MAPPING MICROCLIMATE PH IN BIODEGRADABLE POLYMERIC MICROSPHERES

By

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DEDICATION

To my father Jie Liu, mother Guihua Guo, brother Zhibin Liu

To my husband Xiaofeng Wang

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ABSTRACT

The microclimate inside microspheres prepared from biodegradable polymers (e.g., poly(lactic-co-glycolic acid) PLGA) often becomes acidic owing to the accumulation of water-soluble polymer degradation products, which can induce the destablization of encapsulated therapeutic agents. The objective of this dissertation was to quantitatively evaluate the microclimate pH (µpH) inside biodegradable polymeric microspheres in order to facilitate the development of microsphere formulations that control µpH and stabilize acid-labile drugs. Chapter 1 presents an overview of the background of these studies with a focus on the most common biodegradable polymer, PLGA. In Chapter 2, the upH distribution inside proteinencapsulated PLGA microspheres was accurately quantified using a ratiometric method based on confocal laser scanning microscopy (CLSM). The fluorescent response of Lysosensor yellow/blue® dextran used to map acidic µpH in PLGA was influenced by the presence of encapsulated protein. A method for correction of the interference of protein was developed and validated. The upH kinetics in four different PLGA microsphere formulations during incubation under physiological conditions were determined to be roughly pH 4 to neutral pH depending on the formulation. Based on previous literature findings of enhanced stability of

encapsulated proteins and peptides in hydrophilic and biodegradable poly(lactic-cohydroxymethyl glycolic acid) (PLHMGA) microspheres, the µpH distribution and kinetics in the microspheres prepared from PLHMGAs were evaluated in Chapter 3 by CLSM and compared that with their PLGA counterparts. The PLHMGA microspheres developed a far more neutral upH than PLGA, which was linked to more rapid diffusion of acidic degradation products out of the polymer. In the last chapter, a mathematical model was developed to simulate the µpH kinetics and spatial distribution inside degrading PLGA microspheres by considering the acid production, mass transfer via diffusion and partition of water-soluble acids that contribute to the development of upH. Fundamental parameters in the PLGA microspheres were determined from experiments to validate the model. The model successfully predicted the kinetics of upH development, whereas showing a small difference in distribution compared to experimental results. Hence, these mechanistic approaches may provide valuable experimental and theoretical tools to control upH inside the most commonly used biodegradable polymer for controlled release of acid-labile therapeutics.

CHAPTER 1

Introduction

1.1 Biodegradable polymeric systems for controlled release proteins

Therapeutic proteins currently represent a promising and fast growing market globally. Sales for therapeutic proteins are increasing and their market share is projected to reach around 30% of the industrial drug pipeline by 2017 [1]. However, the successful development of protein therapeutics is still facing tremendous formulation challenges. Non-invasive routes of administration (e.g., oral, nasal, pulmonary) often result in poor absorption and low bioavailability. Therefore, routine parenteral administration with a syringe and needle remains the most common method to deliver therapeutic proteins [2]. However, frequent injections are often needed due to the short *in vivo* half-lives of most proteins, leading to poor patient compliance in addition to causing inconvenience and psychological stress for patients. Hence, there is substantial need to develop new and better methods for successful delivery of therapeutic proteins.

Among these methods, injectable biodegradable polymers that can provide slow and continuous protein release over duration of weeks to months have attracted tremendous research interest. Because of this sustained-release approcach, the injection frequency of therapeutic proteins can be sharply reduced, thus improving patient comfort and compliance. They can also protect proteins from *in vivo* degradation, and reduce toxicity by controlling the drug's blood concentration [3]. The local sustained-release of growth factors and other therapeutic proteins also has demonstrated the potential for clinical use in tissue engineering, where local release of proteins at appropriate times promotes tissue regeneration [4-6]. Moreover, application of controlled-release polymers to antigen delivery presents great promises for their ability of enhancing immune response as well as reducing the number of vaccinations [7, 8].

Various drug-carrying devices formulated from biodegradable polymers have been reported, including: microspheres, microcapsules, nanoparticles, pellets, implant, and films. Among these configurations, polymeric microspheres are the most common dosage form for prolonged delivery of proteins and peptides. They are usually administered as depot via subcutaneous or intramuscular injection, with the size in the range of 1 to 250 μ m (ideally less than 125 μ m) [9]. Specific release profiles of therapeutic drugs may be generated by manipulating the properties of the polymer and microspheres such as polymer composition, molecular weight, microsphere size, porosity, etc. Commercially available examples are the products of

human growth hormone (Nutropin Depot®, Genetech), leuprolide acetate (Lupron Depot®, Takeda Chemical), and octreotide acetate (Sandostatin LAR Depot®, Novartis).

1.2 Poly(lactic-co-glycolic acid) (PLGA)

Following the entry of bioresorbable surgical sutures into the market, poly(lactic-co-glycolic acid) (PLGA) emerged as the most investigated biodegradable polymer during the past few decades for the purpose of sustained delivery of a variety of drug classes such as vaccines, peptides, proteins, nucleic acids, etc. [8, 10-15]. They degrade by bulk erosion via hydrolysis in physiological environment to non-toxic products that are eventually safely eliminated by the body. Because of their favorable biocompatibility, biodegradability and low immunogenicity; they are one of only a few polymers widely used in US Food and Drug Administration (FDA) approved pharmaceutical products and medical devices.

1.2.1 Physico-chemical properties of PLGA

PLGA copolymer is composed of monomers of lactic acid and glycolic acid, whose structures are shown as follows:

Both lactic acid and glycolic acid are α -hydroxy acids with a pK_a at 25°C of 3.86 and 3.83, respectively. Since lactic acid contains an asymmetric carbon atom, it has two optical isomers: the L- form and D- form. Therefore, poly (lactic acid) (PLA) exists in two optically active stereoregular forms: poly (D-lactic acid) (P(D)LA) and poly (L-lactic acid)(P(L)LA) and an optically inactive racemic form poly (D,L-lactic acid) (P(D,L)LA). PLGA generally stands for poly (D,L-lactic acid-*co*-glycolic acid) where the D- and L- lactic acid forms are in equal ratio.

PLGA polymers can be synthesized either by direct polymerization from monomers or by ring-opening polymerization from its cyclic dimers, i.e., lactide and glycolide in the presence of catalyst. Due to the difficulty of removing water that degrades the polymer, the first method is limited to moderate molecular weights [16]. Therefore, ring-opening polymerization is typically employed for obtaining high molecular weight polymers. The end groups of polymer chains are either free carboxylic acid or terminated with an aliphatic alcohol via an ester linkage.

The physico-chemical properties of PLGA polymers, such as crystallinity, hydrophobicity, solubility in organic solvents, molecular weight and polydispersity play crucial roles in determining polymer water uptake, hydrolysis, drug release and

hence the performance of drug delivery systems manufactured from PLGA. These PLGA properties can be manipulated via a number of variables, such as monomer stereochemistry, monomer ratio, molecular weight, and end-group chemistry. Poly (glycolic acid) (PGA) is highly crystalline due to the lack of any methyl side groups. It is only soluble in the most aggressive solvents (e.g., hexafluroisopropanol), thereby limiting its use. P(L)LA and P(D)LA are semicrystalline in nature and can be soluble in methylene chloride but not in ethyl acetate or acetone. By contrast, P(D,L)LA and PLGA are amorphous and can dissolve in a wide range of common solvents chloroform including methylene chloride. acetone. ethvl acetate. tetrahydrofuron, hence increasing the possible conditions of drug encapsulation [17]. Due to the introduction of methyl side groups, PLA is more hydrophobic than PGA, so PLGA copolymers rich in lactide content are more hydrophobic. PLGAs with free carboxylic acid end groups are more hydrophilic than those with capped endgroups. Commercially available PLGA polymers are commonly characterized in terms of intrinsic viscosity, as a direct indicator of molecular weight. The glass transition temperature (Tg) of PLGA polymers are in the range of 40-60°C, rendering their glassy state above the physiological temperature of 37 °C. Therefore, they possess significant mechanical strength to be formulated into delivery devices of various size and geometry.

1.2.2 PLGA degradation and erosion

It is well accepted that aliphatic polyesters like PLGA degrade via hydrolysis of their ester linkage *in vivo*, producing both a primary alcohol and a carboxylic acid [18]. The polymer degradation is catalyzed by the carboxyl end groups initially present or produced upon ester bond cleavage, a phenomenon known as autocatalysis effect [19]. The enzymatic activity is suggested not likely to be involved in PLGA biodegradation [20]. PLGA devices degrade by bulk erosion mechanism, meaning the degradation occurs throughout the system. It is often characterized by a continuous decline in polymer molecular weight and lagging mass loss of the matrices due to the dissolution and diffusion of degradation products [21]. A number of factors can affect the rate of PLGA hydrolysis, and therefore can be modulated to adjust the rate of drug release from PLGA matrix.

One important factor determining degradation rate is the polymer composition [22]. Increasing the ratio of lactic acid to glycolic acid in PLGA decreases the rate of degradation due to the higher hydrophobicity of lactic acid relative to glycolic acid, which reduces the water uptake by polymer. Moreover, the introduction of additional methyl group on the lactic acid monomer hinders the water attack of ester bonds sterically. The morphology state (i.e. amorphous vs. semi-crystalline) of polymers also plays a critical role in PLGA degradation. It is well established that the amorphous regions in polymers are preferentially degraded due to the accessibility of water penetration [19]. Compared to semi-crystalline poly (L-

lactide, the degradation time of amorphous poly (D,L-lactide) is much shorter [23]. Lower molecular weight polymers usually relate to faster degradation due to the greater easiness of change in polymer from glassy to rubbery state [24]. The end group chemistry of PLGA also affects its degradation. PLGA with free carboxylic acid groups shows a more rapid degradation than that of end-capped PLGA, owning to the increased hydrophilicity (thus higher water uptake) and increased autocatalysis of ester bond hydrolysis by carboxylic acids [25]. In addition to the afore-mentioned factors; polymer molecular weight distribution, the presence of additives, device dimensions, glass transition temperature, the site of implant, etc., can all influence PLGA hydrolytic behavior [20].

There are two mechanisms responsible for chain scission of polyesters: random chain scission and chain-end scission [26]. Since the microclimate pH inside PLGA matrix is determined by the concentration of total water-soluble acids (see section 1.5.1), chain-end scission rate, which liberates one monomer per chain cleavage is more important in terms of governing microclimate pH than random chain scission rate, which does not necessarily yield a water-soluble acid for each chain cleavage.

1.2.3 PLGA water-uptake kinetics

Polymer hydration represents a fundamental step of initiating and sustaining the hydrolytic degradation and erosion of polymer matrices. After PLGA microspheres are injected into the body or placed in incubation media, water will rapidly penetrate the polymer by simple diffusion. The water absorbed by PLGA microspheres can be divided in two parts, the water associated with the polymer phase (bound water) and the water filling up the pores in the polymer matrix (bulk water). The content of bound water is related to the hydrophilicity of the polymer, which is dependent on polymer properties such as molecular weight, polydispersity, end-group capping and glycolide/lactide ratio, etc. For PLGA RG503H (M_w 30,000 Da), it is reported to be less than 3% w/w of bound water during the first 21 days of degradation. Being in the same phase as the polymer, the bound water acts as an effective plasticizer, causing the decrease of glass transition temperature (Tg) of the polymer [27]. Such plasticization effect increases polymer flexibility or mobility, facilitating the transport of incorporated drug or polymer degradation products. It is also suggested that the bound water results in similar polymer degradation as in bulk water [27]. As PLGA erodes, the pores enlarge and coalesce with each other, leading to increased water content. Moreover, the osmotic pressure created by the dissolved encapsulated drug and/or excipents as well as the accumulation of watersoluble polymer degradation products leads to additional water uptake. At late stages of erosion, with the mass loss and changes in the physical state of the

polymer, the water pores would be closed or reduced in total volume so that the water uptake may be ceased or even declined [28]

1.3 Microspheres preparation

Several methods have been reported and developed for preparation of polymeric microspheres to date. The fabrication method can greatly influence the characteristics of the microspheres, such as size, morphology, drug loading, drug stability (particularly protein therapeutics) and drug release. Thus the choice of preparation method for polymeric microspheres should be wisely made based on the desired properties of microsphere formulation products. Ideally, the microspheres should be manufactured in a way to achieve optimal size and drug loading, high encapsulation efficiency, preserved stability of encapsulated drug, high yield of microspheres, batch uniformity and reproducibility, and free flowing property of microspheres [29]. Furthermore, it is desirable that the produced polymeric microspheres are capable of sustained-release of encapsulated drugs for a specified period with low initial burst release and minimal incomplete release effect. The most widely used manufacturing techniques for polymeric microspheres loaded with protein are: i) spray drying; ii) phase separation (coacervation); and iii) emulsion solvent evaporation method.

i) Spray drying

In this process, the drug in solid form is dispersed in a polymer solution of a volatile organic solvent, such as dimethylene chloride or acetone. The suspension is then sprayed into a chamber supplied with heated air stream. As the organic solvent evaporates instantaneously, the polymer solidifies around the drug forming microspheres. The typical size of microspheres prepared from this method ranges from 1 to 100 µm, which is dependent on the atomizing conditions that are related to temperature, feed flow rate, etc [29]. The final microspheres are collected by a cyclone separator. This method is very rapid, convenient, easy to scale-up and provides good reproducibility. The particle size can be well controlled within a narrow range and the encapsulation efficiency of drug is high [30-32]. Considerations on the use of this method should include the possible significant loss of products during the process due to the adhesion of the microspheres to the inside walls of the apparatus [33]. In addition, the encapsulated protein must be relatively stable against high temperatures and lyophilization processes before dispersion.

ii) Phase-separation (coacervation)

In principle, protein in solid form or in aqueous solution is first dispersed in a polymer solution of organic solvent. Then by mixing with another organic nonsolvent, the polymer solubility in its solvent is gradually decreased, leading to phase separation. Consequently, the polymer rich liquid phase (coacervate) encapsulates the drug and the formed droplets are then transferred to a larger

volume of second organic nonsolvent to harden the microparticles [34-37]. DCM, acetonitrile, and ethyl acetate are typically used to dissolve the polymer. The nonsolvent added should be miscible with the polymer solvent while not dissolving the polymer or the drug. Examples include silicone oil, vegetable oils, and light liquid paraffin. Aliphatic hydrocarbons like hexane, heptane, and petroleum ether are usually employed as the second nonsolvent [38]. In the process, factors including polymer properties, polymer concentration, the stirring rate, the addition rate and viscosity of first nonsolvent, the ratio between polymer solvent and first nonsolvent can affect the coacervation process and thereby the characteristics of final microsphere products [39]. This anhydrous method can circumvent the problem of protein partitioning into the aqueous continuous phase, thereby increasing the protein's encapsulation efficiency. However, residual organic solvent is a major concern for this method and particle agglomeration can be a frequent problem due to the lack of emulsion stabilizer [38].

iii) Emulsion solvent evaporation

The water/oil/water (w/o/w) double emulsion method is most widely employed for preparation of microspheres containing proteins and peptides. Briefly, protein is dissolved in a buffered or plain aqueous medium and mixed with a polymer solution in organic solvent (usually methylene choloride) under vigorous stirring using a homogenizer or sonicator, thus creating the primary w/o emulsion

droplets. Then, a second aqueous phase containing a surfactant (e.g. poly(vinyl alcohol), PVA) is gently added to the first emulsion followed by intensive mixing to generate the second w/o/w emulsion. The organic solvent is removed by transferring the formed emulsion to a large volume of water bath (with or without surfactant), into which the organic solvent is diffused out and eventually evaporated under constant stirring. Finally, the hardened microspheres are collected through filtration or centrifugation, washed with water, sieved for size and lyophilized to the final product. The properties of final microspheres in terms of size, porosity, drug encapsulation efficiency and release depend on a number of formulation variables including polymer type, polymer concentration, volume ratio of internal aqueous to organic phase, homogenization speed, concentration of surfactant, solvent removal rate, the incorporation of salts, etc [40-45]. This double emulsion method is appropriate for numerous proteins, however, one issue often associated with this process is the presence of various stresses on protein stability such as high shear/cavitation force, water/organic interface, and elevated temperatures.

1.4 Instability of protein in PLGA delivery systems

Proteins are relatively large molecules with labile bonds, reactive side chains and sophisticated secondary, tertiary and even quaternary structures. And they have to maintain their specific, folded, three dimensional structures in order to deliver proper function. However, unlike most small molecule drugs, proteins easily

lose chemical and structural integrity, resulting in loss of bioactivity and/or increase of immunogenicity. Protein stabilization is, therefore, regarded as a principle difficulty hindering successful development of PLGA based formulations for sustained-release of therapeutic proteins [14].

1.4.1 Instability mechanisms

Generally, protein instability mechanisms can be divided into two classes: chemical instability and physical instability [46]. The chemical instability involves the covalent modification of amino acid groups on peptide chains. The reaction includes hydrolysis, deamidation, racemization, oxidation, disulfide interchange and β-elimination. Physical instability of protein refers to the disruption of proteins' higher order (e.g. secondary, tertiary or quaternary) structures, including denaturation (protein unfolding), aggregation, precipitation and adsorption to surfaces. Detailed reviews on the protein degradation pathways can be found in other literature [25, 47-49]

1.4.2 Stresses for protein instability

Proteins are exposed to a variety of damaging stresses over the entire life of PLGA delivery systems, which can occur through different stages including: encapsulation, lyophilization and storage, and long-term release [14, 50-52].

During encapsulation

During the loading process with emulsion method, proteins in an aqueous solution are usually emulsified in a polymer organic solution. The presence of water/organic solvent interfaces is a major cause of protein denaturation and aggregation during the process [53-55]. Protein molecules tend to adsorb to the interfaces, leading to unfolding with the exposure of their hydrophobic core to the organic solvent and subsequently aggregation. Moreover, the high shear forces generated by the emulsification process (e.g. homogenization, vortex) is another common destabilizing factor [56]. Emulsification of protein solution by sonication create cavitation stress that is detrimental to proteins because of local temperature extremes and free radical formation [51]. Adsorption to solid surfaces, including containers, solid excipients and homogenizer components, also play a detrimental role in protein stability.

Microspheres prepared from anhydrous conditions (e.g. solid/oil/oil (s/o/o) method) often exhibit better stability of encapsulated proteins than those involved an aqueous medium [14], due to the higher stability of protein in solid state and the elimination of the water/organic solvent interface. However, in order to obtain the anhydrous powder, proteins usually undergo freeze-drying or spray drying first if a reduced particle size is required, which can also potentially damage protein integrity.

