Two-step nanoprecipitation for the production of protein-loaded PLGA nanospheres

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One of the first methods to encapsulate drugs within polymer nanospheres was developed by Fessi and coworkers in 1989 and consisted of one-step nanoprecipitation based on solvent displacement. However, proteins are poorly encapsulated within polymer nanoparticles using this method because of their limited solubility in organic solvents. To overcome this limitation, we developed a two-step nanoprecipitation method and encapsulated various proteins with high efficiency into poly(lactic-co-glycolic)acid (PLGA) nanospheres (NP). In this method, a protein nanoprecipitation step is used first followed by a second polymer nanoprecipitation step. Two model enzymes, lysozyme and α-chymotrypsin, were used for the optimization of the method. We obtained encapsulation efficiencies of >70%, an amount of buffer-insoluble protein aggregates of typically <2%, and a high residual activity of typically >90%. The optimum conditions identified for lysozyme were used to successfully encapsulate cytochrome c(Cyt-c), an apoptosis-initiating basic protein of similar size, to verify reproducibility of the encapsulation procedure. The size of the Cyt-c loaded-PLGA nanospheres was around 300–400 nm indicating the potential of the delivery system to passively target tumors. Cell viability studies, using a human cervical cancer cell line (HeLa), demonstrate excellent biocompatibility of the PLGA nanoparticles. PLGA nanoparticles carrying encapsulated Cyt-c were not efficient in causing apoptosis presumably because PLGA nanoparticles are not efficiently taken up by the cells. Future systems will have to be optimized to ascertain efficient cellular uptake of the nanoparticles by, e.g., surface modification with receptor ligands.

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1. Introduction

Nanoparticles can be used to design or even comprise excellent drug delivery systems [1,2]. For example, due to the enhanced permeability and retention (EPR) effect, nanoparticles can passively target tumors and accumulate in them [1,3]. Nanoparticles can increase the stability of drugs including proteins in blood, are secreted less readily by the kidney, which often results in increased therapeutic efficacy and can reduce side effects of other therapies. [1,3–8]. In this work, we focus on nanoparticles comprised of poly(lactic-co-glycolic acid) (PLGA) because the polymer is an intensely studied material in the field of sustained release, has received FDA approval in various invasive applications including drug delivery, and is biocompatible, biodegradable, and non-toxic [2,9–11].

A promising method to obtain PLGA nanoparticles is by nanoprecipitation, a procedure that was developed by Fessi and coworkers and enables production of particles in the 100–300 nm range [12]. Advantages of this method include that it is a single step not requiring extended shearing/stirring rates, sonification, or high temperatures. The method is characterized by the absence of an oil–aqueous interface which is detrimental to protein structure and function [13,14]. However, the nanoprecipitation method, as developed, is mostly suitable for hydrophobic compounds that are soluble in ethanol or acetone, but display limited solubility in water. For example, Barichello et al. obtained encapsulation efficiencies close to 100% with the lipophilic drugs cindomethacin and cyclosporine A, but less than 15% for the hydrophilic drugs vacomycin and phenobarbital [15].

In order to overcome these limitations, the original nanoprecipitation method was modified by Bilati et al. using a wide range of water-miscible organic solvents [13]. This work provided evidence that nanoprecipitation could occur with solvents other than acetone or ethanol, and that an accurate solvent and non-solvent selection could be extended to enable nanoprecipitation of more hydrophilic drugs. It remains difficult to identify two suitable solvents, because one of them must be able to dissolve both drug and polymer (solvent or diffusing phase), while the polymer should be insoluble in the second solvent (non-solvent or dispersing phase). In a second study,
they selected the water-miscible organic solvent DMSO as the dif-
fusing phase and tested the encapsulation using the model proteins
lysozyme and insulin [16]. The authors were able to load nanoparticles
efficiently with lysozyme, but not with insulin. Note that the study
by Bilati et al. [13] did not include protein stability experiments. This
is troublesome because DMSO is reported to irreversible unfold most
proteins [17,18] and it is therefore unlikely that the developed method
generally applicable.

