown in 2D lack the extracellular matrix (ECM), as well as cell-cell contact and cell-matrix interactions. Thus, 3D culture models have emerged as an additional *in vitro* step study to assist drug product development and tend to reduce the requirement for unnecessary animal studies [49–57]. Among various methods to engineer 3D cancer model such as spheroids, bio-printing and assemblies, we developed multicellular tumor spheroids (MCTS) cell culture by aggregation of cells in hanging drop which is one of the easiest ways to form 3D tumor model from cancer cell lines.

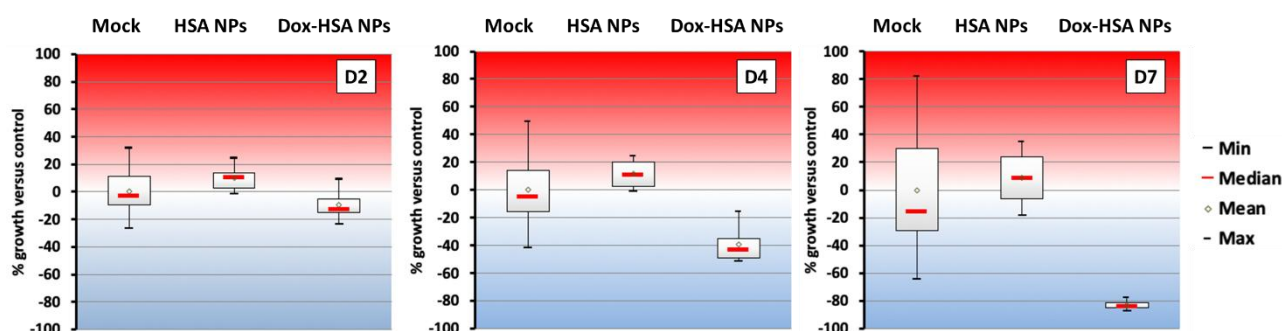
Thus we evaluated first the toxicity of Dox-HSA NPs and corresponding (Dox free) HSA NPs, by imaging the growth of the MCTS at various timepoints (2, 4 and 7 days). Huh7 MCTS were incubated with Dox-HSA and HSA NPs diluted in complete medium with concentrations of  $35.4 \mu\text{g}\cdot\text{mL}^{-1}$  of HSA for both samples and a Dox concentration of  $31 \mu\text{g}\cdot\text{mL}^{-1}$  for the Dox-HSA NPs sample. **Figure 7** shows representative optical microscopy pictures of Huh7 MCTS at different days and with both NPs in comparison to control experiment (Mock). It shows that the HSA NPs did not modify the growth compared to non-treated MCTS

however the cell growth for Dox-HSA NPs MCTS exhibited a strong inhibition and a different morphology that could be related to a massive cell death. Such images are characteristics of the Dox effect on MCTS.[58,59]

In the second series of experiments, measurements of MCTS surface area were also done after incubation of Dox-HSA NPs ( $17.7 \mu\text{g.mL}^{-1}$  of HSA and  $15.5 \mu\text{g.mL}^{-1}$  Dox) and of HSA NPs ( $17.7 \mu\text{g.mL}^{-1}$ ). The growth percentage was also calculated compared to the Mock. As shown on the results presented in **Figure 8**, Huh7 MCTS growth in presence of Dox-HSA NPs was readily inhibited. At this concentration tested, we can observe a decrease of 80 % compared to the control. As the growth is not modified when Huh7 MCTS are incubated with HSA NPs, the effect may be related to Dox release. Our particles are made of human serum albumin and can be taken up by cells at least those of the outer layer of the cell aggregate by non-specific endocytosis and or pinocytosis. Dox could therefore be released after HSA degradation by lysosomal proteases. Furthermore, we expect from other experiments (not published yet) that Dox can diffuse through the entire Huh7 MCTS even if the NPs internalization occurred only at the spheroid borders. So the effect recorded here is a proof that Dox can be released from the nanocapsules.