During lyophilization and storage

To remove the residual solvent in microspheres left from the preparation process, lyophilization (freeze-drying) is most often used. However, proteins are susceptible to a variety of stresses during the freezing and subsequent drying process, leading to degradation (mainly by denaturation and aggregation) [57]. For example, increased protein concentration and ionic strength, pH changes, and formation of ice crystals during the freezing step can have deleterious effects on maintaining a protein's native state. In addition, the removal of the hydration shell of a protein during the drying process can facilitate the protein-protein hydrophobic interactions, causing protein aggregation [58].

During storage, proteins are not necessarily stable in the solid state [58-60]. The moisture level plays an important role in protein degradation. Moisture can induce premature polymer hydrolysis, which will produce acidic monomers and oligomers, lowering the microenvironment pH and potentially degrading the protein. Moreover, moisture can also cause aggregation by providing a medium for thiol-disulfide exchange [61] or induce formalinized antigen aggregation through formaldehyde-mediated cross-linking [62]. Additionally, temperature and interaction with excipients or PLGA can also lead to protein aggregation or chemical degradation reactions.

During release

Generally, three principle stresses present in the microenvironment of PLGA matrix are recognized to be responsible for protein instability during prolonged *in vivo* release from PLGA matrix: moisture, acidic pH, and adsorption to PLGA surface [50].

When PLGA microspheres are injected into the body, water will penetrate into the polymer matrix and dissolve the encapsulated protein rapidly. Generally, it is well accepted that proteins are most stable in their solid state [14, 50]. The rehydration of protein will mobilize the protein and enhance its reactivity significantly, resulting in destabilization.

Another important detrimental factor for protein stability is the microclimate pH inside aqueous pores of the PLGA matrix. The presence of acid impurities (often monomers and dimers of glycolic acid and lactic acid) plus the degrading products of PLGA containing carboxylic acids create an acidic microenvironment, which could be deleterious to acid labile proteins. Acid-induced instability mechanisms for proteins include acid-catalyzed peptide bond hydrolysis, deamidation, aggregation, and denaturation [48, 63]. For example, simulations of BSA in a very acidic microclimate pH (pH=2) showed denaturation, peptide bond hydrolysis, and noncovalent aggregation [64]. Evidence for acidification within degrading PLGA microparticles has become increasingly notable recently, and will be discussed in detail in section 1.5.

An additional cause of protein destabilization involves polymer surfaces. Protein's hydrophobic interior often has tendency to interact with the hydrophobic polymer chain, resulting in adsorption and subsequently irreversible conformational changes of protein [65]. Since there is large internal surface area inside PLGA matrix due to its porous structure after degradation, the interaction between polymer and protein can induce protein inactivation and incomplete release.

1.5 Acidic Microenvironment in PLGA Microspheres

1.5.1 Physical-chemical description of microclimate pH (µpH)

The microclimate pH (µpH) refers to the pH in the aqueous pores inside the PLGA delivery matrix. Ding *et al.*[28] described the physical-chemical basis of the development of µpH and developed an equilibrium model to quantitatively predict the µpH in PLGA films for the first time. Briefly, upon immersing the PLGA carriers in a physiological buffer, water will be imbibed into the polymer matrix rapidly and fill up the pores that are generated during the preparation process. During the bioerosion period, the pores will close, open, grow in size and coalesce with the degradation of polymer. Two separate phases are assumed to coexist within the polymer matrix, the polymer phase and the aqueous phase. In the polymer phase, the degradation of the polymer occurs after the brief hydration, producing acid

monomers and oligomers that could be either water-soluble or water-insoluble depending on the chain length of the acid. The water-soluble ones can be released out by diffusion through the polymer matrix and partitioning into the aqueous pores, where dissociation takes place, producing protons that lower the μpH . Besides the PLGA degradation products, the acidic impurities in the polymer that are left from polymer synthesis and storage could also contribute to the development of acidic μpH . In summary, the acidic microclimate stems from the accumulation of water-soluble acids in PLGAs.

1.5.2 Factors affecting µpH

From the above discussed model, it can be concluded that a number of factors from the physical-chemical processes could contribute to the development of acidic microclimate, namely the water-soluble acid production rate, the water-soluble acid liberation rate, acids partition between polymer phase and aqueous phase, and the dissociation of water-soluble acids in the aqueous medium. Any formulation and processing variables that affect these factors could directly or indirectly impact the $\mu p H$ distribution and kinetics in PLGAs. Therefore, it is crucial to recognize the effect of different variables on $\mu p H$ in order to wisely design and develop the PLGA delivery systems with controlled $\mu p H$ for the pH-sensitive therapeutic substances.

Ding *et al.* reported that μpH inside PLGA microspheres is dependent upon the molecular weight (MW) and the lactic/glycolic acid ratio of PLGA [66]. Lower μpH was observed in microspheres made from higher MW PLGA after 2 weeks incubation in phosphate buffer, which was explained by the reduction in polymer permeability to water-soluble acids as the molecular weight increases. Increasing the lactic acid content in PLGA, the μpH became less acidic during the first two weeks incubation. This was attributed to the slower degradation rate for higher lactic acid content polymers. However, the μpH developed in PLGA 85/15 and 100/0 microspheres were more acidic after two weeks than that in PLGA 50/50 formulations because the lower permeability to water-soluble acids in the high lactic acid-content polymers impedes the liberation of pH-lowering acids.

The size of the microspheres also plays an important role in µpH kinetics. Li et al. [67] reported that the larger the microspheres, the lower pH they displayed due to the longer diffusion path for water-soluble acids to be released out. However, Ding et al. [66] suggested that the effect of microsphere size on µpH distribution is dependent on polymer materials. For lactic-rich polymers (e.g. PLGA 85/15 and PLA), the size effects become insignificant because of both the low production rate and low diffusion rate of water-soluble acids in such polymers.

Porosity of the microspheres is likely to be of great significance in determining the acidity in PLGAs. It is anticipated that the more porous the inner structure of particles, the more rapid the produced acids are transported out of the

polymer, since the diffusivity of water-soluble acids in water is many orders of magnitude higher than that in the polymer. Furthermore, the buffer salts in the incubation medium are more likely to diffuse into the aqueous pores, thereby helping to neutralize the $\mu p H$.

The method of microsphere preparation could influence the µpH kinetics as well. It is found that microspheres fabricated by the oil-in-oil emulsion method have a lower acidic microenvironment than those made by water-oil-water double emulsion method [66]. Possible reasons speculated were that the oil-in-oil microspheres have higher porosity and amount of residual solvent, both of which could lead to the facilitated release of water-soluble acids from the polymer matrix.

The protein encapsulated can also contribute to the pH of PLGA microenvironment. The side chains on the amino acids of protein molecules can participate in the acid-base equilibrium in the aqueous phase and act as a buffer to neutralize the acidity.

1.5.3 Formulation strategies of controlling µpH

Several studies have shown success in controlling μpH and stabilizing encapsulated proteins as a consequence. For example, introducing poorly soluble bases such as Mg(OH)₂ and MgCO₃, can counteract the often acidic environment and prevent the structural loss and aggregation of proteins [63, 67]. Blending PLGA 50/50 with 20% of polyethylene glycol (PEG) significantly increased the μpH to

above 5 during incubation up to 4 weeks, supporting the observed improved stability of ovalbumin and bovine serum albumin (BSA) in PLGA/PEG microspheres [68, 69]. The incorporation of PEG could increase the water uptake of the polymer due to its high hydrophilicity and solubility, which would dilute the acid concentration within the polymer system. In addition, PEG also enhanced the polymer permeability to water-soluble acids by its plasticization effect, thus promoting acid release. In another study, poly(ethylene glycol)-poly(L-histidine) diblock polymer (PEG-PH) was added as an excipient to the PLGA microspheres containing BSA [70]. This excipient formed ionic complex with BSA that stabilizes the protein. A neutral microenvironment was observed inside these microspheres. The poly (histidine) is a weak base that can neutralize the local acidity, and PEG can reduce acid accumulation as previously discussed.

1.5.4 Evidence of acidic microclimate in PLGA

1.5.4.1 Indirect evidence

It has long been recognized that a low-pH microenvironment commonly exists in the aqueous pores of PLGA delivery systems. The following are examples of indirect evidence indicating the presence of acidic microclimate in PLGA devices. Heterogeneous degradation in large (~1-2 mm dimensions) PLGA specimens was observed where the degradation proceeded more rapidly in the center than at the

surface [23]. Since PLGA degradation is acid catalyzed, it was concluded that acidic PLGA degradation products accumulated inside the matrix core, which created a low-pH environment and accelerated ester bond cleavage. Shenderova et al. [71] found that camptothecin was stabilized in its acid-stable (and active) lactone form when encapsulated in PLGA microspheres, and later confirmed an acidic microclimate inside the PLGA matrix. Insulin incubated under acidic conditions was analyzed using HPLC by Uchida et al. [72] and found to share the same retention time with the degradation products of insulin extracted from the microspheres, indicating the existence of an acidic environment within PLGA. Furthermore, it was proven that co-encapsulating antacids (e.g., Mg (OH)₂, and MgCO₃) in PLGAs could improve the stability and release kinetics of encapsulated proteins due to the inhibition of acid-catalyzed degradation of proteins, as demonstrated in studies with bovine serum albumin [63], basic fibroblast growth factor (bFGF) [63], bone morphogenetic protein-2 (BMP-2) [63], tetanus vaccine antigen (i.e., tetanus toxoid) [73], and tissue plasminogen activator [74].

1.5.4.2 Techniques for direct measurement of µpH

To accurately measure, predict and control the μpH in PLGA delivery systems, several techniques have been developed to directly quantify μpH . In one study, ^{31}P nuclear magnetic resonance (NMR) was employed to examine the microenvironment in PLGA microspheres incubated in sheep serum [75], which

served as a model physiological fluid containing endogenous inorganic phosphate, ³¹P. The study was based on the hypothesis that under these conditions, phosphate solutes diffuse into the microsphere from the outside medium, and the ³¹P chemical shifts between internal and external phosphate populations indicate the pH change. Results showed that pH stabilized around 6.4 over the course of 45 days. A major pitfall associated with this study was that the author failed to discuss the extent of penetration by phosphate ions into microsphere matrix interior. It is highly possible that the phosphate ions only partitioned into the surface pores, since there was little data on the permeability of phosphate ions in the polymer phase. Later on, the study was modified by encapsulating phosphate- and histidine- containing porogen excipients inside PLGA microspheres and then characterizing the internal environment using ³¹P and ³¹C NMR spectroscopy [76]. Results indicated that the pH was maintained below 4 but above the phosphoric acid pKa through the erosion period of 14 days. This study addressed the issue of phosphate distribution within microspheres and the results given by encapsulated solutes are representative of the PLGA microenvironment proteins resides. However, it is limited by the ability of only reporting a rough μpH range from recording the chemical shifts of ³¹P and ³¹C.

Electron paramagnetic resonance (EPR) was another technique developed to measure the microenvironment, including µpH in PLGA implants *in vivo* [77] and PLGA microspheres *in vitro* [78, 79] by incorporation of either free spin probes or spin-labeled drugs. The hyperfine splitting constant of the probe is sensitive to the

pH changes of environment due to the changes in the protonation state of the imidazolidine-derived structure of the probe, allowing the determination of pH after calibration. It was reported that the pH in PLGA microspheres dropped to a value equal or less than 4.7 within 13 hours. Measuring pH inside microspheres with EPR is a non-invasive and continuous method. However, since this technique relies on the mobility of spin probes, a reliable measurement is not possible in PLGA microspheres after 50 hours of erosion in the study due to the increase of microviscosity and decrease in signal to noise ratio causing changes in the spectral shape.

Potentiometric measurement was reported as a rapid and reliable way of determining μpH values in thin polymer films [80]. Briefly, standard glass pH electrodes were coated with PLGA films, and the zero-current potential was measured with respect to a reference electrode. This technique was developed based on the assumption that after hydration of the PLGA film, an aqueous layer is formed between the electrode and the PLGA coating that mimics the microenvironment in the aqueous pores inside the polymer. The μpH was deduced from the measured cell potential after correction of the diffusion and interfacial potentials in the electrochemical cell. It was found that μpH in PLGA 50/50 films with thickness of 30-100 μm declined to less than 3 after 1 day of incubation in a physiological buffer at 37°C and remained acidic for 4 weeks. And the thickness of PLGA films can influence the μpH development, as low pH (pH 2-3.5) was developed

for coatings with thickness above 30 μ m while neutral pH (pH 6.5) was observed if the coating was very thin (~7 μ m) after 1 week of incubation. Although potentiometric method is a simple and fast way for screening formulation strategies with desired controlled μ pH in PLGA polymer, the pH measured could not be extrapolated to the μ pH in smaller delivery devices such as microspheres and nanospheres, due to the different geometry, structure and transport characteristics inside such systems. Overall, the above-mentioned measurements all suffer from the limitation of only providing an averaged, general picture of the microenvironment pH inside PLGA matrix. It is highly likely that the μ pH is unevenly distributed and that some pores of extreme high acidity are present despite an overall neutral measured pH, leading to the destabilization of entrapped proteins in those low-pH regions.

Confocal microscope imaging techniques, on the other hand, can directly visualize the spatial µpH distribution within microspheres or throughout a device by encapsulation of pH-sensitive fluorescent probes. Shenderova *et al.* [71] first employed confocal laser scanning microscope to monitor the µpH by correlating the pH with fluorescent intensity. Fluorescein was encapsulated in microspheres, whose emission intensity decreases with the decrease in pH. However, since dye partitioned in the polymer and the measured fluorescence intensity was also dependent on dye concentration, this method by itself was only semi-quantitative. In order to eliminate the effect of poorly controlled dye concentration on

fluorescence, Fu et al. [81] improved this confocal microscopic imaging method by co-encapsulating two fluorescent dye-dextran conjugates (SNARF-dextran and NERF-dextran) in the microspheres and correlating their intensity ratios at two respective emission wavelengths with pH. Hence, quantitative pH information that is independent of dye concentration could be acquired from this ratiometric method. The presence of acidic microenvironment with minimum pH as low as 1.5 was suggested and the formation of pH gradients within the microspheres were demonstrated using this technique. Results also suggested that the µpH development was dependent on the size of microspheres. However, since both dves emit in the green range (535 nm for NERF and 580nm for SNARF), the resolution of images was very poor. Besides, the ratio images were not properly processed; giving rise to high noise-to-signal ratio that undermines the assay's accuracy. In addition, the range of pH measurement was relatively narrow (pH from 1.5 to 3.5 in standard curve), restricting its application to systems with less acidic pH.

Another quantitative ratiometric method based on confocal imaging was developed by Li *et al.* by encapsulation of dextran-SNARF-1® conjugate as the fluorescent probe [67]. The ratio of fluorescent intensities of the dye at two emission wavelengths, typically 580 and 640 nm is responsive to pH change in the neutral range (pH 5.8-8.0), thereby providing pixel-by-pixel neutral µpH maps inside PLGA microspheres. This technique was applied to determine µpH in both acid-neutralized and non-neutralized PLGA microspheres during extended

incubation in physiological buffer. The noise-to-signal ratio was significantly reduced after performing a series of image processing steps. Hence, the accuracy of this method was greatly improved. The measured pH in the neutral range may serve well for formulation screening purposes; nevertheless, it is inadequate to provide information on understanding the acidic μpH development in PLGA degrading matrices.

Later, mapping of µpH in acidic pH range in PLGA microspheres was reported using the same technique [66]. Lysosensor yellow/blue® dextran conjugate was employed as the fluorescent dye to sense µpH over a broadly acidic range (2.8<µpH<5.8) inside PLGA microspheres by relating pH to ratio of its fluorescent intensities under two emission wavelengths. This method is accurate to within ± 0.2 pH units, as found with dextran SNARF-1® in the neutral pH range. The ratiometric method is advantageous in that it eliminates artifacts resulted from the variations of dye concentration inside microspheres, including the leakage of the dye, non-uniform distribution of dye within a microsphere and/or differences in dye loading among populations of microspheres. The high molecular weight dye conjugation prevents the dye from quickly releasing out of the microspheres. It also localizes the dye within the aqueous pores of the microspheres, similar to the partition behavior of proteins in the polymer.

Combining the above two pH mapping techniques using confocal laser scanning microscopy (CLSM) could allow us to obtain accurate µpH distribution

over the entire useful pH range (2.8-8.0), which is beneficial to further our understanding of μpH development and promote formulation designs for optimized PLGA delivery of acid-labile bioactive drugs.

1.6 Thesis Overview

In this dissertation, the μpH inside biodegradable polymeric microspheres was quantitatively evaluated. The overall objective of this dissertation is to employ μpH mapping and related tools to further our mechanistic understanding of μpH development and control in order to improve formulations of protein encapsulated biodegradable microspheres.

Despite numerous studies on measuring μpH , accurate μpH mapping has not been accomplished in the presence of protein. The purpose of Chapter 2 was to develop a method to map the μpH using CLSM in protein-encapsulated PLGA microspheres. Correction of the interference of encapsulated proteins on dye's fluorescent reporting of pH was performed based on the estimation of protein concentration in PLGA pores. This technique was then applied in examining the μpH distribution and kinetics in different PLGA microspheres formulations with the incorporation of pH-modifying excipients or adjusting the formulation variables.

Chapter 3 presents the µpH mapping using CLSM in microspheres of a hydrophilic biodegradable polymer, poly(lactide-co-hydroxymethyl glycolide) (PLHMGA) that have previously displayed better stability of encapsulated

biomacromolecules than PLGAs with similar degradation time-scales. The transport of a fluorescent probe, bodipy, in microspheres of PLHMGA and PLGA was studied to illustrate the relevance of the faster diffusion of water-soluble acids in the development of less acidic μpH .

Previous studies have implicated the role of water-soluble impurities and degradation products in governing the µpH in large PLGA specimens, but mathematical models for µpH prediction were equilibrium-based without considering the concentration gradients of diffusing acids. In Chapter 4, a mathematical model for the simulation of µpH distribution in more commonly used PLGA microspheres as a function of degradation time was developed. The goal here was to gain a better understanding of the physicochemical mechanisms of µpH development and to evaluate the effect of different variables on µpH distribution and kinetics. Key parameters in the model involving the mean size of microspheres, the initial concentration of acids in polymer, acid production rate, and acid diffusion rate in polymer matrix was estimated from independent experiments. The

In Chapter 5, the thesis concludes by proposing various studies that warrant future investigation to build upon the μpH mapping and simulation studies described herein.