We set out to overcome the aforementioned problems by develop-
ing a new nanoencapsulation procedure. Overcoming the obstacles
in protein encapsulation by one-step nanoprecipitation is challenging.
First, it is difficult to find a common solvent for the quite hydrophobic
PLGA and the hydrophilic protein. Second, the organic solvent can
induce deleterious protein structural and functional loss. We there-
fore designed a novel two-step nanoprecipitation method (Fig. 1) and
tested its capability to encapsulate two model proteins, lysozyme and
α-chymotrypsin, into PLGA nanoparticles. The two model prote-
ins were chosen because we have employed them frequently in
the past to follow encapsulation procedures [14]. While lysozyme is
quite stable, α-chymotrypsin easily denatures and is an excellent sen-
sor for the potential impact of the procedure on protein structure and
function [14]. The first step in this new method consists in solvent-
induced nanoprecipitation of the protein. Then, encapsulation was
accomplished by a subsequent polymer nanoprecipitation step. In
contrast to Bilati et al. who used DMSO to dissolve the proteins [16],
we suspended the dehydrated protein nanoparticles obtained by sol-
vent precipitation in organic solvents incapable of dissolving proteins,
but capable of dissolving PLGA. Results from solid-state protein for-
mulations show that in the absence of water, protein conformational
mobility is reduced so that the stability of proteins in contact with
the organic solvent is enhanced [14,19,20]. Results from non-aqueous
enzymology support this assumption [14,21–23]. By determining pro-
tein aggregation and function after encapsulation, we tested whether
our assumptions with respect to the advantages of reduced protein
structural mobility were correct or not.

After optimizing the methodology, we employed the processing
parameters established for lysozyme to encapsulate an unrelated ba-
sic protein of similar size, horse heart cytochrome c(Cyt-c), in PLGA
nanospheres to test the potential of the drug delivery system for ap-
lications in cancer treatment [24]. Cyt-c is an important mediator of
apoptosis when it is released from the mitochondria to the cytoplasm.
This process normally takes place in response to DNA damage, but in
many cancer cells it is inhibited. The targeted delivery of Cyt-c directly
to the cytoplasm of cancer cell could selectively initiate apoptosis.

2. Materials and methods

2.1. Chemicals

Poly[(d,l-lactic-co-glycolic)acid (PLGA) with a co-polymer ratio of
50:50 and 65:35 [lactate-to-glycolide] and a MW of 10,000 (not end-
capped), was from Lakeshore Biomaterials (Birmingham, AL). The MW
is an average value determined by the supplier. Bovine pancreatic
α-chymotrypsin, hen egg-white lysozyme, equine heart cytochrome
c(Cyt c), micrococcus cells, and poly(vinyl)alcohol (PVA, 87%–89% hy-
drolized with a MW of 13,000–23,000) were from Sigma-Aldrich (St.
Louis, MO). Acetonitrile (ACN, HPLC grade) was from Fisher Scientific
(Pittsburgh, PA). Succinyl-Ala-Ala-Pro-Phe-p-nitroanilide was from
Bachem Laboratories (Torrens, CA).

2.2. Protein precipitation and encapsulation

Protein nanoparticles were obtained using a similar method as
described by Weber et al. [25]. Briefly, lysozyme and α-chymotrypsin
were solvent-precipitated from 0.8 and 1 ml of aqueous solutions
at concentrations of 25 and 15 mg/ml, respectively, by adding the
water-miscible solvent acetonitrile at a 1:4 volume ratio. The result-
ing protein suspension was stirred for 5 min with a magnetic stir bar.
PLGA was dissolved in acetonitrile at 190 and 28.5 mg/ml and 2 and
10 ml added to the lysozyme and α-chymotrypsin suspensions, re-
spectively. The resulting mixtures (6 and 14 ml) were added directly
through a syringe needle into 240 and 560 ml of a 10% PVA solution
under stirring (60 ml/min) with a magnetic stir bar (5.08 cm length).
The volume ratio of dispersing phase to diffusing phase was 1:40.
Polymer nanoprecipitation was immediately visible upon injection
of the protein suspensions. The PLGA nanoparticles formed were imme-
diately centrifuged for 10 min at 8000 rpm, the supernatant discarded,
and the pellet re-suspended in distilled water. This washing step
was thrice repeated and the samples subsequently freeze-dried by first
rapidly freezing them in liquid nitrogen followed by lyophilization
at a condenser temperature of −45 °C and a pressure of <60 μm of
Hg [26]. Cyt-c encapsulation was performed using the same optimum
conditions established by us for lysozyme since it has a similar size
and net charge.