**Figure 7.** Photographs, taken by optical microscope (10X magnification), of A) non treated spheroids or Mock, B) spheroids treated with HSA NPs at  $35.4 \mu\text{g.mL}^{-1}$  of HSA and C) spheroids treated with Dox-HSA NPs at  $35.4 \mu\text{g.mL}^{-1}$  of HSA and  $31 \mu\text{g.mL}^{-1}$  of Doxorubicin at 0, 4 and 6 days of incubation (D0, D4 and D6).



**Figure 8.** Box plot representing percentage of growth of Huh7 MCTS either treated with HSA NPs or Dox-HSA NPs in comparison with non-treated ones (Mock) after 2, 4 and 7 days (D2, D4, D7) of incubation at concentrations of  $17,7 \mu\text{g.mL}^{-1}$  of HSA and  $15,5 \mu\text{g.mL}^{-1}$  Dox.

## Conclusion

In this study, we reported the synthesis of small HSA nanocapsules loaded with doxorubicin (Dox) using mesoporous silica RB NPs as sacrificial templates. This sacrificial template based synthesis allowed hence i) HSA nanocapsules self-assembly without the use of toxic covalent cross-linkers, ii) nanometric sizes below 50 nm and iii) a high DLC of ca. 88%. The efficacy of Dox-HSA nanocapsules was confirmed in multicellular tumor spheroid models (MCTS) of human hepatocarcinoma cells by their significant growth inhibition with respect to controls. In future works, modification of the protein surface of the capsules for cell targeting or to improve colloidal stability and low fouling properties are envisioned. As shown herein, our approach opens a new synthesis route in the field of protein-based nanocarriers for drug delivery applications.

## Conflicts of interest.

There are no conflicts to declare.

## Acknowledgements.

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## Supporting information:

S1.  $\text{N}_2$  adsorption-desorption isotherms; S2. FTIR spectra of RB and RB-NH<sub>2</sub>. And S3. FTIR spectra of RB-Dox-HSA and RB-HSA samples compared to RB.



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# SUPPORTING INFORMATION

## **Mesoporous silica templated-albumin nanoparticles with high doxorubicin payload for drug delivery assessed with a 3-D tumor cell model**

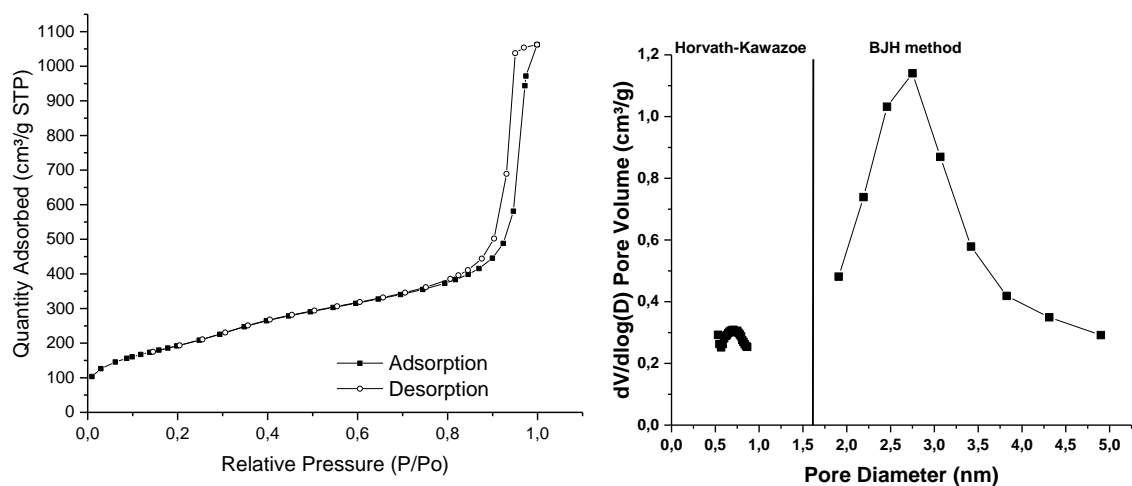
Mathilde Ménard<sup>a,b</sup>, Florent Meyer<sup>\*,b</sup>, Ksenia Parkhomenko<sup>c</sup>, Cédric Leuvrey<sup>a</sup>, Grégory Francius<sup>d</sup>, Sylvie Bégin-Colin<sup>a</sup>, Damien Mertz<sup>\*,a</sup>