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CHAPTER 2

Mapping Microclimate pH Distribution inside Proteinencapsulated PLGA Microspheres Using Confocal Laser Scanning Microscopy

2.1 Abstract

The pH in the aqueous pores of poly(lactide-co-glycolide) (PLGA) matrix, also referred to microclimate pH (μpH), is often uncontrolled ranging from highly acidic to neutral pH range. The μpH distribution inside protein-encapsulated PLGA microspheres was quantitatively evaluated using confocal laser scanning microscopy. The fluorescent response of Lysosensor yellow/blue® dextran used to map μpH in PLGA was influenced by the presence of encapsulated protein. The non-protonated form of the pyridyl group on the fluorescence probe at neutral pH was responsible for the interference, which was dependent on the type and concentration of protein. A method for correction of this interference based on estimating protein concentration inside the microspheres was established and validated. After correction for this influence, the μpH distribution and kinetics inside microspheres was evaluated for different PLGA 50/50 microsphere

formulations under physiological conditions for 4 weeks. Generally, the μpH acidity increased with the increasing of incubation time. The co-incorporation of a poorly soluble base, magnesium carbonate, in the microspheres postponed the appearance of detectable acidity for up to 3 weeks. Co-addition of an acetate buffer was able to control the μpH over a slightly acidic range (around pH 4.7) after two weeks of incubation. Microspheres prepared from a lower polymer concentration exhibited a higher μpH , likely owing to reduced diffusional resistance to acidic degradation products out of the microspheres. The stability of protein was enhanced by addition of MgCO₃, acetate buffer, or by reduced polymer concentration in the preparation, as evidenced by more soluble protein recovered after incubation. Hence, the μpH imaging technique developed can be employed in the future for optimization of formulation strategies for controlling μpH and stabilizing encapsulated proteins.

KEY WORDS: microclimate pH; microspheres; confocal laser scanning microscopy; poly(lactide-*co*-glycolide); pH distribution

2.2 Introduction

Poly(lactide-*co*-glycolide) (PLGA), as one of the most important classes of biodegradable and biocompatible polymers, has long been the research focus of controlled delivery of biomacromolecules, including peptides, proteins, and vaccines [1-6]. Despite its excellent safety and versatility, a major drawback associated with

this polymer is the common acidification and lack of control of its microenvironment inside the polymer matrix during erosion, as a result of acidic polymer impurities and the build-up of acidic monomers and oligomers generated from polymer hydrolysis. Consequently, the integrity of encapsulated acid-labile proteins can be greatly compromised during release [7-9].

Several studies, using indirect methods, have shown evidence of an acidic microclimate within degrading PLGA devices. For example, faster degradation in the center of large PLGA specimens (~1-2 mm dimensions) was observed due to the accelerated hydrolysis of ester bondage catalyzed by the acids accumulated at the matrix core [10]. Additionally, Shenderova *et al.* [11] found that camptothecin was stabilized in its acid-stable (and active) lactone form when encapsulated in PLGA microspheres. Furthermore, co-incorporation of antacids, such as Mg(OH)₂, MgCO₃ and ZnCO₃ in PLGAs could strongly inhibit protein structural losses and aggregation for over one month, as demonstrated in studies with model as well as therapeutic proteins [12-15].

Moreover, techniques have been developed to directly quantify μpH inside PLGA delivery systems, including ³¹P nuclear magnetic resonance (NMR) [16], electron paramagnetic resonance (EPR) [17], potentiometry [18], and confocal microscopy imaging [19-21]. The first three methods are limited to providing an averaged μpH . Confocal microscopy imaging, on the other hand, by encapsulating

fluorescent pH-sensitive probes, is capable of delineating a detailed μpH map noninvasively within the polymer matrix.

After early attempts to develop a quantitative ratiometric method of µpH measurement using confocal laser scanning microscopy [19], our group found that SNARF-1® dextran [20] and Lysosensor yellow/blue® dextran [21], as fluorescent probes encapsulated into PLGA microspheres, could sense pH changes from pH 5.8 to 8.0, and pH 2.8 to 5.8, respectively. Thus, after confocal image processing, an accurate pixel-by-pixel µpH distribution map either in the neutral or acidic range could be created [20, 21]. This ratiometric method is advantageous in that it eliminates artifacts including photo bleaching, leakage of the dye probe, and non-uniform distribution of dye within microspheres. The dextran-conjugated probes employed are water-soluble macromolecules, thereby localizing themselves in the aqueous pores where protein resides [19].

In the present study, μpH inside PLGA microspheres encapsulating both protein and Lysosensor yellow/blue® dextran was accurately quantified using confocal microscopy imaging. To accomplish this, significant interference of the dye response from the presence of protein was corrected by estimating protein concentration inside the PLGA pores to perform the measurement. The acquired knowledge is beneficial to further our understanding of μpH development and promote formulation designs for optimized delivery of pH-sensitive biomacromolecules.

2.3 Materials and Method

2.3.1 Materials

Poly(D,L-lactide-*co*-glycolide), end capped, 50/50 with inherent viscosity (i.v.) of 0.6 dl/g in hexafluoroisopropanol at 25 °C was purchased from Durect Corporation (Birmingham, AL). The fluorescent pH sensitive probe, Lysosensor yellow/blue® dextran (MW=10 kDa) was purchased from Invitrogen (Eugene, OR). Bovine Serum Albumin (BSA, fraction V), was purchased from Sigma Chemical Co. (St. Louis, MO). Polyvinyl alcohol (PVA, 80% hydrolyzed, MW 9-10 kDa) was supplied by Polysciences Inc. (Warrington, PA). All other chemicals were of analytical grade or higher were obtained from commercial suppliers.

2.3.2 Preparation of microspheres

Protein-encapsulated PLGA microspheres containing Lysosensor yellow/blue® dextran as an acidic pH sensitive probe were prepared using the w/o/w double emulsion-solvent evaporation method. Briefly, 100 µl of 300 mg/ml BSA with 25 mg/ml dye in double distilled water was added to 1 ml of 400 mg/ml PLGA solution (40% w/v) in methylene chloride. The mixture was then homogenized using a Tempest IQ² homogenizer (The VirTis Co., Gardiner, NY) at 7,500 rpm for 1 min to generate first w/o emulsion, followed by quickly adding 1 ml of PVA solution (2% w/w). After vortexing for 20 s, the formed w/o/w emulsion

was poured slowly into 100 ml of PVA solution (0.5% w/w) and stirred at room temperature for 3 hours to extract and evaporate the organic solvent. Then, the hardened microspheres were harvested and sieved for $45-63 \mu m$ size. After washing with double distilled water three times, the microspheres were freeze-dried on a FreeZone 2.5 Liter Benchtop freeze dry system (Labconco, Kansas City, MO).

For microspheres containing BSA of a specific pH, 100 mg/ml BSA solution was first titrated with HCl to pH 3, 4, and 5, respectively, followed by freeze-drying. The lyophilized powder was then reconstituted with water and encapsulated with dye in PLGA microspheres as described above. Microspheres containing magnesium carbonate were prepared by suspending 3% (w/w) of the base to the polymer solution with all other conditions as described above. To prepare microspheres encapsulating acetate buffer, BSA and dye were dissolved in 100 μ l of 0.1 M sodium acetate buffer of pH 4.6 to make the water phase, with other conditions unchanged. Microspheres with a lower polymer concentration (30% w/v) were also prepared following the same procedures.

2.3.3 Confocal laser scanning microscopy for microspheres imaging

A ratiometric method based on a confocal microscopy imaging technique was employed similarly as described by Ding *et al.* [21]. A Carl Zeiss LSM 510-META laser scanning confocal microscope (LSCM, Carl Zeiss Microimaging, Inc., Thornwood, NY) was equipped with an Enterprise UV laser and a Carl Zeiss

Axiovert 100 M inverted microscope. The fluorescent dye that was encapsulated in the microspheres was excited at 364 nm, and the emission at two wavelengths, 450 nm and 520 nm were recorded. All measurements were conducted using a C-Apochromat 63X water immersion objectives lens with a numerical aperture of 1.2. The detection gain was set at 650, and the pinhole was 328 μ m, which resulted in optical slice of thickness of 5 μ m. The laser power was set at 40% of its full power. The image size was 512x512 pixels and the images were scanned by 8 bit plane mode at a scan speed of 6.40 μ s/pixel.

2.3.4 Calibrating fluorescence intensity ratio vs. pH in the presence of protein

A set of universal buffers with pH ranging from 2.8 to 5.8 were prepared using combined 0.1 M citric acid and 0.2 M Na_2HPO_4 solutions. A certain concentration of protein solutions (BSA or lysozyme, e.g. 100 mg/ml) was prepared by dissolving protein in the buffers and then titrating the solution to its original pH. Lysosensor yellow/blue® dextran was then dissolved in the protein buffer solutions with a concentration of 1.2 mg/ml.

Images of dye solution were obtained under confocal microscope at 450 nm and 520 nm. The acquired images (n=8) were processed by frame averaging, followed by neighborhood averaging, and applying a median filter as described by Li et al. [20] using Image J software (developed by National Institutes of Health and available on the internet at http://rsbweb.nih.gov/ij/) to eliminate the signal noise

and obtain accurate pixel value. The standard curves were established by plotting the ratio of mean pixel intensity of the dye solutions at two emission wavelength, 450 nm and 520 nm *vs.* the pH of that solution.

2.3.5 Microclimate pH mapping inside microspheres

Microspheres (20–25 mg) were incubated in 1 ml phosphate buffer saline (7.74 mM Na₂HPO₄, 2.26 mM NaH₂PO₄, 137 mM NaCl and 3 mM KCl) containing 0.02% tween 80 (PBST, 10mM, pH=7.4) at 37°C under mild agitation at 320 rpm by a KS 130 basic shaker (IKA® Works Inc., Wilmington, NC). At pre-determined time points, the release medium was replaced with fresh buffer and a small amount of microspheres were collected and placed under confocal microscope while focusing at the center of microspheres to obtain images (n=5). After image processing [20], the ratio of fluorescence intensity I_{450nm}/I_{520nm} at each pixel having intensity above the threshold value (indicating the fluorescence from release media) of the images was calculated and assigned to a pH from the standard curves independent of dye concentration. In the processed images, each pixel was converted to a color corresponding to specific pH. When plotting the upH distribution curves, the probability of a specific pH inside microspheres was calculated by taking the amount of pixels corresponding to that pH divided by the total pixels of the microspheres. For intensity ratios exceeding the limit of standard curve, the pH was

assigned to either below 2.8 or above 5.8. In such cases, their percentage was plotted as the boundaries of the distribution curves accordingly.

The μpH could be accurately mapped within \pm 0.2 pH unit over pH from 2.8 to 5.8 (see Supporting Information 2.6.1 for statistical analysis).

2.3.6 Determination of protein loading and encapsulation efficiency

The amount of protein encapsulated in PLGA microspheres was determined by direct recovery from the polymer matrix [13]. Eight mg of microspheres were dissolved in 2 ml acetone. The mixture was vortexed and centrifuged at 8,000 rpm for 10 min, followed by removal of the acetone. After repeating the above procedures three times, the BSA pellet was air-dried and reconstituted in PBST and incubated at 37° C for 1 h. The protein concentration was then determined using Coomassie* Plus protein assay reagent. The working range in this study was from 25 µg/ml to 500 µg/ml (assay sensitivity is from 1 µg/ml to 1500 µg/ml) and not interfered by reagents used in our experiments. Protein loading was calculated from the amount of protein recovered divided by the mass of microspheres. Encapsulation efficiency was obtained from the ratio of actual protein loading to the theoretical protein loading. All measurements were performed in triplicate (n=3).

2.3.7 Release and stability of protein from microspheres

Microspheres (20–25 mg) were incubated in 1 ml PBST (10mM, pH=7.4) at 37 °C under mild agitation at 320 rpm. At pre-determined time points, the release media was removed after centrifugation at 5,000 rpm for 5 min and replaced with fresh buffer. The protein concentration in the release media was determined using Coomassie® Plus protein assay reagent.

At the end of the release study, soluble protein was recovered from PLGA microspheres as described in the loading study. Any remaining insoluble aggregates were collected by centrifugation and dissolved in denaturing solvent (PBST/6M urea/1mM ethylenediaminetetraacetic acid (EDTA)) and incubated at 37°C for 30 min to dissolve non-covalent bonded aggregates. Finally, any insoluble aggregates were collected again and dissolved in reducing solvent (denaturing solvent plus 10 mM dithiothreitol (DTT)) to dissolve any disulfide-bonded aggregates. The protein content in each step was all analyzed with Coomassie® Plus protein assay reagent using the appropriate solvent as diluent for protein standards. All measurements were performed in triplicate (n=3).

2.3.8 Water uptake of microspheres

Microspheres (20–25 mg) were incubated in 1 ml PBST (10mM, pH=7.4) at 37 °C under mild agitation at 320 rpm. At pre-determined time points, the microspheres were collected and the surface water was removed by filtration and

the wet weight (W_1) of the microspheres was recorded. The samples then were dried under vacuum to a constant weight and the dry weight (W_2) was recorded.

To correct for the interparticle water, dry microspheres were suspended in PBST at room temperature and rapidly filtered and dried as described above. Assuming little water uptake by the microparticles between suspension and filtration, the weight differences between wet and dry particles accounted for the portion of interparticle water (W_i) , as defined by

$$W_{i} = \frac{W_{1}^{'} - W_{2}^{'}}{W_{2}^{'}} \tag{1}$$

Where W_1 ' and W_2 ' are the weights of wet microspheres and dry microspheres after immediate collection (t=0), respectively. The water uptake of microspheres at time t ($W_P(t)$) was estimated by:

$$W_{P}(t) = \frac{W_{1} - W_{2} - W_{2} \times W_{i}}{W_{2}}$$
 (2)

Where W_1 and W_2 are the wet and dry microsphere weights at time t. Note that in control experiments the interparticle water estimation did not significantly depend on the temperature of water used, e.g. 4°C, 25°C and 37°C (data not shown). All measurements were performed in triplicate (n=3).

2.3.9 Correction of protein interference on µpH mapping

To account for the influence of protein on dye emission, corrections were necessary to acquire an accurate estimation of the μpH in the presence of significant BSA. The influence of lysozyme was significantly less, and therefore, its correction was not considered further. Since protein concentration inside microspheres changes during incubation due to the protein release and water uptake by the polymer matrix, corrections were done for each time point of pH mapping. The average protein concentration ($C_p(t)$) inside the microsphere aqueous pores at time t can be estimated by the following equation:

$$C_p(t) = \frac{M_p}{V_{pores}} = \frac{M_{P,0} \times l \times (1 - f(t)) \times \rho_w}{M_P(t) \times W_P(t)}$$
(3)

where M_p is the mass of protein in microspheres. V_{pores} is the volume of aqueous pores. $M_{P,0}$, $M_P(t)$ are the initial and time dependent mass of microspheres respectively. l is the fraction of protein loaded. f(t) is the fraction of protein release from microspheres. And ρ_w is the density of water.

When the estimated protein concentration was not the same as those in known standard curves (Figure 2.1B), the corresponding fluorescence ratio *vs.* pH curve was interpolated. (see Supporting Information 2.6.2)

2.4 Results and discussion

2.4.1 Interference of protein on fluorescent response of the dye

Lysosensor yellow/blue® dextran, which is sensitive to changes in acidity from roughly pH 2.8 to 5.8, was selected as a fluorescence probe to investigate the μpH inside PLGA microspheres, as previously reported [21]. Adding protein to the dye solutions also provided a fluorescent intensity emission ratio ($I_{450 \text{ nm}}/I_{520 \text{ nm}}$) vs. pH standard curve well fitted to a third-order polynomial function (r^2 =0.999) from pH 2.8 to 5.8 (Figure S2.3). The pH sensitivity of the dye is concentrationindependent as well, which ensures the standard curve is not affected when the dye concentration changes in microspheres during incubation. However, as protein concentration was raised to elevated levels (e.g., >25 mg/ml of BSA), the fluorescent response of the dye was significantly influenced by the presence of protein, and this interference was dependent on the specific protein. For example, as shown in Figure 2.1A, the pH sensitivity differed for dye solutions with or without presence of protein, with the presence of BSA giving more pronounced changes in emission intensity ratio compared to that of lysozyme. As expected from Figure 2.1A, the fluorescence ratio was significantly affected by the protein concentration over wide range. As shown in Figure 2.1B, the intensity ratio at a certain pH for BSA concentration of 0 to 500 mg/ml rose as protein concentration was increased, with little or no influence at pH 2.8 to an extensive effect at pH 5.8. The sensitivity of dye at high pH corresponding to the presence of the non-protonated form of the dye's pyridyl group implicates this dye species as responsible for the protein interference. Note that by 500 mg/ml BSA at highly acidic pH resulted in a gel formation, consistent with the low pH unfolding of the protein [22] and noncovalent aggregation of BSA in PLGA [12, 13], which was associated with an unexpected increase in the intensity ratio at the pH of 2.8 (Figure 2.1B).

The mechanism of protein interference on the fluorescent response of the non-protonated form of dye is not well understood, although it was demonstrated that the presence of protein would quench the emission of dye at 520 nm and shift the emission peak at 450 nm slightly to a shorter wavelength in the fluorescence spectrum (Figure S2.4). Ground state interactions between dextran-dye and protein, e.g., binding, was not likely to cause the interference, considering the emission ratio did not depend on dye concentration in presence of either lysozyme or BSA (Figure S2.3). Processes involving excited state of fluorophores, such as energy transfer or collisional quenching induced by the protein, were more likely responsible for the interference.