2.3. Determination of the precipitation yield

After protein nanoprecipitation, the resulting protein suspension
was centrifuged at 5000 rpm for 10 min. The supernatant was dis-
carded and the pellet vacuum dried for 30 min. Protein concentration
and protein aggregates in the pellet were determined as described
by us in detail [26–29]. In brief, the protein pellet was suspended in
2 ml of potassium phosphate buffer for 2 h to dissolve the buffer-
soluble fraction. The samples were then subjected to centrifugation
at 5000 rpm for 5 min and the supernatant used to determine the
concentration of soluble protein. Next, 1 ml of 6 M urea was added to
the pellet to dissolve the buffer-insoluble protein fraction and used to
determine the concentration of aggregated protein by measuring the
UV absorbance at 280 nm. The precipitation yield was calculated from
the actual and theoretical quantity of protein recovered after nano-
precipitation and rehydration. The experiments were performed in
triple, the results averaged, and the standard deviations calcu-
lated.
The activity was obtained by plotting the time-dependent absorbance changes vs. time. The linear portions of the graphs at less than 10% substrate conversion were used to obtain the initial velocities ($V_0$). In all cases the specific activity (mM of substrate converted into product per min and per mg of protein) was calculated. The experiments were performed in triplicate and the results averaged and the standard deviations calculated.

2.7. In vitro release studies

In vitro release studies were conducted as described by us in the past [26–29]. In brief, nanospheres (30 mg) were placed in 1 ml of 10 mM phosphate buffered saline (PBS) at pH 7.3 and incubated at 37 °C. At pre-determined times (typically every 24 h) the supernatant was removed after a short centrifugation. The concentration of the released protein in the supernatant was determined by absorbance measurement at 280 nm (the absorbance was corrected for the very small absorbance produced by degrading empty PLGA nanospheres). The concentration of the released protein was used to construct cumulative release profiles. Release experiments were performed at least in triplicate, the results averaged, and the standard deviations calculated.

2.8. Cell culture

Human HeLa epithelial adenocarcinoma cells (American Type Culture Collection (ATCC), Manassas, VA, catalog number CCL-2) were cultured in Eagle’s Minimum Essential Medium (Invitrogen Corp.) supplemented with 10% (v/v) fetal bovine serum (FBS; Invitrogen Corp.), penicillin (100 U/ml), and 1% glucose as described by the ATCC. Cells were maintained at 37 °C in a humidified atmosphere of 5% CO$_2$ and 95% air.

2.9. In vitro cell studies – non-radioactive cell proliferation assay

The non-radioactive cell proliferation assay was performed according to the manufacturer instructions (Promega). HeLa cells were seeded into 96-well plates at 7.5 × 10^4 cells/well and incubated at 37 °C for 24 h. Then, cells were subjected to medium replacement containing 1% FBS and incubated overnight. Various concentrations of Cyt-c-PLGA NPs dispersed in the cell culture medium were added to cells followed by further incubation at 37 °C for 24, 48, 72, or 96 h. The tested concentrations of Cyt-c-PLGA NPs are equivalent to Cyt-c concentrations of 0.61, 1.21, 3.10, 6.19 and 12.38 μg/ml. Control experiments were performed using blank PLGA NPs. At the day of the experiment, the cells were washed once with PBS and 100 μl fresh medium was added. Background values were recorded at 492 nm using a microplate reader. Then, cells were treated with 20 μL of MTS (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl-tetrazolium; 5 μg/μl) for 1 h and the absorbance was measured at 492 nm. The results were based on at least three independent experiments and the data averaged.