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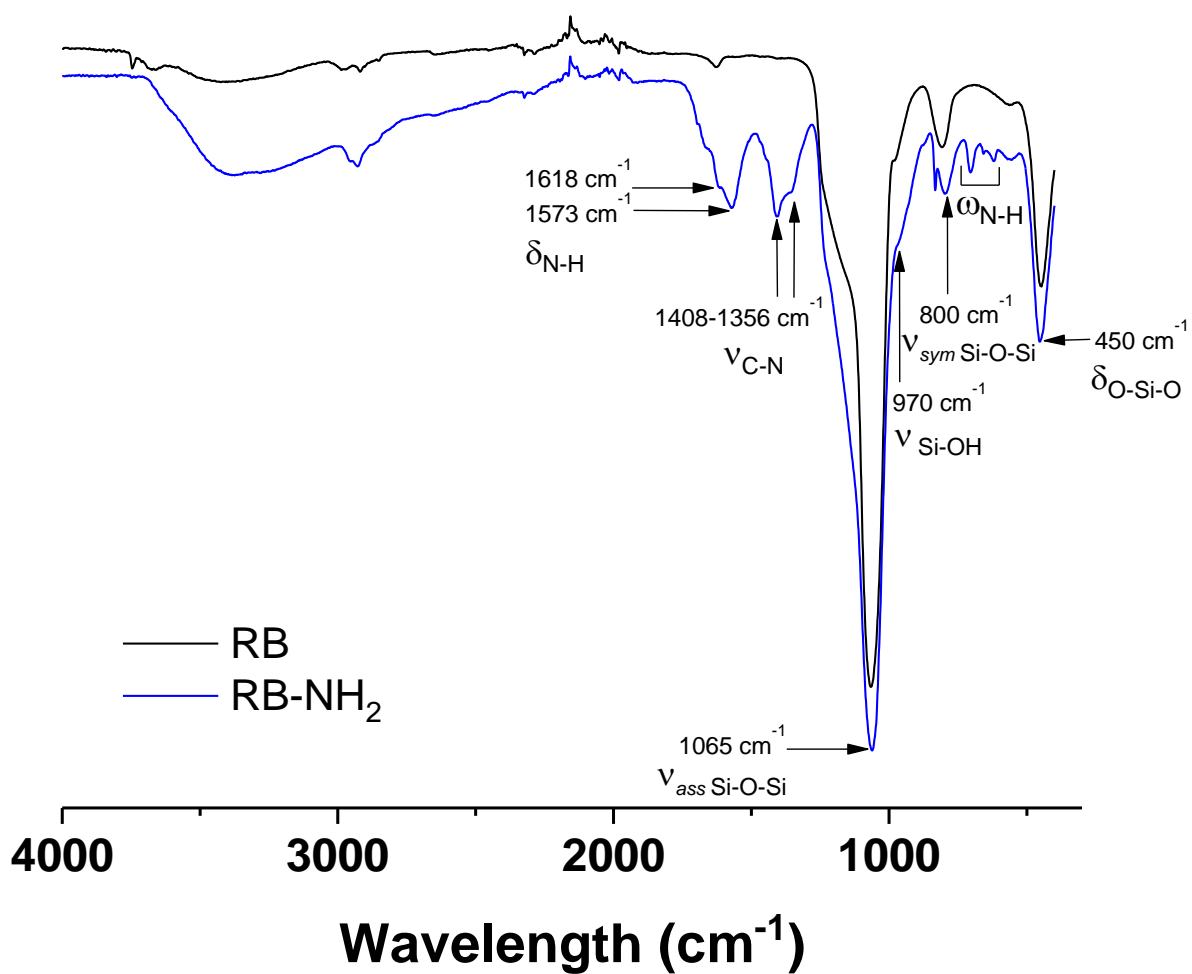
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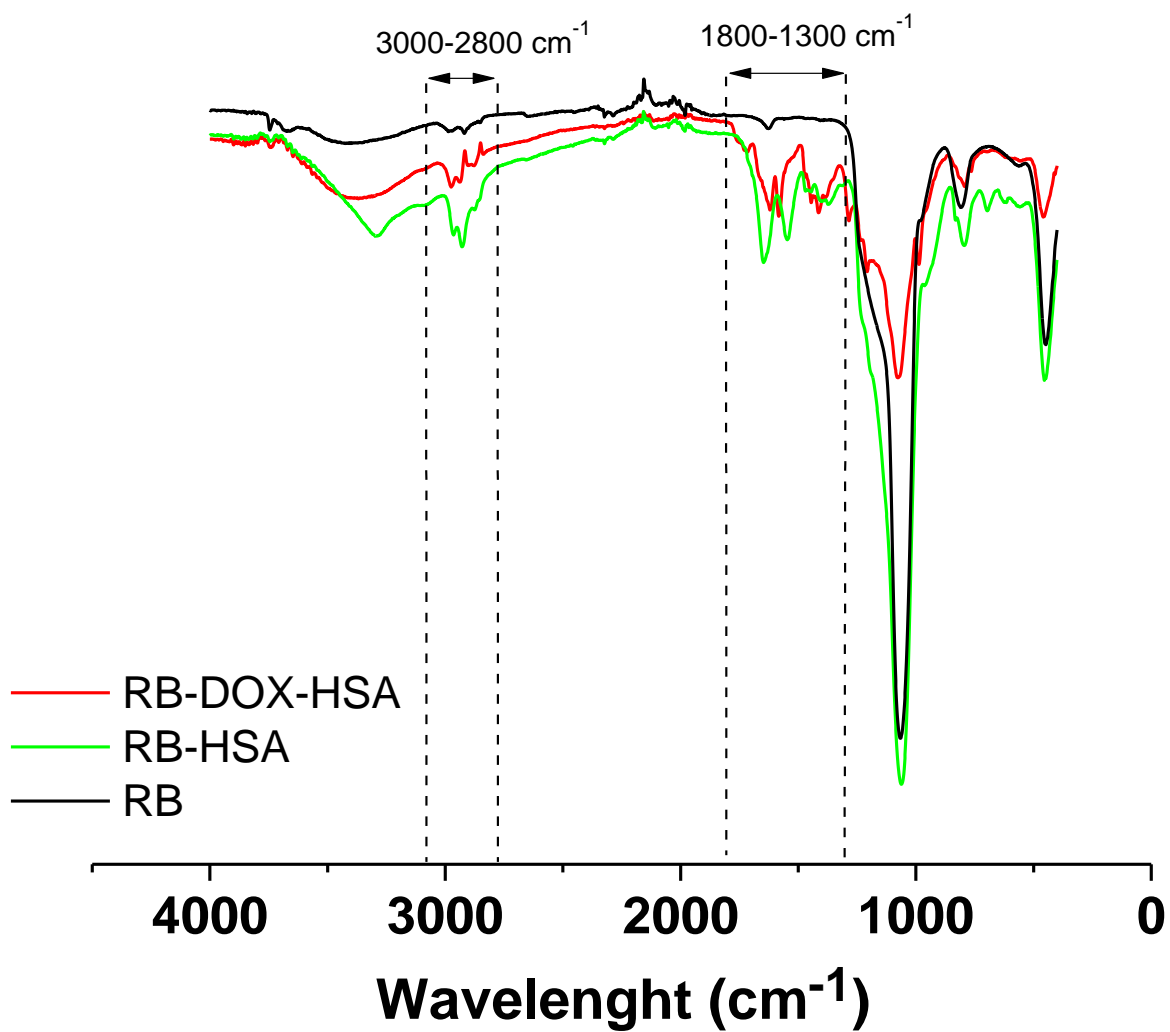
<sup>d</sup>*CNRS - Université de Lorraine, Laboratoire de Chimie Physique et Microbiologie pour l'Environnement, LCPME, UMR 7564, Villers-lès-Nancy, F-54600, France*



**Figure S1.** A) N<sub>2</sub> adsorption-desorption isotherms of RB NPs and B) their corresponding pore size distribution curves in inset calculated by the BJH method for pore sizes above 2 nm and the Horvath-Kawazoe method for pore sizes <1 nm.



**Figure S2.** FTIR spectra of RB and RB-NH<sub>2</sub>.



**Figure S3.** (A) ATR-IR analysis of RB-Dox-HSA and RB-HSA samples compared to RB.