2.4.2 Correction of BSA effect on dye interference and BSA buffering capacity

As described in the Materials and Methods, we estimated the BSA concentration in the microspheres to correct for the BSA interference on the µpH reporting of the dye. Key assumptions involved estimates of interparticle water and assuming uniform BSA concentration in the pores. In addition, for this polymer molecular weight, very little water partitions in the polymer phase until late stages

of polymer erosion [23]. To validate our approach of correction, we compared the upH measured inside microspheres encapsulating BSA after 1 day incubation in PBST at 37°C as a function of various pH of BSA solutions used to form the primary emulsion during microsphere preparation. We hypothesized that after 1-day incubation, a concentrated protein solution would be formed due to the water penetration into the polymer matrix. Therefore, the µpH would be dictated by the pH of the encapsulated protein solution in aqueous pores as significant degradation of polymer is not expected at such an early time of incubation [23, 24]. Moreover, from upH measurement of microspheres without encapsulating protein, little acidity was observed (See Figure 2.3B), indicating the lack of significant acid impurities. Encapsulated BSA of different pH was prepared by titrating 100 mg/ml BSA solution to a specific pH and then freeze-drying. The estimated protein concentration after 1day incubation inside polymer pores was roughly 500 mg/ml (within ±10%) for each formulation, as calculated from (3). Thus, upH values were estimated from fluorescence ratio vs. pH standard curve with 500 mg/ml BSA. The processed confocal images and µpH distribution curves are shown in Figure 2.2 & Figure 2.3, respectively, and corresponding results are summarized in Table 2.1. The estimated upH after correction was very close (difference within 0.1 pH unit) to the pH of the concentrated protein solution, strongly supporting the approach of correction.

Table 2.1 pH comparison of concentrated BSA solution and average µpH after 1 day incubation from confocal microscopy after correction of protein interference.

_			
	pH of 100 mg/ml BSA solution	pH of 500 mg/ml BSA solution ^a	Average μpH from confocal imaging ^b
	3.0	3.2	3.2
	4.0	4.1	4.2
	5.0	5.1	5.0
	7.0	7.0	>5.8

^a The solution was made by reconstitution of lyophilized BSA power from 100 mg/ml solution of specific pH, as described in Materials and Methods.

A slight acidity was observed in PLGA microspheres encapsulating only dye after the 1-day incubation (Figure 2.2A), which could be ascribed the existence of a very low level of acidic impurities in the polymer. However, upon incorporation of BSA, the pH in most aqueous pores were raised above 5.8, with more than 95% of pixels out of detection limit compared to 80% in microspheres without protein (Figure 2.3B), consistent with significant buffering capacity of the encapsulated BSA. Moreover, the µpH was more homogenously distributed inside microspheres encapsulating protein (Figure 2.2E).

2.4.3 Mapping µpH distribution and kinetics in degrading PLGA microspheres

After the successful test of the correction for BSA interference on µpH measurement, µpH distribution and kinetics were examined and compared in degrading PLGA microspheres prepared from different formulations during a one-

^b The μpH was controlled by the inner water phase pH, as described in Materials and Methods.

month incubation. The processed confocal images visualized the μpH distribution by color, and the distribution curves provided a quantitative illustration. It should be noted that the μpH mapped refers to the μpH distribution within the region of optical slice of confocal imaging (5 μm under the condition of this experiment). Presumably the fluorescence could only be observed in areas of aqueous pores; however, since pores at different planes within this region do not vertically overlay with each other, fluorescence could be observed through the entire image at the beginning of incubation.

Little acidity was observed in microspheres made from polymer concentration of 40% (w/v) at the beginning of incubation, as evidenced by more than 95% of pixels in the images out of detectable limit of the dye (pH 5.8). As the incubation progressed, the µpH decreased steadily until day 21 in accordance with the accumulation of water-soluble acids generated by the degradation of the polymer. After 21 days, the µpH maintained mostly in the range of 4 to 5.8 (Figure 2.5A), probably due to the onset of polymer erosion and liberation of water-soluble acids out of the polymer to balance acid production rate. From the processed confocal images (Figure 2.4A), the acidity was observed to be higher in the center of microspheres than in the peripheral regions, consistent with development of an expected diffusion/reaction mechanism governing polymer distribution of acidic degradation products. The blank regions inside the microspheres indicate no

detectable fluorescence, corresponding to regions of free dye loading or extensive dye release from the polymer.

Addition of magnesium carbonate delayed the appearance of detectable acidity inside microspheres up to 3 weeks (Figure 2.4B & 2.5B) possibly by multiple mechanisms, including: i) dissolution and direct neutralization of PLGA-produced acids and ii) water uptake and pore formation imparted by the osmotic Mgcarboxylate salts resulting from acid-base titration, which increases liberation of sequestered acids. The minimization of the pH drop conferred by incorporation of base in PLGAs was supported indirectly in a previous study from decreasing the degradation rate of the polymer [12]. Moreover, this effect was further confirmed in a quantitative way using a neutral pH sensitive dye [20]. In this study, the effect of MgCO₃ incorporation was examined by the changes in acidic pH in BSA-containing PLGA 50/50 microspheres. Consistent with previous studies with larger millicylidrical implants [13], elevated water uptake was observed in microspheres upon incorporation of MgCO₃ (Figure 2.6B), as a result of the osmotic pressure generated by the Mg-carboxylate salts described above. Since more water channels were created, encapsulated protein was released slightly faster than that of without base (Figure 2.6A). The same trend was also expected for dye release in microspheres with MgCO₃. The rapid release of dye, therefore, was associated with a higher fraction of blank regions in confocal images than in base-free formulations. In addition, the decrease of the µpH by 28 days of incubation may have been caused

by the depletion of base from the polymer. Note that water uptake (Figure 2.6B) and protein release kinetics (Figure 2.6A) were used to estimate protein concentration kinetics inside polymer pores (Figure 2.6C) for correction of protein interference on µpH mapping, as described in the Materials and Methods.

In order to control upH over a moderate acidic range, water-soluble buffering species were co-encapsulated with protein into PLGA microspheres. This was achieved by adding to the protein inner water phase a buffer solution (0.1 M acetic acid and sodium acetate, pH=4.6). During incubation, the aqueous pores inside microspheres would be filled with buffering species with protein so long as these species are retained in the polymer. From confocal images (Figure 2.4C) and upH distribution curves (Figure 2.5C), the µpH gradually dropped from neutral pH as incubation progressed. After 14 days incubation, the average µpH was maintained from 4.6 to 4.8. The relatively high pH during the initial stage of incubation was due to the very high protein concentration in aqueous pores, which acted also as a buffer, undermining the buffering capacity of acetate salts. As water imbibed into and protein released out of polymer matrix, protein concentration decreased. Meanwhile, despite the possible removal of acetic acid, water-soluble acids were generated from polymer degradation, leading to $\mu p H$ approach to the pH of acetate's maximal buffer capacity (pKa=4.7).

The effect of polymer concentration on µpH distribution kinetics was also examined in our study. Microspheres were prepared from a lower polymer

concentration of 30% (w/v). Compared to that made of 40% (w/v), the μpH was much less acidic upon 14 days incubation (Figure 2.4D & 2.5D). This could be rationalized by the fact that microspheres made from solution of lower polymer concentration usually possess more porous internal structure [25], which likely caused a higher effective diffusivity of acidic degradation products through the polymer matrix [26] and facilitated their liberation as a result. After 21 days incubation, the effect of polymer concentration on μpH was not apparent, which was presumably due to the changes in polymer properties, e.g. degradation rate, so that the difference in diffusion rate of acids was not significant.

2.4.4 Assumptions for correction and anticipated error

The correction for BSA's effect on the dye's fluorescent response was based on multiple assumptions. One important assumption for correction is that protein is evenly distributed inside aqueous pores in microspheres. This was supported by lysozyme's homogeneous distribution inside PLGA microspheres prepared by w/o/w double emulsion method, as observed by confocal laser scanning microscopy and infrared microscopy [27]. In the confocal images recorded in this study (Figure 2.4), the fairly homogeneous distribution of dye (being a water-soluble macromolecule as protein) was observed with the exception of formulations with base added and at the very last time point of 28 days, indicating the similar behavior of protein inside microspheres. Another assumption is that the protein

concentration in the cavities of microspheres was relatively uniform. In the case that local protein concentration in some pores were higher than the estimated average protein concentration, the $\mu p H$ distribution curves would generally shift to a lower p H.

Since the average protein concentration in PLGA pores at each point during controlled release was based on estimation from equation (3), errors may be associated with the deviation of estimation from the actual protein concentration, affecting the µpH measurement. The kinetics of estimated protein concentration inside microspheres during one month incubation are displayed in Figure 2.6C. Among all the parameters used to calculate the averaged protein concentration, the time dependent water uptake is most variable because the fraction of interparticle water may be changing during incubation depending on the property of polymer. Assuming there was 20% of error involved in experiments of estimating water uptake of microspheres, the upH kinetics corrected from protein concentration accounting for this \pm 20% of error (-17% to +25% of protein concentration) for microspheres with or without encapsulating MgCO₃ are shown in Figure S2.5. As demonstrated, the resulted µpH was not significantly affected (within 0.2 pH unit). Hence, the correction was only modestly influenced by small deviations based on the interparticle water assumption.

2.4.5 Formulation effects on protein stability

Insoluble protein aggregation has been observed when encapsulated in PLGAs, which has been linked to the acidic environment in PLGAs. For example, BSA was found to become hydrolyzed and form noncovalent aggregates by hydrophobic interactions when encapsulated in PLGA 50/50 millicylindrical implants [12, 13, 15]. Therefore, we analyzed the composition of residual protein inside PLGA microspheres after 4 weeks incubation in terms of soluble and insoluble protein [12] and the results were summarized in Table 2.2.

Table 2.2 Release and stability of various microsphere formulations after 28 days incubation.

Formulation Polymer conc. (w/v)/excipient	Loading (%)	Encapsulation efficiency (%)	Released (%) ^b	Soluble residue (%) ^b	Insoluble residue (non-covalent) (%)b	Insoluble residue (covalent)	Recovery (%) <i>b, c</i>
40 % (w/v)	4.7 ± 0.2 ^a	68 ± 3	18.1 ± 0.2	9 ± 2	34 ± 5	8 ± 1	71
40 % (w/v) w/MgCO ₃	5.2 ± 0.1	75 ± 2	20.9 ± 0.3	44 ± 5	21 ± 2	4 ± 1	90
40 % (w/v) w/acetate buffer	4.9 ± 0.1	69 ± 1	17.1 ± 0.7	41 ± 2	16 ± 3	5 ± 1	79
30 % (w/v)	3.9 ± 0.1	62 ± 1	23.3 ± 0.4	46 ± 1	12 ± 2	2 ± 1	83

^a All data are reported as mean \pm SD, n=3.

^b Determined after 28-day release.

 $^{^{}c}$ Recovery (%) = released (%) + soluble residue (%) + insoluble residue (%).

Incorporation of poorly soluble base (MgCO₃) significantly improved protein stability in terms of aggregation, as 44 ± 5 % of soluble residue remained after 28 days release relative to $9 \pm 2 \%$ in microspheres without any excipients. The mechanism of stabilization is believed to occur primarily via raising the acidic µpH in degrading PLGA matrix, as displayed in our confocal images and μpH distribution curves. This stabilization effect conferred by antacids was also shown in other therapeutic proteins [12, 15]. Some degree of aggregation persisting in these formulations is consistent with our previous data with microspheres prepared with the ester-end-capped PLGA 50/50 [12]. The 90% recovery in this formulation also suggested reduced protein hydrolysis than the other samples (71-83% recovery), as low recovery likely results from a lack of recognition of hydrolyzed protein by the Coomassie® Plus protein assay reagent [28]. Note that higher water content and slightly faster release (and thus, less remaining encapsulated protein to become damaged) are also potential effects to decrease the 28-day aggregation values [13, 15].

Addition of acetate buffer also reduced protein aggregation inside PLGA microspheres during one-month incubation. In this case, the pH profile was very similar to that recorded in the most unstable preparation (prepared with 40% w/v polymer concentration), albeit just slightly higher in the vicinity of the high buffering capacity of the acetate buffer. This data suggests perhaps other factors involved, e.g., the elevated water content and strongly reduced protein

concentration (see Figure 2.6) or the different ionic strength anticipated in the microclimate of this formulation affecting protein's stability. We also note that the small changes in pH in the vicinity of the first unfolding transition of BSA [22] may have been important.

Preparing microspheres from a lower polymer concentration also resulted in enhanced protein stability. This can be attributed to a less acidic microclimate developed in degrading PLGA specimen during the course of incubation. A more porous internal structure can take more water, thereby increasing the effective diffusion coefficient of the detrimental water-soluble acids in the polymer matrix [26] and accelerating their release from the microspheres.

2.5 Conclusions

An uncontrolled and often acidic µpH is regarded as one of the most deleterious factors responsible for the instability of encapsulated protein in PLGA delivery systems. Therefore, it is important to develop methods of quantitative description of the microenvironment in PLGA. In our study, we demonstrated that µpH mapping in the polymer was affected by the presence of encapsulated protein, whose interference on fluorescent response of dye depends on the type and concentration of protein. µpH distribution in microspheres with protein and/or excipients could be quantitatively evaluated using confocal laser scanning microscopy after correction of the interference of protein. This µpH mapping

technique is presented as a valuable tool for the study of $\mu p H$ development mechanisms and design of formulation methodologies that control $\mu p H$ with stabilized biomacromolecules.

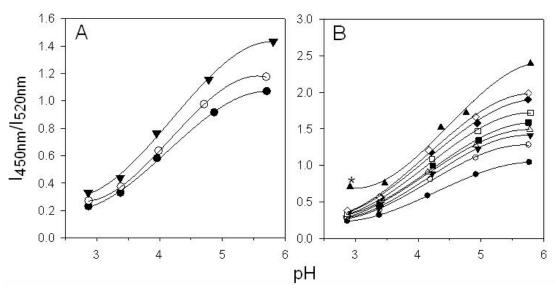


Figure 2.1 Interference of confocal pH measurement of Lysosensor yellow/blue[®] dextran as a function of pH **(A)** by the presence of 100 mg/ml of BSA (\blacktriangledown), 100 mg/ml of lysozyme (\bigcirc), or absence of protein (\bullet); **(B)** by the presence of BSA at the concentration of 0 mg/ml (\bullet), 25 mg/ml (\bigcirc), 50 mg/ml (\blacktriangledown), 75 mg/ml (\triangle), 100 mg/ml (\blacksquare), 150 mg/ml (\square), 200 mg/ml (\bullet), 250 mg/ml (\bigcirc), and 500 mg/ml (\blacktriangle). The concentration of fluorescence dye was 1.2 mg/ml. Lines represent best fits to a third order polynormial function of the experimental data. SD for all data points were less than 2% of mean (n=8). * BSA formed a gel-like phase at this protein concentration and pH.

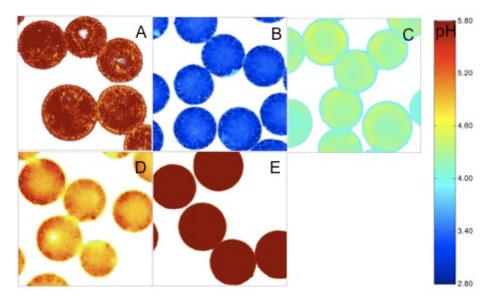


Figure 2.2 Processed confocal images of microspheres encapsulating dye only **(A)**; dye and BSA of pH of 3 **(B)**; dye and BSA of pH of 4 **(C)**; dye and BSA of pH of 5 **(D)**; and dye and BSA of pH of 7 **(E)** after incubation at 37° C in PBST buffer for 1 day. The μ pH was controlled by the inner water phase pH, as described in Materials and Methods.

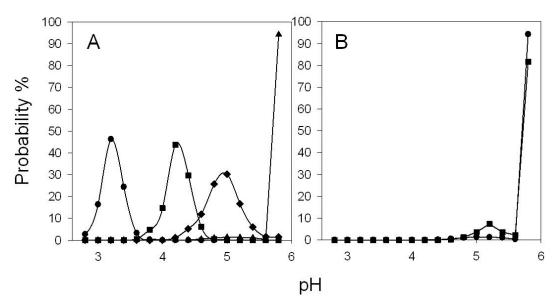


Figure 2.3 µpH distribution kinetics of microspheres encapsulating **(A)** dye and BSA of pH of 3 (•); dye and BSA of pH of 4 (•); dye and BSA of pH of 5 (•); and dye and BSA of pH of 7 (•) **(B)** dye only (•) and dye with BSA (pH of 7)(•) after incubation at 37°C in PBST buffer for 1 day. The µpH was controlled by the inner water phase pH, as described in Materials and Methods.

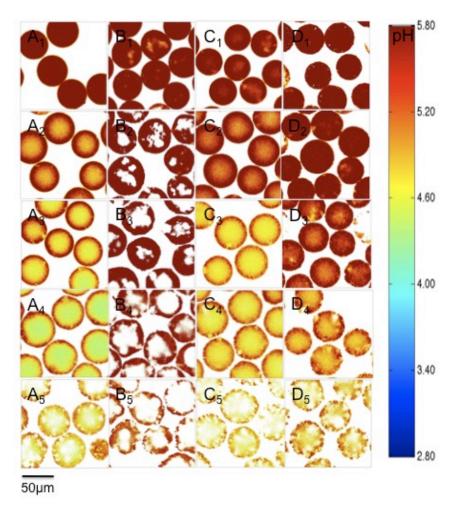


Figure 2.4 Processed confocal images of microsphere formulations during incubation in PBST at 37° C for 4 weeks. Microspheres were prepared from 40% (w/v) PLGA (A), 40% (w/v) PLGA + MgCO₃ (B), 40% (w/v) PLGA + acetate buffer (C) and 30% (w/v) PLGA (D). Images were taken at 1 (A₁-D₁), 7 (A₂-D₂), 14 (A₃-D₃), 21 (A₄-D₄) and 28 (A₅-D₅) days.

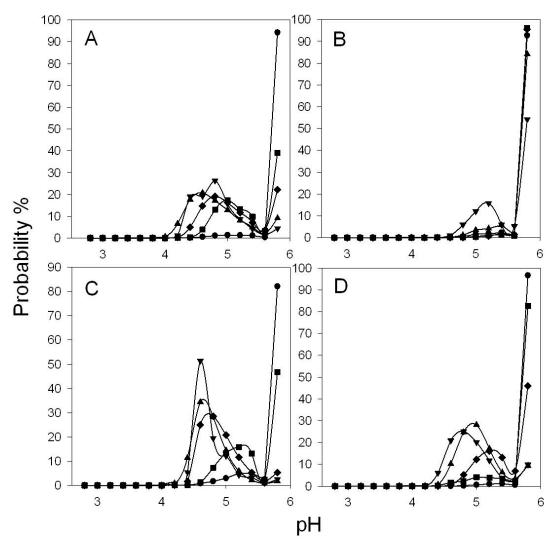


Figure 2.5 μpH distribution kinetics of microsphere formulations during incubation in PBST at 37°C for 1 day (•), 7 days (•), 14 days (•), 21 days (•), and 28 days (\checkmark). Microspheres were prepared from 40% (w/v) PLGA (**A**), 40% (w/v) PLGA + MgCO₃ (**B**), 40% (w/v) PLGA + acetate buffer (**C**) and 30% (w/v) PLGA (**D**).