2.10. Statistical analysis

All experiments were performed at least in triplicate, the data averaged, and the standard deviations calculated. The standard deviations are included in all tables as ± values. To establish statistical significance when comparing multiple groups we used one-way multiple Tukey comparison post-test ANOVA. A P value of <0.05 was considered statistically significant.

### Table 1

Properties of the protein precipitates using acetonitrile (ACN) and acetone as desolvent agent.*

| Protein/solvent | Precipitation yield (%) | Insoluble aggregates (%) | Residual activity (%) |
|-----------------|-------------------------|--------------------------|----------------------|
| Lysozyme        |                         |                          |                      |
| ACN             | 79 ± 4                  | 0 ± 0                    | 96 ± 6               |
| Acetone         | 54 ± 28                 | 6 ± 3                    | 81 ± 3               |
| α-Chymotrypsin  |                         |                          |                      |
| ACN             | 80 ± 5                  | 3 ± 2                    | 73 ± 1               |
| Acetone         | 82 ± 3                  | 1 ± 2                    | 75 ± 8               |

* Protein concentration: 10 mg/ml; volume ratio of water-to-organic solvent: 1:4.

2.4. Dynamic light scattering

The size of protein nanoparticles and PLGA nanospheres was determined by dynamic light scattering using a DynaPro Titan with MicroSampler from Wyatt Technology Corporation (Santa Barbara, CA) as described by us in detail [20]. Protein nanoparticles were measured as a suspension in acetonitrile and the PLGA nanospheres as a suspension in water at 100% power intensity. Data analysis was performed using the Dynamic 6.7.6 software supplied with the instrument. The instrument was periodically calibrated using BSA as a standard. In the past, we found that scanning electron microscopy images and size data from dynamic light scattering were consistent [20].

2.5. Determination of actual protein loading and encapsulation efficiency

The actual protein loading of nanospheres was determined following a methodology developed in our laboratory [27]. In brief, 20 mg of PLGA nanospheres were dissolved in 2 ml of ethyl acetate and stirred for 2 h followed by centrifugation at 9000 rpm for 10 min. The supernatant was discarded and the pellet vacuum dried for 30 min. This pellet, consisting mostly of protein, was suspended in 2 ml of potassium phosphate buffer for 2 h to dissolve the buffer-soluble protein fraction. The samples were then subjected to centrifugation at 9000 rpm for 10 min and the supernatant used to determine the concentration of soluble protein. Next, 1 ml of 6 M urea was added to the pellet to dissolve the water-insoluble protein fraction. In all cases, a clear solution without noticeable light scattering was obtained and used to determine the concentration of aggregated protein by measuring the UV absorbance at 280 nm and by BCA assay at 562 nm. The encapsulation efficiency was calculated from the actual and theoretical loading of protein in the nanospheres. The experiments were performed in triplicate, the results averaged, and the standard deviations calculated to highlight the reproducibility of the experiments.

2.6. Determination of enzyme activity

To determine the enzyme activity after encapsulation, ethyl acetate was used to dissolve PLGA because it does not cause enzyme inactivation in the process [27]. Activity of α-chymotrypsin was determined using succinyl-Ala-Ala-Pro-Phe-p-nitroanilide as the substrate [28]. The reaction was carried out in 1 ml of 0.1 M Tris–HCl buffer containing 0.05 mg/ml enzyme, 0.35 mM substrate, and 10 mM CaCl$_2$ at pH 7.8 and our data for α-chymotrypsin as purchased are compared to those reported [28]. The activity of 0.01 mg/ml lysozyme was determined by measuring the decrease in turbidity at 450 nm of a 0.015% (w/v) suspension of Micrococcus lysodeikticus cells in 1 ml of 66 mM potassium phosphate buffer at pH 6.2 and 25 °C as described by us [29]. The peroxidase-like activity of Cyt-c which is not a natural enzyme was obtained as described [30]. Briefly, the reaction was followed at 415 nm using 0.25 ml of 0.01 mg/ml Cyt-c, 0.2 ml of 300 mM H$_2$O$_2$, and 0.55 ml of 0.05 mM 2,2'-azino-bis(3-ethylbenzothiazoline-6-sulphonic acid) in 20 mM potassium phosphate buffer at pH 7. The data obtained by us for commercial Cyt-c are comparable to those reported in the literature [30].
3. Results and discussion