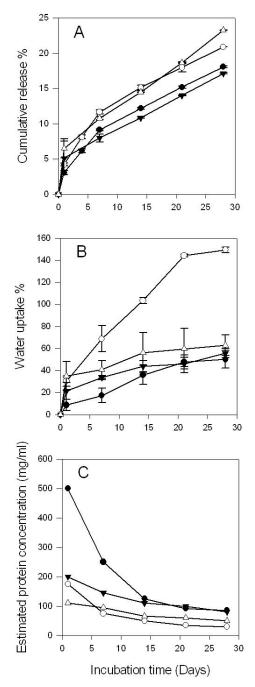


Figure 2.6 Kinetics of protein release **(A)**, water uptake of microspheres **(B)**, and estimated protein concentration in polymer pores **(C)** from PLGA microsphere formulations during incubation in PBST at 37°C for 4 weeks. Microspheres were prepared from 40% (w/v) PLGA (\bullet), 40% (w/v) PLGA + MgCO₃ (\circ), 40% (w/v) PLGA + acetate buffer (\blacktriangledown) and 30% (w/v) PLGA (\triangle). Symbols represent mean \pm SD, n=3 for A and B, SD is not applicable for C because the value is calculated from independent parameters from equation 3.

2.6 Supporting Information

2.6.1 Analysis of deviation in pH distribution from confocal images

To assess errors that may be caused from generating the pixel-by-pixel pH map from confocal images, the following procedures were performed. First, the acquired confocal images (n=8) of standard pH buffer solutions were processed as reported previously [20]. The resulted image is a 512×512 matrix of pixel intensities. Then, by taking the ratio of pixel intensities of processed image at two emission wavelengths (I_{450nm}/I_{520nm}), a ratio matrix was generated. Next, each ratio pixel was converted to a pH according to the standard curve of intensity ratio vs. pH. After that, the probability density function of pH was fit with Gaussian function to obtain mean and standard deviation of pH distribution.

Table S2.1 displayed the variability of pH distribution yielded from this ratiometric measurement of confocal images. Except for the pH approaching detection limit (pH 5.8), in presence or absence of protein, the mean of measured pH was very closed to the actual pH. All pH distributions of standard solutions have a narrow Gaussian distribution with reasonably low standard deviation. Therefore, pH could be accurately mapped within the deviation range of ± 0.2 pH unit over pH from 2.8 to 5.8.

Table S2.1 pH distribution of mapped image from standard pH solution

BSA	0 mg/ml				100 mg/ml					
Concentration										
pH ^a	2.87	3.38	4.15	4.92	5.77	2.88	3.39	4.23	4.96	5.76
Mean pH ^b	2.92	3.36	4.13	4.91	5.60	2.91	3.31	4.20	4.90	5.61
SD b	0.10	0.04	0.04	0.07	0.17	0.04	0.03	0.01	0.07	0.15

^a pH of the standard buffer solutions used to establish the standard curve of intensity ratio *vs.* pH

Dye concentration in the standard solutions was 1.2 mg/ml.

2.6.2 Interpolation of standard curves from estimated protein concentrations

As shown in Figure S2.1, the fluorescence ratio increased with increasing the BSA concentration at constant pH. For BSA concentration from 0 to 250 mg/ml, a linear relationship of ratio *vs.* concentration could be assumed in the range of 0 to 25 mg/ml, 25 to 100 mg/ml, and 100 to 250 mg/ml. BSA concentration higher than 250 mg/ml is rare except after 1-day incubation, in that case, standard curves were fitted from experimental data.

Equations for standard curves of known concentration from experiment:

BSA=0 mg/ml,
$$y_0$$
=-0.0572 x^3 +0.7352 x^2 -2.7529 x +3.4356

BSA=25 mg/ml,
$$y_1$$
=-0.0731 x^3 +0.911 x^2 -3.2801 x +3.91

BSA=100 mg/ml,
$$y_2$$
=-0.0769 x^3 +0.9682 x^2 -3.4631 x +4.1084

BSA=250 mg/ml,
$$y_3$$
=-0.0935 x^3 +1.1604 x^2 -4.0344 x +4.5951

Where x is the pH and y is the fluorescence ratio.

^b mean pH and standard deviation were determined by fitting the pH distribution curve of standard pH solutions with Gaussian distribution function.

If BSA concentration falls to the range of 25 to 100 mg/ml, the slope of linearity between concentration and fluorescence ratio (k) is given by:

$$k = \frac{y_2 - y_1}{100 \, mg / ml - 25 \, mg / ml}$$

$$= \frac{(-0.0769 \, x^3 + 0.9682 \, x^2 - 3.4631 x + 4.1084) - (-0.0731 x^3 + 0.911 x^2 - 3.2801 x + 3.91)}{75 \, mg / ml}$$

$$= \frac{-0.0038 \, x^3 + 0.0572 \, x^2 - 0.183 x + 0.1984}{75 \, mg / ml}$$

For a calculated concentration α mg/ml, the corresponding standard curve can be predicted as:

$$y = y_2 - k \times (100 \, mg/ml - \alpha \, mg/ml)$$

For example, if α =75 mg/ml, then

$$y = (-0.0769x^{3} + 0.9682x^{2} - 3.4631x + 4.1084) - \frac{-0.0038x^{3} + 0.0572x^{2} - 0.183x + 0.1984}{75 mg/ml}$$

$$\times (100 mg/ml - 75 mg/ml)$$

$$= -0.0756x^{3} + 0.9491x^{2} - 3.4021x + 4.0423$$

The predicted curve was well aligned to the experimental curve. (Figure S2.2). Standard curves of concentrations fall within other ranges (0-25 mg/ml, 100-250 mg/ml) could be obtained similarly.

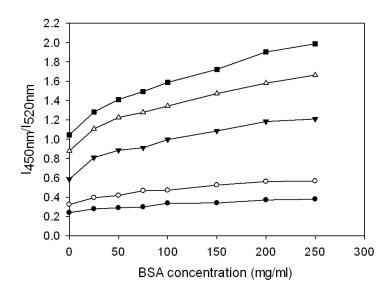


Figure S2.1 The BSA concentration dependency of fluorescence intensity ratio of Lysosensor yellow/blue $^{\circ}$ dextran at pH 2.8 (\bullet), 3.4 (\circ), 4.2 (\blacktriangledown), 4.9 (\triangle) and 5.7 (\blacksquare). The concentration of dye was 1.2 mg/ml.

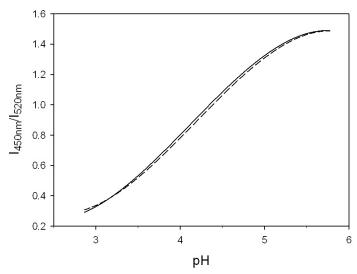


Figure S2.2 The pH sensitivity curves of Lysosensor yellow/blue[®] dextran in presence of 75mg/ml BSA plotted from fitting experiment data (— solid line) and predicted equation (-- dashed line).

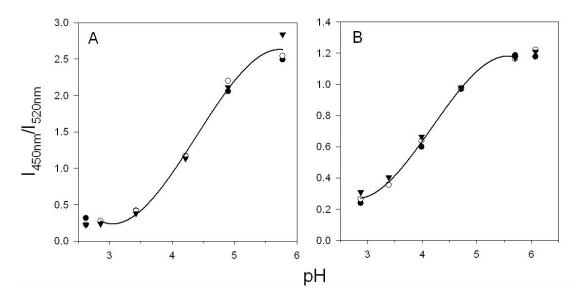


Figure S2.3 The pH sensitivity of confocal pH measurement of Lysosensor yellow/blue $^{\circ}$ dextran at concentration of 0.8 mg/ml ($^{\bullet}$), 1.2 mg/ml ($^{\circ}$), and 2.0 mg/ml ($^{\bullet}$) in presence of 100 mg/ml of BSA **(A)** and lysozyme **(B)**. Lines represent best fits to a third order polynormial function of experimental data.

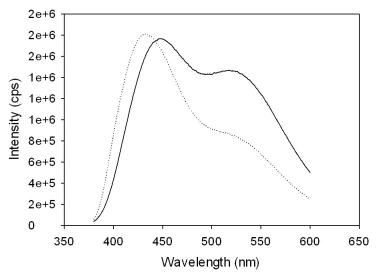


Figure S2.4 Fluorescence spectrum of Lysosensor yellow/blue[®] dextran in the absence (— solid line) and presence of 10 mg/ml BSA (--dashed line) in PBST (pH=7.4). The concentration of dye was 1.0 mg/ml.

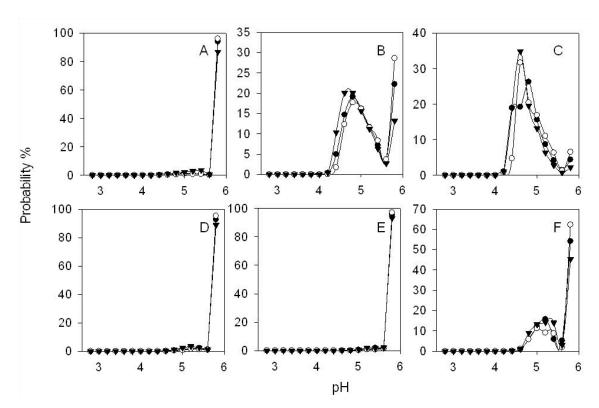


Figure S2.5 Comparison of μpH kinetics in microspheres estimated from protein concentration calculated from measured water uptake (•), 120% of measured water uptake (•), 80% of measured water uptake (\blacktriangledown) at 1 day **(A, D)**, 14 days **(B, E)** and 28 days **(C, F)**. Microspheres were prepared from 40% (w/v) PLGA **(A-C)** and 40% (w/v) PLGA + MgCO₃ **(D-F)**.

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CHAPTER 3

Investigation of the Microclimate pH in Degrading Microspheres of Hydrophilic Poly(D,L-lactide-co-hydroxymethyl glycolide) and PLGA

3.1 Abstract

The microclimate pH (μpH) in biodegradable polymers, such as poly(D,L-lactic-co-glycolic acid) (PLGA) 50/50, commonly falls to deleterious acidic levels during biodegradation, resulting in instability of encapsulated acid-labile molecules. The μpH distribution in microspheres of a more hydrophilic polyester, poly(D,L-lactide-co-hydroxymethyl glycolide) (PLHMGA), was measured and compared to that in PLGA 50/50 of similar molecular weight and degradation time scales. pH mapping in the polymers was performed after incubation under physiological conditions by using a previously validated ratiometric confocal laser scanning microscopic (CLSM) method. Confocal μpH maps revealed that PLHMGA microspheres, regardless of copolymer composition, developed a far less acidic μpH during 4 weeks of incubation compared with microspheres from PLGA. A pH-independent fluorescent probe marker of polymer matrix diffusion of μpH-

controlling water-soluble acid degradation products, bodipy, was observed by CLSM to diffuse ~3-7 fold more rapidly in PLHMGA compared to PLGA microspheres, consistent with much more rapid release of acids observed from the hydrophilic polymer during bioerosion. Hence, PLHMGA microspheres are less susceptible to acidification during degradation as compared to similar PLGA formulations, and therefore, PLHMGA may be more suitable to deliver acid labile molecules such as proteins.

KEY WORDS: microclimate pH; confocal laser scanning microscopy; hydrophilic polyesters; microsphere; pH distribution; poly(lactic-*co*-glycolic acid)

3.2 Introduction

Poly(D,L-lactic-*co*-glycolic acid) (PLGA) is a biodegradable aliphatic polyester that has been widely investigated for controlled delivery of peptides, proteins and vaccine antigens [1-6]. PLGA degrades in aqueous medium via hydrolysis of ester bounds connecting the monomer units in the polymer chain and the final degradation products are lactic and glycolic acid [7]. A major drawback of PLGA systems is the accumulation of acid degradation products inside degrading matrices, which is associated with a drop in microclimate pH (μpH, i.e., the pH in the aqueous pores of the polymer) [8-10] and unwanted instability of acid-labile PLGA-encapsulated species [11, 12]. The use of poorly soluble bases such as magnesium

carbonate, magnesium hydroxide and zinc carbonate as well as blending PEG with the PLGA has been investigated to minimize the drop of pH in protein-loaded PLGA microspheres and to enhance protein stability and release [13-17]. Although coencapsulation of these additives has shown to improve protein stability and release kinetics [18], the release of peptide/protein drugs from PLGA systems is still commonly incomplete and/or difficult to control [19, 20].

Recently, polyesters with functional pendant hydroxyl groups have been developed and showed attractive degradation and release properties for drug delivery purposes [21-25]. Poly(D,L-lactide-co-hydroxymethyl glycolide) (PLHMGA) in particular, showed tailorable degradation kinetics and release of proteins and peptides from PLHMGA microspheres, which was governed by degradation of the microspheres [22, 23, 25], and reduced aggregation of encapsulated lysozyme and formation of less acylated peptide adducts compared to comparable PLGA formulations [22, 25]. The introduction of hydroxyl groups in the backbone of the PLHMGA copolymers makes the polymer more hydrophilic than PLGA and as a consequence, PLHMGA microspheres have a higher water absorbing capacity than their PLGA counterparts. This increase in hydrophilicity might facilitate the more rapid release of the formed acid degrading products into the release medium, which in turn could inhibit the drop of µpH and subsequently improve the stability of encapsulated species. For example, increased upH was observed inside PLGA/PEG blend microspheres [10], which showed increased water uptake due to the

hydrophicility of PEG, and significantly improved stability of encapsulated ovalbumin and BSA [17, 26].

In order to have a better insight into the μpH distribution of PLHMGA microspheres, confocal laser scanning microscopy (CLSM), as a noninvasive tool capable of providing detailed μpH mapping, was used to monitor μpH changes during degradation of the microspheres. In the present study, the effects of PLHMGA copolymer composition and polymer solution concentration used to prepare the microspheres on μpH kinetics during polymer bioerosion were studied and compared to that in PLGA 50/50 formulations. The underlying factors accounting for the μpH differences were also explored.

3.3 Materials and Methods

3.3.1 Materials

Poly(D,L-lactide-*co*-hydroxymethyl glycolide)s with copolymer ratios of 65/35 and 75/25 were synthesized and characterized as described before [23, 24]. Poly(D,L-lactic-co-glycolic acid), end capped, 50/50 with an inherent viscosity (i.v.) of 0.19 dl/g (M_w=19 kg/mol) was generously provided by Alkermes Inc. (Cambridge, MA). Polyvinyl alcohol (PVA, MW 9-10 kDa, 80 % hydrolyzed) was from Polysciences (Warrington, PA). The fluorescent probes, Lysosensor yellow/blue® dextran (MW=10,000 kDa) and BODIPY®, FL (MW 292.1) was purchased from

Invitrogen (Eugene, OR). Unless otherwise stated, all chemicals were analytical grade or higher and used as received.

3.3.2 Preparation of microspheres

Lysosensor yellow/blue® dextran as an acidic pH-sensitive fluorescent probe was encapsulated in PLHMGA and PLGA microspheres by a double emulsion evaporation technique, as described previously [22]. Briefly, 125 µl of dye solution (12 mg/ml) was added to a polymer solution with 350 mg of copolymers/PLGA in 500 µl methylene chloride (35 % w/w). The mixture was homogenized with Tempest IQ² homogenizer (The VirTis Co., Gardiner, NY) at 20,000 rpm for 30 s to form the w/o emulsion. Next, 500 µl of an aqueous PVA solution (1 % w/w) was slowly added to the first emulsion and a w/o/w was formed by homogenizing the mixture at 20,000 rpm for 30s. The prepared w/o/w was slowly transferred into 5 ml of an aqueous PVA solution (0.5 % w/w) and stirred at room temperature for 2 h to extract and evaporate methylene chloride. The formed microspheres were sieved for 20-45 µm size (USA standard test sieve, sieve No.325 and 635, Newark Wire Cloth Co, Newark, NJ) and washed three times with 100 ml double distilled water and thereafter freeze dried on a FreeZone 2.5 Liter Benchtop freeze dry system (Labconco, Kansas City, MO). Microspheres of copolymer PLHMGA 75/25 from three different polymer concentrations (25, 30 and 35 % w/w) were also prepared followed the same procedure.

3.3.3 Microspheres morphology

The morphology of the microspheres was studied using a Hitachi S3200 scanning electron microscope (SEM, Hitachi Ltd., Tokyo, Japan). Approximately 1-2 mg of lyophilized microspheres was evenly sprinkled onto a brass stub with double-adhesive conductive tape. Samples were sputter coated with gold under vacuum using DESK II sputter coater (Denton Vacuum LLC, Moorestown, NJ). The images of microspheres were taken at an excitation voltage of 15.0 kV.

3.3.4 Confocal laser scanning microscopy for microspheres imaging

A ratiometric method was employed as essentially described by Ding *et al.* [10] to map microclimate pH distribution inside microspheres using Zeiss LSM 510-META confocal laser scanning microscope (Carl Zeiss Microimaging, Inc., Thornwood, NY). This instrument was equipped with four laser systems and a Zeiss Axiovert 100M inverted microscope. Lysosensor yellow/blue® dextran was excited at 364 nm by an Enterprise UV laser and two filters (450 nm and 520 nm) were used to build images. For assessing bodipy diffusion in the microspheres, bodipy was excited at 488 nm by an Argon laser and LP 505 filter was used to construct images, as described by Kang *et al.* [27]. Other instrumental parameters were set up as stated elsewhere [10, 27]. All measurements were conducted using a C-Apochromat 63X water immersion objectives lens with numerical aperture of 1.2.

3.3.5 Standard curve of fluorescent intensity ratio vs. pH

Buffers of pH from 2.8 to 5.8 were prepared using combined 0.1 M citric acid solutions and 0.2 M Na₂HPO₄ solutions. Lysosensor yellow/blue® dextran was dissolved in buffer solutions with concentration of 0.8, 1.2, and 2.0 mg/ml. The standard dye solutions were scanned by CLSM. The acquired confocal images were first processed by frame averaging, followed by neighborhood averaging, and applying a median filter as described by Li *et al.* [28] using Image J software (developed by National Institutes of Health and available on the internet at http://rsbweb.nih.gov/ij/) to eliminate signal noise. The standard curve was established by plotting the ratio of mean fluorescent intensities of the dye solutions under two emission wavelengths, 450 nm and 520 nm, versus pH of that solution.

3.3.6 Microclimate pH distribution kinetics inside microspheres

Roughly 15 mg microspheres were suspended into 1 ml phosphate buffer saline (7.74 mM Na₂HPO₄, 2.26 mM NaH₂PO₄, 137 mM NaCl and 3 mM KCl) containing 0.02 % Tween 80 (pH 7.4) (PBST) and incubated in a Glas-Col^{*} vial rotator (Glas-Col LLC, Terre Haute, IN) at 40 rpm at 37 °C. At predetermined time points, a small amount of microspheres was separated for confocal imaging study. The release media was also removed for pH measurement using a Corning 430 pH meter (Corning, NY), followed by replacing with fresh media. The ratio of

fluorescent intensities of each pixel having intensity above the threshold value (indicating the fluorescence from release media) at two emission wavelengths (450 nm and 520 nm) was then calculated and related to a pH from the standard curve. In the processed images, each pixel was converted to a color corresponding to pH. The probability of specific pH value inside microspheres was obtained by dividing the amount of pixels corresponding to a specific pH to the total pixels in the images. Pixel ratios that exceeded the limit of standard curve range referred to a pH of either above 5.8 or below 2.8. In such cases, the percentages were plotted as the boundaries of the µpH distribution curves.

3.3.7 Quantification of water-soluble acids inside PLHMGAs and PLGA

Microspheres (80-90 mg) were incubated in PBST buffer under mild agitation at 320 rpm by a KS 130 basic shaker (IKA® Works Inc., Wilmington, NC) at 37 °C for pre-determined times. After incubation, the microspheres were separated from PBST by a brief centrifugation, followed by washing with double distilled water three times. Then, the microspheres were freeze-dried.

Eighty mg of PLHMGA and PLGA copolymers or dried microspheres were dissolved in 0.5 ml chloroform before adding 3 ml of double distilled water. After a mild vortex mixing, the biphasic solution was left for 10 min and then was centrifuged at 4°C at 4,000 rpm for 5 min. The upper water layer was then quickly

removed. The extraction was repeated for 4 times, and finally the water phases were combined.

The water phase was then titrated with 0.1 M NaOH solution to determine the amount of total water-soluble acids. The electromotive force (EMF) was recorded as a function of the moles of titrant added using a pH meter. The quantity of acid was determined by the total added titrant at the end point, which corresponds to the inflection point of the first derivative of potentiometric titration curve. All measurements were performed in triplicate (n=3).

3.3.8 Determination of bodipy diffusivity in PLHMGA and PLGA microspheres

About 1 mg of PLHMGA and PLGA microspheres were suspended in 1 ml of PBST and incubated at 37°C under mild agitation for pre-determined times. After incubation, the microspheres were separated from PBST by a brief centrifugation, followed by adding 1 ml of bodipy in PBST (5 μ g/ml), which was pre-incubated at 37°C. After incubating the mixture at 37°C for 3 hours, a small amount of microspheres was separated for CLSM observation.

Monitoring bodipy uptake in PLHMGA and PLGA microspheres by CLSM and image and data analysis were carried out following procedures reported by Kang et al. [27]. Briefly, the acquired images were analyzed using Image J software to extract intensity profiles along the diameter. The pixel intensity (I)-position (r) data pairs were then normalized by the surface intensity (I₀) and radius (a) of the

microsphere, respectively. Because of the linear relationship between fluorescence intensity and probe concentration, the normalized intensity (I/I_0)-position (r/a) data pairs were then fit to the following solution to Fick's second law of diffusion to obtain the effective diffusion coefficient (D) of bodipy in the polymer matrix:

$$\frac{C}{C_0} = \frac{1}{r/a} \sum_{n=0}^{\infty} \left(erfc \frac{(2n+1) - r/a}{2\sqrt{Dt/a^2}} - erfc \frac{(2n+1) + r/a}{2\sqrt{Dt/a^2}} \right)$$
 (1)

where t is the diffusion time. The fitting was done according to a least-squares nonlinear regression using n=12 by DataFit software (Oakdale Engineering, Oakdale, PA). All measurements were performed in eight replicates (n=8).

3.4 Results

3.4.1 Characteristics of PLHMGA copolymers

In Table 3.1, the characteristics of the protected poly(D,L-lactic acid-ran-benzyloxymethyl glycolic acid) (PLBMGA) and deprotected PLHMGA are displayed. The molecular weight and thermal behavior of the copolymers were measured and the characteristics are comparable with those reported in previous studies [22, 23, 29].

Table 3.1 Characteristics of PLBMGA and PLHMGA copolymers

polymer	feed ratio D,L ^a /M ^b	copolymer composition (NMR)	M _n (kg/mol)	M _w (kg/mol)	T _g (°C)
PLBMGA	75/25	78/22	16	35	36
	65/35	70/30	24	51	41
PLHMGA	75/25	80/20	13	30	49
	65/35	69/31	22	45	47

 $aD_{,L} = D_{,L}$ -lactide

3.4.2 Preparation of microspheres loaded with an acidic pH sensitive probe

Microspheres were prepared from PLHMGA with different copolymer compositions (65/35 and 75/25) and PLGA 50/50 using a w/o/w double emulsion-solvent evaporation method. Additionally, for PLHMGA 75/25, solutions with different polymer concentrations were employed to prepare microspheres. Since the development of microclimate pH depends on the size of microspheres [10], the microspheres used for confocal microscopy imaging were sieved to yield particles with a similarly narrow size distribution of 20-45 μ m. As can be seen from the scanning electron micrographs (Figure 3.1), all microspheres displayed spherical shape and a non-porous surface.

3.4.3 Microclimate pH distribution inside degrading PLHMGA and PLGA microspheres

^bM = BMMG (benzyloxymethyl methyl glycolide)

The microclimate pH distribution inside degrading PLHMGA and PLGA microspheres during incubation under physiologically conditions for one-month was monitored using a CLSM imaging technique. The encapsulated fluorescent dextran-conjugated probe, Lysosensor yellow/blue® dextran, is sensitive to pH change from 2.8 to 5.8, and this dextran conjugate dye partitions into aqueous pores in polymer similar to encapsulated proteins [8, 10]. A standard curve of the dye correlating its fluorescence intensity ratio at wavelength of 450 nm and 520 nm and pH from 2.8 to 5.8 was established and fitted to a third order polynomial function (r²=0.999) (Figure 3.2). This figure shows that the pH sensitivity of the dye is concentration independent as previously reported [10], which ensures that the standard curve is not affected even though the dye concentration changes during incubation. Some important instrument parameters (e.g., detection gain, pinhole, laser power) were adjusted so that within the concentration range from 0.8 mg/ml to 2.0 mg/ml, the images of dye solutions gave fluorescence intensity from 10 to 255 (units of the instrument). In the microspheres images, any value below 10 was regarded as background and the value exceeding 255 was considered saturated. Intensities within the range of 10 to 255 indicated the existence of entrapped dye. The blank regions in the processed images suggested either dye-free pores or a pure polymer phase.

3.4.3.1 Effect of polymer composition on µpH distribution kinetics

μpH changes were compared in degrading microspheres prepared from PLHMGA of different compositions (65/35 and 75/25) and PLGA 50/50 during incubation in PBST at 37 °C for four weeks, as shown in processed confocal images (Figure 3.3) and µpH distribution curves (Figure 3.4). Within four weeks incubation of microspheres prepared from PLHMGA 65/35, more than 95 % of pixels in the images gave a fluorescence ratio corresponding to pH out of detection range (pH>5.8), indicating these microspheres did not develop any detectable acidity during that time (Figure 3.4A). For PLHMGA 75/25 microspheres, acidity in most aqueous pores with an average pH of 4.8 was observed after one-day incubation, indicative of some acidic impurities in the polymer, although the acidity decreased rapidly with increasing incubation time and almost disappeared by 14 days incubation (Figure 3.4B). By contrast, PLGA microspheres developed µpH as low as 4 during 28 days of study and maintained at an acidic µpH as the degradation of the microspheres continued. The µpH was most acidic after one day of incubation, with around 55 % of pixel domains giving a pH below 5.8. The µpH rose until 2 weeks of incubation before decreasing again. Also note that microspheres made from PLHMGAs were observed to be larger than PLGA ones especially at the later stage of incubation, indicative of the higher water-absorbing capacity of PLHMGAs.

3.4.3.2 Effect of polymer concentration used for preparation of microspheres on $\mu p H$ distribution kinetics

To investigate the effect of polymer concentration used during microsphere preparation on μpH distribution kinetics, microspheres were prepared using methylene chloride solutions of PLHMGA 75/25 of three different polymer concentrations (25, 30 and 35 % w/w). As shown in the processed confocal images (Figure 3.5) and μpH distribution curves (Figure 3.6), increasing the polymer concentration decreased the initial μpH after one day of incubation. As the incubation continued, the acidity inside of the microspheres decreased and disappeared completely after 2 weeks. The μpH is typically found to be more acidic in the center of microspheres than the peripheral regions, due to the relatively shorter diffusion length of formed acid degradation products in polymer regions near the microsphere surface.

3.4.4 Quantification of water-soluble acids in PLHMGAs and PLGA

The water-soluble acids were extracted from PLHMGA and PLGA raw polymers as well as microspheres and quantified, as the total concentration of these species in the polymer pores is predictive of µpH [30]. The water-soluble acids existing in the raw polymer of PLHMGAs and PLGA are attributed to the acid impurities from synthesis and storage, whose quantities were comparable among three polymers (Figure 3.7A). The amount of acids in microspheres, on the other hand, were much less than in the raw polymers, consistent with anticipated diffusion out of acids into the outer water phase during the in-liquid hardening of

microspheres preparation process. However, the quantity of acids extracted from PLHMGA microspheres after incubation in PBST at 37°C was too negligible to be accurately determined, in contrast to PLGA showing the presence of water-soluble acids throughout the incubation period (Figure 3.7A). Since this study was conducted using a different batch of polymer from μpH mapping, it did not show detectable acids in PLHMGA 75/25 microspheres during initial incubation, as expected from the mild acidity recorded by CLSM after 1 day of incubation in Fig. 4B. Nevertheless, it is consistent with the μpH measured using CLSM, which demonstrated non-detectable acidity (pH>5.8) in PLHMGA 65/35 microspheres but an acidic environment in PLGA counterparts during one-month incubation.

3.4.5 pH kinetics in the release media

The pH in the erosion media of microspheres made from different polymers was monitored at the same time point of each μpH mapping, with the buffer being changed weekly. Generally, the pH maintained relatively constant for the release medium of PLGA microspheres. By comparison, the pH declined with the progression of incubation for PLHMGA polymers, with pH of the release medium containing PLHMGA 65/35 lower than that of PLHMGA 75/25 (Figure 3.7B).

3.4.6 Determination of diffusion coefficient of bodipy in PLHMGA and PLGA microspheres

In order to test the hypothesis that acid diffusion out of the polymer was responsible for higher upH in PLHMGA polymer compared to PLGA, the diffusivity of a small hydrophobic fluorescent probe, bodipy inside PLHMGA and PLGA polymer microspheres was determined using CLSM as previously reported [27]. The probe is similar in molecular weight and polymer/water partition coefficient to the trimer of lactic acid [27, 30]. As a result, its diffusion behavior is expected to be a relative indicator of the diffusion of the acid degradation products. The confocal micrographs of bodipy uptake in different polymer microspheres are displayed in Figure 3.8. The dark regions signified the aqueous pores in the microspheres, as the probe concentration in the polymer phase is much higher [27]. The bodipy concentration gradient inside individual microspheres was accurately fit by the solution to Fick's second law of diffusion. (R2 invariably >0.90, see Supporting Information Figure S3.1). The determined diffusion coefficients (D) of bodipy in degraded microspheres are summarized in Table 3.2. The diffusivity of bodipy for each polymer maintained relatively constant during the early state of microsphere incubation (≤3 days). After one week of degradation, the PLHMGA microspheres became porous and the consequent heterogeneous probe distribution caused a poor fit of the confocal images to the solution of the diffusion equation. The diffusion coefficient of bodipy was highest in PLHMGA microspheres made from PLHMGA 65/35, followed by PLHMGA 75/25 microspheres, and lowest in PLGA microspheres.

Table 3.2 Diffusion coefficient of bodipy in degraded microspheres after 3 hours incubation in bodipy solution.

Pre-incubation time (day)	PLHMGA65/35 D (×10 ⁻¹² cm ² /s)	PLHMGA75/25 D (×10 ⁻¹² cm ² /s)	PLGA50/50 D (×10 ⁻¹² cm ² /s)
0	2.6 ± 0.6^{a}	1.6 ± 0.6	0.40 ± 0.05
1	2.8 ± 0.4	1.6 ± 0.3	0.5 ± 0.1
3	2.6 ± 0.8	1.6 ± 0.4	0.46 ± 0.12
7	b	b	0.5 ± 0.10

a: values represent mean ± SD, n=8.

3.5 Discussion

The acidic microenvironment is often regarded as one of most deleterious factors responsible protein instability inside PLGA delivery systems, particularly rapidly degrading PLGA 50/50. The μpH depends on the concentration of watersoluble acids in the aqueous cavities of polymer matrix. According to the equilibrium model for prediction of μpH developed by Ding et~al.~[30], a number of factors may contribute to the development of μpH , including: the amount of acidic impurities, the production rate of water-soluble acids, the liberation rate of water-soluble acids out of polymer, the partition coefficient of acids between polymer and aqueous phases, and the acids' dissociation constant (pKa).

In the present study, we observed that PLHMGA microspheres developed a less acidic microclimate than that of PLGA 50/50 during one-month incubation

b: poor fit of D (R^2 <0.90) due to the increased aqueous pores and a consequent heterogeneous probe distribution.

under physiological conditions as monitored by CLSM. Since the pKa of monomer acids of PLHMGA and PLGA are very close, (3.86, 3.82 and 3.53 for lactic acid, glycolic acid and hydroxymethyl glycolic acid, respectively), the pKa contribution to the µpH differences is negligible. As compared to PLGA, PLHMGAs are more hydrophilic due to the pendant hydroxyl groups on polymer backbone. Because of this more hydrophilic characteristic, monomer acids likely partition more favorably in the water phase of PLHMGA as compared to PLGA. Moreover, since the degradation times of this hydrophilic polyester were shorter than those of PLGA [29], the production of water-soluble acids in PLHMGA is expected to be faster. Therefore, we hypothesized that the upH-determining water-soluble acid degradation products should be released faster from the more hydrated PLHMGA microspheres in order to counteract the above-mentioned unfavorable factors causing higher acidity in the polymer pores. One important piece of evidence was the lower pH observed in the release media of PLHMGA microspheres (Figure 3.7B). Since those acids were quickly released, they did not accumulate in the polymer, leading to an overall less acidic microenvironment in PLHMGAs than in PLGA.

To test our hypothesis, we examined and compared the diffusivity of a small hydrophobic fluorescent probe, bodipy inside PLHMGA and PLGA polymer matrix at the early period of polymer incubation. Bodipy is a good candidate to investigate the transport behavior of small molecules, because i) it is pH-insensitive over the pH

range of 2 to 10; ii) its transport through PLGA is typically limited by transport through the polymer phase; and iii) it displayed a linearity of emission intensityconcentration relationship under our experimental conditions [27]. The effective diffusion coefficient (D) of bodipy in the polymer matrix was determined after incubating degraded microspheres in bodipy solution for 3 hours and then fitting the concentration gradients inside individual microspheres to the solution of Fick's second law of diffusion. This method was confirmed by the fact that i) the bodipy uptake time did not change the value of D (see Supporting Information Table S1), which is consistent with our previous data [27] and ii) the excellent fit of experimental data to equation (1). Since the liberation of water-soluble acids was controlled mainly by polymer diffusion at initial stage of polymer erosion when the pore connectivity in the polymer matrix was still low [27], the higher mobility of bodipy in PLHMGAs should indicate a similar trend in the transport behavior of water-soluble acids (Table 3.2).

The high permeability of PLHMGAs to water-soluble acids relative to PLGA could be explained as follows. First, due to the introduction of pendant hydroxyl group on PLHMGA's backbone, an elevated amount of water is expected to associate with the polymer phase upon microsphere incubation, which could act as a strong plasticizer [31], leading to the relaxation of polymer chains, thereby increasing the diffusivity of degraded acids through polymer phase. Moreover, the more porous

internal structure in PLHMGA microspheres appearing with the progression of incubation, suggested by more isolated dark spots in Figure 3.8, caused a higher effective diffusivity of acidic degradation products through the polymer matrix, as the diffusion across the aqueous pores is several orders of magnitude higher than diffusion in the polymer phase.

Increasing the ratio of hydroxymethyl glycolic acid from 25 % to 35 % increases the hydrophilicity of PLHMGA. Despite the hydrolysis rate of the polymer is raised [23], the diffusion of acid degradation products out of the polymer was further facilitated, causing an even more neutral µpH inside PLHMGA 65/35 microspheres than in PLHMGA 75/25. This was supported by the higher diffusion coefficient of bodipy measured in PLHMGA 65/35 microspheres than in that of PLHMGA 75/25 (Table 3.2) and the more acidic pH in the corresponding release media (Figure 3.7B).

The mild acidity in PLHMGA 75/25 microspheres recorded after one-day incubation described in Figure 3.4 and Figure 3.6 was likely due to the acid impurities existing in the polymer following polymer synthesis, purification and storage. However, as the incubation proceeds, those acids were gradually released, giving rise to a μpH increase and a neutral microenvironment after 2 weeks. Using polymer from a different batch, acidity was not observed during the entire course of incubation, (see Support Information Figure S3.2), suggesting the role of acid

impurities playing in initial low μpH . Increasing the polymer concentration in methylene chloride when fabricating microspheres decreased the initial μpH after one-day incubation. This can be rationalized by the fact that microspheres made from solutions with higher polymer concentration possessed more acidic impurities. Additionally, such microspheres usually have denser structures [32], which impede the liberation of these water-soluble acids at initial incubation. However, because of the higher permeability of PLHMGA copolymer, the effect of polymer concentration had no significant influence on μpH kinetics as the incubation continued. Therefore, the μpH inside of the microspheres all increased to neutral range (above 5.8) consistent with the release of acid impurities as well as degradation products into the incubation medium.