3.1. Development and optimization of a two-step nanoprecipitation method

3.1.1. Protein nanoprecipitation

First we explored the effects of different desolvating agents, different excipients, and the protein concentration on the protein nanoprecipitation process using lysozyme and α-chymotrypsin as model proteins. In order to optimize the processing conditions, the precipitation yield, protein particle size, and residual enzyme activity were determined. While protein precipitates were obtained with acetonitrile and acetone, propanol and ethanol were inefficient for both enzymes (data not shown, for conditions see footnotes in Table 1). We also tested the addition of common stabilizing excipients on the process outcome (poly(ethylene glycol) with a MW of 8000, methyl-β-cyclodextrin, and trehalose at a 1:1 excipient-to-protein weight ratio). It was found that excipients did not improve the process outcome in the case of lysozyme and hindered precipitation in the case of α-chymotrypsin. We therefore did not employ excipients subsequently to avoid such complications—at.

Focusing on the best solvents identified, ACN and acetone, the outcome of the nanoprecipitation process was characterized in detail (Table 1). While the precipitation efficiency was comparable for both solvents, acetonitrile caused less enzyme aggregation and inactivation in the case of lysozyme. This solvent was therefore chosen for all subsequent work.

In order to further optimize the precipitation conditions, we varied the volume ratio of acetonitrile-to-water. Similar to Weber et al. [25] who used ethanol as desolvating agent, we found that a 1:4 water-to-ACN volume ratio was sufficient to precipitate both proteins (data not shown).

Next, we tested the effect of the protein concentration on precipitation results (Table 2). The precipitation yield and particle size increased at increasing protein concentration under otherwise constant precipitation conditions. While for both proteins, no significant amounts of buffer-insoluble aggregates were formed regardless of the protein concentration, the residual activity increased at increasing protein concentration. We interpret this as an indication, that protein molecules close to the solvent-interface are more prone to denaturation than molecules buried in the interior of the precipitates. Such molecules close to the solvent-interface are more prone to denaturation than molecules buried in the interior of the precipitates.

3.1.2. Protein nanoparticle encapsulation

After optimizing the protein precipitation conditions, we proceeded to encapsulate the model proteins into PLGA nanospheres.

Table 2
Characterization of the nano-precipitation results at various protein concentrations at a 1:4 volume ratio of water-to-acetonitrile.

| Protein         | Concentration (mg/ml) | Precipitation yield (%) | Insoluble aggregates (%) | Residual activity (%) | Diameter (nm) |
|-----------------|------------------------|-------------------------|--------------------------|----------------------|---------------|
| Lysozyme        | 10                     | 70 ± 11                 | 2 ± 1                    | 87 ± 5               | 86 ± 16       |
|                 | 20                     | 95 ± 4                  | 1 ± 0                    | 83 ± 1               | 10 ± 15       |
|                 | 30                     | 99 ± 1                  | 1 ± 1                    | 96 ± 2               | 168 ± 14      |
| α-Chymotrypsin  | 10                     | 77 ± 5                  | 0 ± 0                    | 78 ± 0               | 86 ± 16       |
|                 | 20                     | 83 ± 3                  | 0 ± 0                    | 100 ± 1              | 174 ± 16      |
|                 | 30                     | 83 ± 4                  | 0 ± 0                    | 100 ± 2              | 200 ± 12      |