3.6 Conclusion

The microclimate pH (μ pH) inside degrading microspheres prepared from a novel hydroxylated aliphatic polyester; poly(lactic-co-hydroxymethyl glycolic acid) (PLHMGA) was quantitatively mapped by confocal laser scanning microscopy during incubation under physiological conditions. Consistent with previous data of improved stability of PLHMGA encapsulated proteins/peptides, we observed a reduced μ pH in PLHMGA microspheres made from copolymer 65/35 and 75/25 during one-month incubation relative to comparable PLGA formulations. The μ pH inside PLHMGA microspheres made from copolymer 75/25 during the first two

weeks incubation decreased with increasing the polymer concentration. By comparing the pH of release media of PLHMGA and PLGA microspheres and the effective diffusivity of a small fluorescent probe, the data strongly suggests the faster liberation of water-soluble acids in PLHMGA was responsible for its more neutral microenvironment. This study shows that PLHMGA microspheres are potential carriers for controlled delivery of acid-labile biomacromolecules.

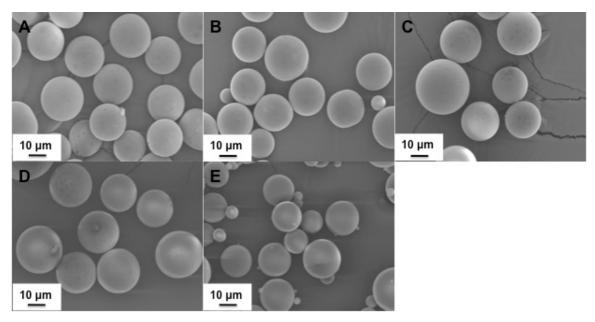


Figure 3.1 Scanning electron micrographs of microspheres prepared from PLHMGA 75/25 with 25 % w/w **(A)**, 30 % w/w **(B)**, 35 % w/w **(C)** polymer solution concentration, and PLHMGA 65/35 **(D)** and PLGA 50/50 **(E)** prepared from a 35 % w/w solution concentration.

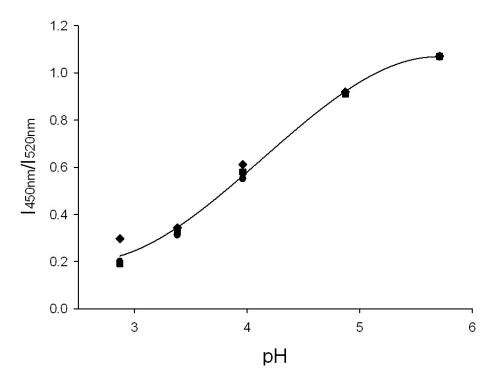


Figure 3.2 The pH sensitivity of confocal measurement of Lysosensor yellow/blue® dextran at concentration of 2 mg/ml (\bullet), 1.2 mg/ml (\bullet) and 0.8 mg/ml (\bullet). The third-order polynomial curve fitting the data was Y = -0.0582 x³+0.7221 x²-2.5676 x+3.0213, where Y = I_{450nm}/I_{520nm} and x = pH, r²=0.999.

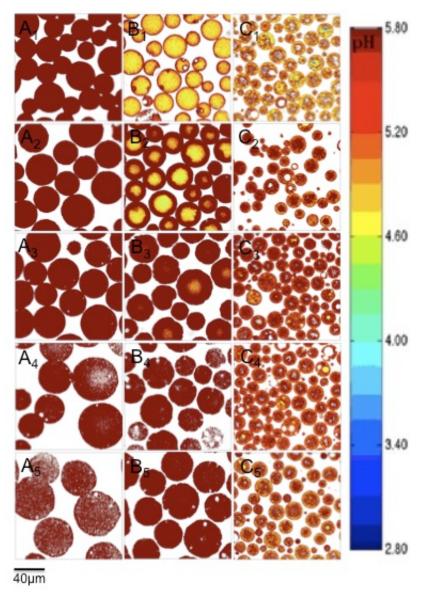


Figure 3.3 Processed confocal images of **(A)** PLHMGA 65/35, **(B)** PLHMGA 75/25 and **(C)** PLGA 50/50 microspheres during incubation in PBST at 37 °C for 4 weeks. Images were taken at 1 (A_1 - C_1), 7 (A_2 - C_2), 14 (A_3 - C_3), 21 (A_4 - C_4) and 28 (A_5 - C_5) days.

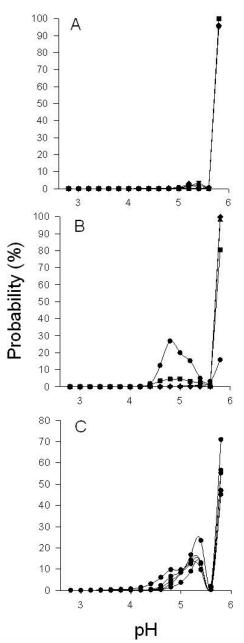


Figure 3.4 The μpH distribution kinetics of microsphere formulations during incubation at 37 $^{\circ}$ C in PBST for 1 day (●), 7 days (■), 14 days (▲), 21 days (▼) and 28 days (♦). Microspheres were prepared from **(A)** PLHMGA 65/35, **(B)** PLHMGA 75/25 and **(C)** PLGA 50/50, and sieved to 20-45μm size for the confocal pH mapping study.

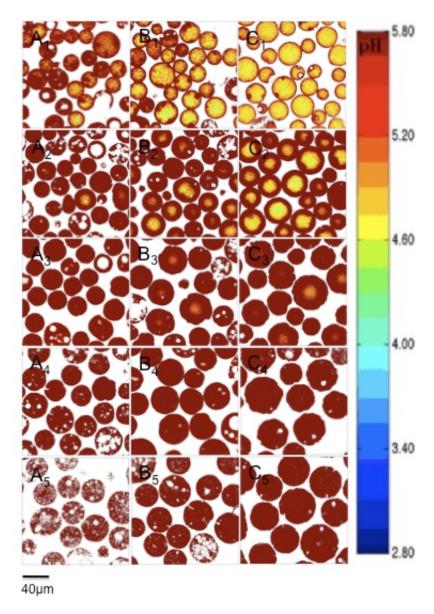


Figure 3.5 Processed confocal images of PLHMGA 75/25 microspheres made from **(A)** 25% w/w **(B)** 30% w/w **(C)** 35% w/w of polymer concentration during incubation in PBST at 37 °C for 4 weeks. Images were taken at 1 (A_1 - C_1), 7 (A_2 - C_2), 14 (A_3 - C_3), 21 (A_4 - C_4), 28 (A_5 - C_5) days.

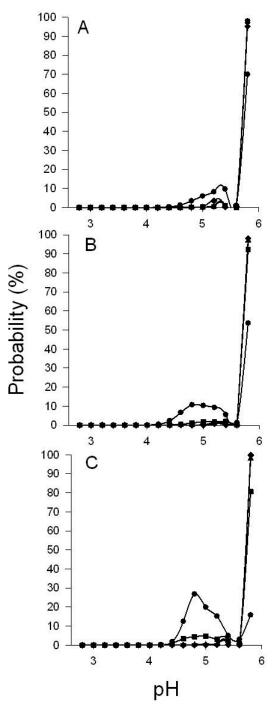


Figure 3.6 The μpH distribution kinetics of microsphere formulations during incubation at 37°C in PBST for 1 day (●), 7 days (■), 14 days (▲), 21 days (▼), and 28 days (♦). Microspheres were prepared from PLHMGA 75/25 of **(A)** 25 % w/w **(B)** 30 % w/w and **(C)** 35 % w/w of polymer concentration, and sieved to 20-45μm size for the confocal pH mapping study.

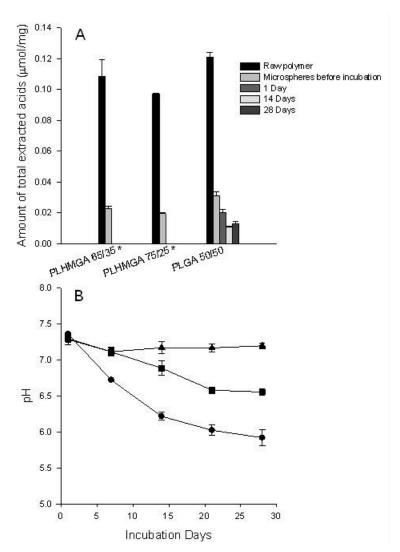


Figure 3.7 Comparison of PLHMGA and PLGA kinetics of total extracted water-soluble acid by titration **(A)** and pH in the erosion medium **(B)** recorded for PLHMGA 65/35 (\bullet), PLHMGA 75/25 (\blacksquare) and PLGA 50/50 (\blacktriangle) microspheres during incubation in PBST at 37 $^{\circ}$ C for 4 weeks. The buffer was changed weekly for both experiments and the pH was measured before each buffer change. Symbols represent mean \pm SD (n=3).

* Acid content in PLHMGAs was below the limit of detection ($<0.002 \mu mol/mg$) (Figure 3.7A) throughout the incubation period.

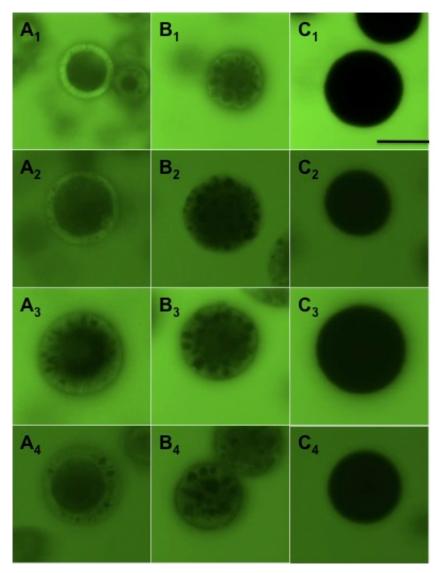


Figure 3.8 Representative CLSM micrographs of the 3-h developed fluorescent intensity gradients of bodipy in **(A)** PLHMGA 6535 **(B)** PLHMGA 7525 and **(C)** PLGA5050 microspheres, which had undergone 0 (A_1 - C_1), 1 (A_2 - C_2), 3 (A_3 - C_3) and 7 (A_4 - C_4) days of degradation under physiological conditions. The scale bar represents 20 μ m.

3.7 Supporting Information

Table S3.1 Diffusion coefficient of bodipy in PLGA 50/50 microspheres ^a after incubating with bodipy solution in PBST at 37°C for various times.

Time of bodipy uptake (h)	3	7	50
D (×10 ⁻¹³ cm ² /s)	3.9±0.6 b	3.6±0.6	3.3±0.3

a: microspheres were not pre-incubated.

b: values represent mean ± SD, n=8.

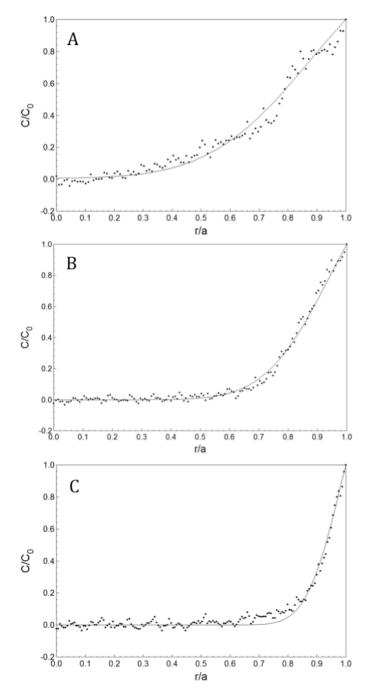


Figure S3.1 Examples of measured and fitted probe concentration profiles inside **(A)** PLHMGA 6535 (R^2 =0.97), **(B)** PLHMGA 7525 (R^2 =0.99) and **(C)** PLGA 5050 microspheres (R^2 =0.98) after 3 hour probe uptake. Microspheres were preincubated in PBST at 37°C for 3 days.

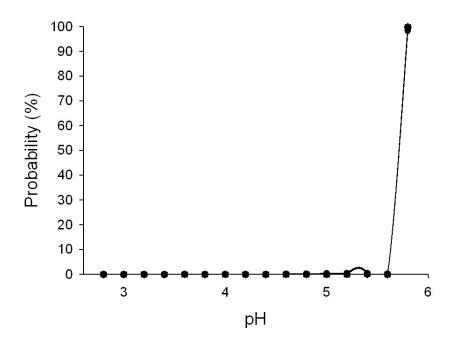


Figure S3.2 µpH distribution kinetics of PLHMGA 75/25 microspheres during incubation at 37° C in PBST for 1 day (\bullet), 7 days (\blacksquare), 14 days (\blacktriangle), 21 days (\blacktriangledown) and 28 days (\spadesuit).

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CHAPTER 4

Simulation of Microclimate pH Distribution and Kinetics inside Degrading PLGA Microspheres

4.1 Abstract

The implications of an acidic microenvironment inside poly(D,L-lactic-co-glycolic acid) PLGA delivery systems on the stability of encapsulated species has gained increasing recognition in recent years. The aim of this chapter is to quantitatively simulate the microclimate pH (µpH) distribution and kinetics inside degrading PLGA microspheres using theoretical tools. A mathematical model based on the production, liberation and partition of the µpH-determining water-soluble acids in spherical geometry was developed. To evaluate the model, fundamental parameters including the size of microspheres, the initial concentration of water-soluble acids in polymer matrix, the production rate constant, and the diffusion coefficient of the water-soluble acids in PLGA microspheres were determined from experiments. This model successfully predicted the µpH development kinetics while showing a small deviation (within 0.5-0.8 pH units) in µpH distribution from

confocal microscopic imaging results, providing us a valuable tool for studying and controlling the μpH in PLGA microsphere formulations.

KEY WORDS microclimate pH distribution; poly(lactic-co-glycolic acid); microspheres; water-soluble acids

4.2 Introduction

Over the past few decades, poly (lactic-co-glycolic acid) (PLGA) is one of the most investigated biodegradable polymers for use in pharmaceutical products and medical devices approved by U.S. Food and Drug Administration [1-6]. Delivery systems (e.g. microspheres, nanoparticles, implant, coated-stent, and others) formulated from this polymer can provide slow and continuous release of bioactive substances over duration of weeks to months during simultaneous biodegradation and erosion of the polymer. However, a major issue associated with this polymer is that an acidic microclimate is commonly formed in the degrading PLGA matrix, which poses a deleterious environment for the stability of encapsulated pH-sensitive biomacromolecules (e.g. peptides and proteins) and drugs [7-11]

The acidic microenvironment has been shown to be a consequence of the accumulation of water-soluble acids in polymer matrix [12], which are present as impurities (e.g., from residual lactide) and during biodegradation. Briefly, upon immersing the PLGA carriers in the physiological buffer, water penetrates into the

polymer matrix rapidly and fills up the pores that are generated during the processing and/or during swelling of the polymer matrix. During the bioerosion period, the microstructure of the polymer can change in a variety of ways with pores often opening and growing in size (e.g., via osmotic forces) or closing and coalescing (e.g., via polymer healing) during the degradation of polymer. Two separate phases are assumed to coexist in the polymer matrix with end-capped PLGAs, which are the polymer phase and aqueous phase. The microclimate pH (µpH) is defined as the pH in the aqueous phase where biomacromolecules reside. In the polymer phase, the degradation of polymer yields acid monomers and oligomers that are either watersoluble or water-insoluble depending on the chain length of the acid. The watersoluble acids will: i) be released out of the polymer matrix by mass transfer, and ii) partition in the aqueous pores, where dissociation occurs producing protons that lower the µpH. Besides the PLGA degradation products, the acidic impurities in the polymer remaining from polymer synthesis, processing and storage can also contribute to the development of acidic upH. Based on such physical-chemical processes, an equilibrium mathematical model has been developed by our group previously and tested to be able to accurately predict the microclimate pH in thin PLGA films based on measured content water-soluble acid [12].

The purpose of this study is to build on the previous mathematical model in order to quantitatively predict the μpH distribution and kinetics in small spherical geometry of injectable PLGA (i.e., microspheres) by considering additionally the

kinetics of water-soluble acid production and mass transfer via diffusion through the polymer matrix. The ultimate goal would be to further our understanding of μpH development and facilitate the formulation design of optimal PLGA systems with controlled μpH for sustained-delivery of stabilized acid-labile therapeutics.

4.3 Theoretical Section

4.3.1 Basic assumptions

The basic assumptions follow our previous treatment [12], as follows: 1. Rapid equilibrium exists between the polymer and aqueous pore solution allowing the maintenance of two separate phases. 2. The penetration of external buffer ions into polymer matrix is minimal, preventing them from interfering with the pore acid-base equilibrium. 3. Negligible water is associated with the polymer phase so that all water-uptake by polymer matrix is localized in aqueous pores. 4. The ionic strength in aqueous pores is relatively low. However, here we make the additional assumptions to accommodate acid production and release from polymer microspheres as follows: 1. After the initial burst release, the water-soluble acids are released from the polymer matrix primarily by diffusion through the polymer matrix, which is limited predominantly by diffusion through the polymer phase. In addition, this diffusivity is assumed to be a constant value. For example, the dye bodipy, which is similar in molecular size (292 Da vs. 234 Da) and polymer/water

partition coefficient (P of 20 vs. 30) to a trimer of lactic acid, had a relatively constant effective diffusion coefficient of in end-capped PLGA 50/50 microspheres over 28 days [12, 13]. 2. We assume that the production rate of each acid follows pseudo-first order kinetics owing to the general acid auto-catalysis [14, 15].

4.3.2 Quantitative treatment

Considering the factors of acid liberation, production and partitioning to aqueous pores contributing to the µpH development, the spatial distribution of the *i*th of water-soluble acids in the polymer matrix as a function of time can be written as the following partial differential equation:

$$\frac{\partial C_{HA_i}^P}{\partial t} = \frac{D_i}{r^2} \frac{\partial}{\partial r} \left(r^2 \frac{\partial C_{HA_i}^P}{\partial r}\right) + k_i C_{HA_i}^P + r_{HA_i} \tag{1}$$

where $C_{HA_i}^P$ is the concentration of the *i*th water-soluble acid in the polymer phase, t is the degrading time of microspheres, D_i is the diffusion coefficient of the *i*th water-soluble acid, r is the radial position of water-soluble acid to the center of microsphere, and k_i is an assumed pseudo-first order production rate constant of the *i*th water-soluble acid in the polymer phase, r_{HA_i} accounts for the loss of *i*th water-soluble acid from the polymer phase due to the partitioning of each water-soluble acid into the aqueous phase.