Previously, Giteau et al. precipitated proteins to ensure their stability upon subsequent encapsulation within PLGA microspheres using a solid-in-oil-in-water (s/o/w) technique [19]. After protein precipitation with glycofurol, proteins were centrifuged and the pellet suspended in acetonitrile (ACN) containing the polymer and encapsulated within PLGA microspheres. Our method used the same desolvating agent (ACN) to precipitate the protein and to dissolve the polymer. Additionally, several steps in the encapsulation procedure were changed systematically to assure obtaining nanosized PLGA spheres with high protein loading while aiming at avoiding enzyme inactivation and aggregation. Initially, we selected PLGA with a co-polymer ratio of 65% lactic acid and 35% glycolic acid, a theoretical loading of 2% (w/w), and ACN as the diffusing phase. We tested two commonly used emulsifying agents, namely, poly(vinyl alcohol) (PVA) and poly(ethylene glycol) (PEG, MW = 8000) using a set of defined conditions (Table 3) [28]. It was found that the highest encapsulation efficiency of ca. 70% was achieved using 10% of PVA without compromising protein stability.

We tried to increase the protein loading to 5%, but surprisingly the encapsulation failed when the protein nanoparticles suspended in PLGA solution were added to the PVA solution. However, using PLGA with a co-polymer ratio of 50:50 resulted in nanocapsulation, but the encapsulation efficiency needed improvement. When we increased the volume of the diffusing phase to accomplish faster particle hardening, the encapsulation efficiency increased substantially to >80% at a 1:40 volume ratio of dispersing-to-diffusing phase (Table 4). We also tested the polymer concentration in this context. It has been shown that a higher polymer concentration leads to higher encapsulation efficiency and larger size of the nanoparticles [31,32]. At a high PLGA concentration, the viscosity of the diffusing phase increases which should result in improved encapsulation by reduction of lysozyme nanoparticles leaking into the dispersing phase. Indeed, we found increasing lysozyme encapsulation efficiency at increasing polymer concentration as expected (Table 5). In a similar fashion encapsulation efficiency was improved for α-chymotrypsin. Changing the polymer concentration proved only somewhat successful in this case, possibly because at increased PLGA concentrations the polymer shell thickness also increased [33]. The encapsulation efficiency remained with a maximum of 30% too low for practical purposes (Table 5). Reducing the particle size of α-chymotrypsin by employing a lower protein concentration of 15 mg/ml (Table 2) resulted in an improved encapsulation efficiency of 74% (Table 6).

Table 3
Effect of the emulsifier on selected properties of lysozyme-loaded PLGA nanospheres.*

| Dispersing phase | Encapsulation efficiency (%) | Protein aggregates (%) | Residual activity (%) |
|------------------|-----------------------------|-----------------------|----------------------|
| Water            | 48 ± 23                     | 0 ± 0                 | 100 ± 12             |
| 10% PEG          | -10                         | n.d.                  | n.d.                 |
| 5% PVA           | 58 ± 28                     | 3 ± 5                 | 88 ± 21              |
| 10% PVA          | 71 ± 15                     | 0 ± 5                 | 90 ± 3               |

* Lysozyme concentration: 25 mg/ml; volume ratio of water to ACN: 1:4; concentration of PLGA 65:35 in acetonitrile: 28.5 mg/ml; total volume of the diffusing phase: 12 ml, and for the dispersing phase: 150 ml; theoretical loading: 2% (w/w).

Weber et al. [25] who used ethanol as desolvating agent, we found that a 1:4 water-to-ACN volume ratio was sufficient to precipitate both proteins (data not shown).

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Table 4  
Effect of the ratio of dispersing-to-diffusing phase on the encapsulation efficiency of lysozyme in PLGA nanoparticles.*

| Ratio of dispersing phase to diffusing phase | Encapsulation efficiency (%) |
|---------------------------------------------|-------------------------------|
| 1:10                                        | 9 ± 3                         |
| 1:20                                        | 49 ± 9                        |
| 1:30                                        | 71 ± 7                        |
| 1:40                                        | 84 ± 8                        |

* Protein concentration: 25 mg/mL; volume ratio between water and ACN: 1:4; PLGA 50:50 concentration in ACN: 90 mg/mL; theoretical loading: 5% (w/w).

Table 5  
Effect of different polymer concentrations on the protein encapsulation efficiency in PLGA nanospheres.*

| Batch | Concentration of PLGA 50:50 in ACN (mg/mL) | Encapsulation efficiency (%) | Protein |
|-------|--------------------------------------------|-------------------------------|---------|
|       |                                            |                               | Lysozyme | α-Chymotrypsin |
| 1     | 38                                         | 26 ± 6                        | 11 ± 4   | 24 ± 4         |
| 2     | 63                                         | 45 ± 12                       | 24 ± 4   | 24 ± 4         |
| 3     | 95                                         | 68 ± 7                        | 30 ± 1   | 23 ± 3         |
| 4     | 190                                        | 94 ± 5                        | 23 ± 3   | 23 ± 3         |

* Protein concentration: 25 mg/mL; volume ratio of water to organic solvent: 1:4; polymer mass: 380 mg; volume ratio of dispersing-to-diffusing phase: 1:40; theoretical protein loading: 5%.

Table 6  
Effect of the polymer concentration on α-chymotrypsin encapsulation efficiency in PLGA nanospheres.*

| Batch | Concentration of PLGA 50:50 in ACN (mg/mL) | Encapsulation efficiency (%) | Protein |
|-------|--------------------------------------------|-------------------------------|---------|
|       |                                            |                               | Lysozyme | α-Chymotrypsin |
| 1     | 28.5                                       | 74 ± 4                        | 48 ± 4   | 48 ± 4         |
| 2     | 47.5                                       | 49 ± 9                        | 48 ± 4   | 48 ± 4         |
| 3     | 71.25                                      | 48 ± 4                        | 48 ± 4   | 48 ± 4         |
| 4     | 142.5                                      | 38 ± 18                       | 48 ± 4   | 48 ± 4         |

* Protein concentration: 15 mg/mL; volume ratio between water and organic solvent: 1:4; polymer mass: 285 mg; volume ratio of dispersing-to-diffusing phase: 1:40; theoretical protein loading: 5%.

The data show how sensitive the results respond to encapsulation conditions in this method highlighting the fact that encapsulation likely has to be optimized in a similar fashion as described here for other proteins. However, there are only a few processing parameters requiring adjustment and the process is straightforward and reproducible as demonstrated by the small standard deviations obtained for encapsulation parameters under optimized conditions.

The optimum conditions to encapsulate lysozyme and α-chymotrypsin in PLGA nanoparticles are summarized in Table 7. The size of the protein loaded PLGA particles obtained by dynamic light scattering was ca. 300–400 nm in diameter (Table 7). However, while lysozyme encapsulation afforded a highly active enzyme, substantial enzyme inactivation and formation of buffer-insoluble aggregates were observed for α-chymotrypsin. The formation of buffer-insoluble aggregates and loss in specific activity found for α-chymotrypsin is similar to results obtained before upon α-chymotrypsin encapsulation in PLGA microspheres using a s/o/w technique [27,28,34–36]. The use of stabilizing additives (e.g., methyl-β-cyclodextrin or poly(ethylene glycol)) was necessary in the latter case to preserve protein integrity. Such strategies have to be developed for the new two-step nanoprecipitation procedure as well.

3.1.3. Cytochrome c-loaded PLGA nanoparticles

Having accomplished protein loaded nano-sized PLGA particles, we tested the development of the sustained release nanoparticles into an application platform. We selected Cyt-c as model protein because it has been employed in experiments geared towards better cancer treatment options [24]. The size of our particles makes them potentially useful in passive and also active targeting of cancer tissues [37,38]. For example, Santra et al. [24] demonstrated recently the therapeutic potential of Cyt-c in nanoparticles by their capability to induce apoptosis in lung carcinoma cells after uptake by the cells by endocytosis. However, their vehicle consisted of a water-soluble hyperbranched polyhydroxyl polymer not approved in medical applications. In contrast, our nanoparticles employ an already FDA approved and commercially available polymer (PLGA) and a straightforward encapsulation method.

We hypothesized that encapsulation of Cyt-c via the two-step nanoprecipitation method should work using the optimum conditions identified for lysozyme (Table 7) because both proteins have a similar molecular weight (12 and 14 kDa, respectively) and are basic [39]. The encapsulation efficiency for Cyt-c was with 72% similar to that obtained for lysozyme under identical conditions (Table 8). The peroxidase activity of Cyt-c was comparable to values prior to precipitation and encapsulation and only few aggregates were formed indicating good preservation of structural integrity during the process. The size of the particles obtained was 340 nm and thus potentially useful to enable passive delivery to cancer tissues based on the EPR effect [37,38].

In vitro release of Cyt-c from the PLGA nanoparticles showed an initial “burst” release within 24 h that was reasonably small with ca. 20% (Fig. 2). Burst release values of >20% are frequently found for such systems, in particular when nanosized systems are being used [40]. During a 100-day incubation period, Cyt c was released completely from the nanospheres. Since the release was slow, the amount of protein released per day was small and the residual activity during release could not be measured with accuracy. Future experiments using cell cultures and animal models will shed light into the bioactivity of the developed system. However, since 100% of the protein was released, we can exclude the formation of buffer-insoluble Cyt-c during the release period.

3.2. Cyt-c-PLGA nanoparticle cytotoxic effects in cancer cells

Since there are some reports that PLGA nanoparticles could be internalized by cells, we investigated whether the Cyt-c-PLGA NPs...
nanoparticles is by one-step nanoprecipitation. Establishment of biocompatibility of the PLGA nanospheres obtained is important in the development of them as drug delivery systems. It has to be pointed out, however, that the effect of the drug-loaded delivery system on cell viability was too low to be clinically relevant. It is likely that PLGA NPs are not taken up effectively and thus Cyt-c is not being effectively delivered to the cell cytoplasm in agreement with recent data [41]. We can conclude from this that PLGA NPs have to be either modified with a homing ligand or release a drug coupled to a homing ligand to enable uptake by receptor-mediated endocytosis. We are currently working on transforming this system in this direction.

4. Conclusions

Nanosized delivery systems hold promise in improving protein delivery, i.e., to target tumors and inflammation. A convenient method to accomplish nanosized polymer particles is by one-step nanoprecipitation. However, encapsulation of proteins into PLGA nanospheres by nanoprecipitation was inefficient prior to our work and/or involved the solvent DMSO which irreversibly denatures most proteins [17,18,42,43]. To overcome these problems, we developed a two-step nanoprecipitation method to allow for efficient protein encapsulation into PLGA nanospheres without causing irreversible functional changes. Cell viability studies using HeLa cells demonstrate excellent biocompatibility of the PLGA nanospheres obtained. Furthermore, we demonstrate reproducible encapsulation of the model proteins lysozyme, α-chymotrypsin, and Cyt-c into PLGA nanospheres. Optimization of the processing parameters involved in the new two-step nanoprecipitation method enabled obtaining high encapsulation efficiencies. While encapsulation of lysozyme and Cyt-c via the two-step nanoprecipitation method did not lead to the formation of insoluble aggregates or activity loss, significant enzyme inactivation and formation of buffer-insoluble aggregates were observed for α-chymotrypsin. Future studies in our laboratory will be directed towards minimizing this problem.

Admittedly, as one reviewer pointed out to us, the results obtained with the therapeutic protein seem not sufficient to justify the preparative efforts. However, we feel that our work and the results obtained constitute a first significant step into the direction of solving a complex problem. Our work clearly demonstrates the feasibility of obtaining nanosized biocompatible protein delivery systems with good yield and reasonable protein stability. This should support approaches aiming at targeted protein delivery using the enhanced permeability and retention (EPR) effect to deliver pharmaceutical proteins to tumors or inflammation sites. This approach has to be augmented by targeted delivery strategies aimed at enabling endocytosis of the nanoparticles, e.g., by attaching folate to their surface.

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