To obtain the mathematical expression for r_{HA_i} , the relationship between the concentration of ith water-soluble acid in the polymer phase and aqueous pores phase is considered as follows:

$$n_{HA_i} = C_{HA_i}^P V_P + C_{HA_i}^w V_w + C_{A_-}^w V_w$$
 (2)

where n_{HA_i} is the total mole of the *i*th water-soluble acid in the polymer matrix at any time, $C_{HA_i}^w$ is the concentration of *i*th water-soluble acid in the aqueous pores, $C_{A_i^-}^w$ is the concentration of the conjugate base of *i*th water-soluble acid, V_P is the volume of polymer phase, and V_w is the volume of aqueous pores phase.

At steady state, $\frac{\partial n_{HA_i}}{\partial t}$ = 0, therefore making derivative of eq 2 gives

$$\frac{\partial C_{HA_i}^P}{\partial t} V_P + \frac{\partial C_{HA_i}^w}{\partial t} V_w + \frac{\partial C_{A_i}^w}{\partial t} V_w = 0 \quad (3)$$

Due to the negligible water uptake by the polymer phase [16], the volume of polymer and aqueous phase can be related to the mass of dry microspheres (M_p) and hydrated microspheres ($(\phi_w + 1)M_P$):

$$V_P = \frac{M_P}{\rho_P} \tag{4}$$

$$V_{w} = \frac{M_{P}\phi_{w}}{\rho_{w}} \tag{5}$$

where $\phi_{_{w}}$ is water uptake (i.e. ratio of water to dry microsphere masses), and $\rho_{_{P}}$ and $\rho_{_{w}}$ are the densities of the polymer phase and pore water, respectively.

We note that the polymer/water partition coefficient of the ith water-soluble acid (P_i) is defined as follows:

$$P_i = \frac{C_{HA_i}^P}{C_{HA_i}^w} \tag{6}$$

Therefore, the concentration of the *i*th water-soluble acid in the aqueous pores $(C_{HA_i}^w)$ is estimated by:

$$C_{HA_{i}}^{w} = \frac{C_{HA_{i}}^{P}}{P_{i}} \tag{7}$$

Considering the dissociation constant of ith water-soluble acid K_{a_i} gives

$$C_{A_{i}^{-}}^{w} = \frac{C_{HA_{i}}^{w} K_{a_{i}}}{C_{HA_{i}}} \tag{8}$$

Inserting eq 4,5,7,8 into eq 3 gives

$$\frac{\partial C_{HA_i}^P}{\partial t} = -\frac{\partial C_{A_i}^w}{\partial t} / (\frac{\rho_w}{\rho_p \phi_w} + \frac{1}{P_i})$$
 (9)

Inserting eq 7 to 8 gives

$$C_{A_{i}^{-}}^{w} = \frac{C_{HA_{i}}^{P} K_{a_{i}}}{P_{i} C_{H^{+}}}$$
 (10)

Making derivative of eq 10 gives

$$\frac{\partial C_{A_i^-}^w}{\partial t} = \frac{K_{a_i}}{P_i} \frac{\partial (C_{HA_i}^P / C_{H^+})}{\partial t}$$
 (11)

Inserting eq 11 to eq 9 gives $\frac{\partial C_{HA_i}^P}{\partial t}$, which corresponds to the term r_{HA_i}

$$\frac{\partial C_{HA_i}^P}{\partial t} = -\frac{K_{a_i}}{(\frac{\rho_w P_i}{\rho_P \phi_w} + 1)} \frac{\partial (C_{HA_i}^P / C_{H^+})}{\partial t}$$
(12)

Inserting eq 12 to eq 1 as follows:

$$\frac{\partial C_{HA_i}^P}{\partial t} = \frac{D_i}{r^2} \frac{\partial}{\partial r} \left(r^2 \frac{\partial C_{HA_i}^P}{\partial r}\right) + k_i C_{HA_i}^P - \frac{K_{a_i}}{\left(\frac{\rho_w P_i}{\rho_P \phi_w} + 1\right)} \frac{\partial (C_{HA_i}^P / C_{H^+})}{\partial t}$$
(13)

Considering the charge balance in aqueous pores as follows:

$$C_{H^{+}} = \sum C_{A_{i}^{-}}^{w} \tag{14}$$

Inserting eq 7, 8 into eq 14 gives

$$C_{H^{+}} = \sqrt{\sum_{i} \frac{K_{a_{i}}}{P_{i}} C_{HA_{i}}^{P}}$$
 (15)

Inserting eq 15 into eq 13 gives the final partial differential equation describing the distribution of *i*th water-soluble acid in the polymer matrix as a function of time:

$$\frac{\partial C_{HA_i}^P}{\partial t} = \frac{D_i}{r^2} \frac{\partial}{\partial r} \left(r^2 \frac{\partial C_{HA_i}^P}{\partial r}\right) + k_i C_{HA_i}^P - \frac{K_{a_i}}{\left(\frac{\rho_w P_i}{\rho_P \phi_w} + 1\right)} \frac{\partial (C_{HA_i}^P / \sqrt{\sum_i \frac{K_{a_i}}{P_i} C_{HA_i}^P})}{\partial t}$$
(16)

The above equation was integrated following a finite difference method (explicit midpoint) using Matlab software (see Appendix for Matlab code) to give $C_{HA_i}^P$ as a function of r at different time t, which was then substituted into eq 15 to acquire C_{H^+} . Finally, μpH was calculated according to the definition of pH ($\mu pH = -\log(C_{H^+})$) as a function of r at specified time t. In obtaining the solution, it is assumed that at time 0, the water-soluble acids exhibit a uniform distribution within the microsphere ($C_{HA_i}^P(r,0) = C_{0,i}$), where $C_{0,i}$ is the initial concentration of water-soluble acid in the polymer phase; and the concentration of water-soluble acid at the surface of microsphere is 0 ($C_{HA_i}^P(R,t) = 0$), where R is the radius of the microsphere.

In addition, the polymer matrix is assumed a homogenous medium with effective diffusion coefficients.

4.4 Experimental Section

4.4.1 Materials

Poly(D,L.lactic-*co*-glycolic acid), end capped, 50/50 with inherent viscosity (i.v.) of 0.6 dl/g in hexafluoroisopropanol at 25 °C was purchased from Durect Corporation (Birmingham, AL). Polyvinyl alcohol (PVA, 80% hydrolyzed, MW 9-10 kDa) was supplied by Polysciences Inc. (Warrington, PA). The fluorescent probe, Lysosensor yellow/blue® dextran (MW=10 kDa) was purchased from Invitrogen (Eugene, OR). All other reagents were of analytical grade or higher and obtained from commercial suppliers.

4.4.2 Preparation of PLGA microspheres

Blank PLGA microspheres were prepared by using the w/o/w double emulsion-solvent evaporation method. Briefly, 100 µl double distilled water was added to 400 mg/ml PLGA solution in methylene chloride, followed by homogenization at 7,500 rpm for 1 min using a Tempest IQ² homogenizer (The VirTis Co., Gardiner, NY) to generate the first w/o emulsion. Then, 1 ml of PVA solution in water (2% w/w) was quickly added to the emulsion and mixed by

votexing for 20s. The formed w/o/w emulsion was then transferred slowly to 100 ml of PVA solution (0.5% w/w), which was constantly stirred at room temperature for 3 hours to extract and evaporate the organic solvent. Finally, the hardened microspheres were harvested and sieved for size of 45-63 μ m (USA standard test sieve, sieve No.325 and 635, Newark Wire Cloth Co., Newark, NJ). After washing with double-distilled water for three times, the microspheres were freeze-dried on a FreeZone 2.5 Liter Benchtop freeze dry system (Labconco, Kansas City, MO).

For preparation of PLGA microspheres encapsulating the fluorescence probe for μpH mapping, 100 μl of 25 mg/ml Lysosensor yellow/blue® dextran solution in double distilled water was used an internal water phase with all other conditions unchanged.

4.4.3 Mean microsphere size determination

The mean microsphere radius (R) was estimated by averaging the size of 50 microspheres in micrographs taken by a Hitachi S3200 scanning electron microscope (SEM, Hitachi Ltd., Tokyo, Japan). Approximately 1-2 mg of lyophilized microspheres was evenly sprinkled onto a brass stub with double-adhesive conductive tape. Samples were sputter coated with gold under vacuum using DESK II sputter coater (Denton Vacuum LLC, Moorestown, NJ). The excitation voltage was set at 15.0 kV.

4.4.4 Separation and quantification of water-soluble acids

The water-soluble acids recovered from PLGA microspheres were separated and quantified following a chromatographic method developed earlier [17]. The dried acids were first dissolved in acetonitrile, followed by adding 2-fold or greater mole excess of triethylamine (TEA) and bromophenacyl bromide (pBPB) solution in acetonitrile. The reaction was carried out at 50°C in amber glass threaded vials in an oven for 5 h to convert the acids to stable bromophenacyl esters, which was then quantified by reversed phase high performance liquid chromatography (RP-HPLC, Waters Alliance, Midford, MA). The resulting solution was loaded to a 5 µm Symmetry® C-18 column (2.5 cm×4.6 mm i.d.; Waters) with mixture of acetonitrile and water as mobile phase at the flow rate 1.0 ml/min. A linear gradient of 70% to 80% of acetonitrile in 5 min was used, and a UV detector at 254 nm detected the eluent absorbance. Identification of each water-soluble acid was done by comparing a) the retention time of relevant peaks of the analytes with b) the retention times of peaks of the corresponding standards injected separately. Quantification was carried out by integration of peak areas, using the external standardization method. All measurements were performed in triplicates (n=3).

4.4.5 Estimation of initial concentration of water-soluble acids in polymer for µpH simulation

50 mg of blank PLGA microspheres were incubated in 1 ml of phosphate buffer saline (7.74 mM Na₂HPO₄, 2.26 mM NaH₂PO₄, 137 mM NaCl and 3 mM KCl) containing 0.02 % Tween 80 (pH 7.4) (PBST) at 37°C under mild agitation by a KS 130 basic shaker (IKA® Works Inc., Wilmington, NC) for 4 days to remove the acids lost during the initial burst phase. After that, the microspheres were separated from incubation medium by a brief centrifugation, followed by washing with double-distilled water for 5 times to remove the salts before lyophilization.

The water-soluble acids in the PLGA microspheres were recovered by following procedures. Firstly, the dried microspheres were dissolved in 1 ml of chloroform, followed by adding 1 ml of double distilled-water and mixing by mild vortexing for 1 min. Then, the biphasic solution was left for 10 min and then centrifuged at 4°C at 4,000 rpm for 5 min. The upper water layer was then removed. The extraction was repeated for 3 times, and all water phases were combined. Finally, the water phases containing water-soluble acids were concentrated using a vacuum centrifugal concentrator (Labconco, Kansas City, MO) at 50°C using heat mode, and the dried extracts were analyzed by the prederivatization HPLC method as described above. All measurements were performed in triplicates (n=3).

4.4.6 Production kinetics of water-soluble acids in PLGA microspheres

To estimate the rate of production of acids without release, 50 mg of blank PLGA microspheres were placed in a 97% relative humidity environment (created by saturated solution of potassium sulfate) at 37°C. At pre-determined time points, the microspheres were obtained and freeze-dried. The dried microspheres were then dissolved in chloroform and the water-soluble acids were recovered after repeated $CHCl_3/H_2O$ extraction, which were then derivatized and quantified by RP-HPLC as described above. All measurements were performed in triplicate (n=3).

4.4.7 Estimation of the effective diffusion coefficient of water-soluble acids in PLGA microspheres

50 mg of blank PLGA microspheres were first incubated at 97% relative humidity environment at 37°C for one week for degradation. Then, the microspheres were transferred to 1 ml of double-distilled water and incubated at 37°C under mild shaking at 240 rpm. At pre-determined time points over a short time scale, 0.9 ml of release medium was removed, which was then concentrated by vacuum centrifugal concentrator and the composition of water-soluble acids was analyzed using the pre-derivarization HPLC method as described above. Meanwhile, equal amount of fresh water was added to the medium to maintain the sink conditions. All measurements were performed in triplicate (n=3).

The mean cumulative released water-soluble acid vs. time was fitted to Crank's solution for drug release from a monolithic solution in the matrix of

spherical geometry to obtain the effective diffusion coefficient (D) of the watersoluble acid in the polymer matrix, which is described as follows (as has been accomplished for the release of bodipy [13]):

$$\frac{M_t}{M_{\infty}} = 1 - \frac{6}{\pi^2} \sum_{n=1}^{\infty} \frac{1}{n^2} \exp(-Dn^2 \pi^2 t / a^2)$$
 (17)

where M_t/M_∞ is the fraction of released acid relative to the total amount of acid in the microspheres, t is the diffusion time and a is the radius of microsphere. For this study, the average radius of microspheres was used as determined above. The fitting was done according to a least-squares nonlinear regression using n=40 by DataFit software (Oakdale Engineering, Oakdale, PA). Using values larger than n=40 did not change the fitted value of D.

4.4.8 Water-uptake in degrading PLGA microspheres

Microspheres (20–25 mg) were incubated in 1 ml PBST (10mM, pH=7.4) buffer at 37 °C under mild agitation at 320 rpm. At pre-determined time points, the microspheres were collected and the surface water was removed by filtration and the wet weight (W_1) of the microspheres was recorded. The samples then were dried under vacuum to a constant weight and the dry weight (W_2) was recorded.

To correct for the interparticle water, dry microspheres were suspended in PBST at room temperature and rapidly filtered and dried as described above.

Assuming little water uptake by the microparticles between suspension and

filtration, the weight differences between wet and dry particles accounted for the portion of interparticle water (W_i) , as defined by

$$W_{i} = \frac{W_{1}^{'} - W_{2}^{'}}{W_{2}^{'}} \tag{18}$$

Where W_1 ' and W_2 ' are the weights of wet microspheres and dry microspheres after immediate collection (t=0), respectively. The water uptake of microspheres at time $t(W_P(t))$ was estimated by:

$$W_{P}(t) = \frac{W_{1} - W_{2} - W_{2} \times W_{i}}{W_{2}}$$
(19)

Where W_1 and W_2 are the wet and dry microsphere weights at time t. All measurements were performed in triplicate (n=3).

4.4.9 μpH mapping of degrading PLGA microspheres using confocal laser scanning microscopy (CLSM)

The µpH distribution inside degrading PLGA microspheres was monitored using a ratiometric method based on CLSM as reported previously [10, 18, 19]. Briefly, 40 mg of microspheres encapsulating fluorescence dye were incubated in 1 ml of PBST buffer (10mM, pH=7.4) under mild agitation at 320 rpm with the incubation medium replaced once a week. At predetermined time points, a small amount of microspheres were collected and placed under confocal microscope

while focusing at the center to obtain images at two emission wavelengths (450 nm and 520 nm). The setting for confocal microscope and the method for processing the confocal images were the same as reported previously [19]. Meanwhile, standard curve for calibrating the fluorescent ratio (I_{450nm}/I_{520nm}) of dye vs. pH was established following the same procedure [19].

4.5 Results

4.5.1 Microsphere size and morphology

Since the size of microspheres could affect the μpH development, the prepared microspheres were screened using sieves to a narrow size range of 45 to 63 μm . In Figure 4.1, the morphology and size distribution of blank PLGA microspheres are displayed. Microspheres were all well formed with spherical shape and narrowly distributed size. The mean radius of microspheres (R) was estimated by averaging 50 microspheres on the SEM micrographs, giving value of 25 \pm 3 cm (n=50).

4.5.2 Estimation of the initial concentration of water-soluble acids ($C_{0,i}$) in PLGA microspheres for $\mu p H$ simulation

The initial concentration of water-soluble acids for $\mu p H$ prediction should ideally be determined from the concentration of acids in microspheres before

degradation. However, there is an issue with the burst release of encapsulated species from PLGA microspheres present a rapid mass transfer phase, which has significant contributions from pore diffusion before the erosion phase begins. Therefore, microspheres were incubated in phosphate buffer for 4 days to liberate the acids during this initial burst phase, in order to allow more accurate use of the diffusion-limited assumption to the microspheres. The remaining water-soluble acids in the microspheres after initial incubation were then recovered. It should be noted that the amount of water-soluble acids recovered from the experiment includes the sum of HA_i in the polymer phase, HA_i and conjugate base A_i^- in the aqueous pores. Calculations were made to deduct the concentration of HA_i in the polymer phase for the simulation (see Supporting Information), which was treated as the starting concentration of water-soluble acids in the polymer phase at time zero. Consequently, simulation of upH on 7, 14, 21 and 28 days was performed using *t*=3, 10, 17, and 24, respectively.

4.5.3 Determination of the production rate constant of water-soluble acids (k_i) in degrading PLGA microspheres

Water-soluble acid production kinetics were determined by placing the blank PLGA microspheres in a humid environment (97% relative humidity) at 37°C. It was reported that PLGA exposing to water vapor or aqueous medium results in same degradation behavior [16]. More importantly, no loss of acid by transport out of the

polymer occurs under such a condition. At pre-selected time points of erosion, the water-soluble acids microspheres were in recovered after repeated chloroform/water extraction (>95% recovery rate, data not shown) and the composition was measured by pre-derivatization HPLC. By converting acids into UV-sensitive derivatives, the water-soluble degradation products could be well separated and accurately quantified using this method (Figure 4.2). The principal acids in the microspheres during degradation are glycolic acid, lactic acid, and a linear dimer of lactic acid, lactoyllactic acid, by comparing the retention time of relevant peaks of acid extracts with corresponding standards, consistent with the results of degradation products in PLGA films during incubation in phosphate buffer [17].

The production rate of monomer and oligomer acids is assumed to adhere to the pseudo-first order kinetics due to the acid auto-catalytic effect [14]. As shown in Figure 4.3, the natural log of the concentration of each acid in polymer matrix vs. degradation time could be well fitted into a linear relationship, the slope of which gave us the production rate constant k_i .

4.5.4 Estimation of the effective diffusion coefficient of water-soluble acids (D_i) in degraded PLGA microspheres

Water-soluble acids were first accumulated in blank PLGA microspheres degraded in a humid environment for one week where no acid lost was expected.

From a previous study, it was shown that the diffusion coefficient of a small fluorescent molecule, bodipy, having similar molecular weight and polymer/water partition coefficient with trimer of lactic acid, in blank PLGA microspheres did not vary appreciably over the erosion period of 28 days [13]. Therefore, it was expected the influence of polymer erosion time on the diffusivity of water-soluble acids over the same time was insignificant. The release profiles of the accumulated watersoluble acids from such degraded PLGA microspheres were studied during 24 hours incubation in water (Figure 4.4). Within such a small time frame, the acids produced relative to the initial quantity was insignificant, as calculated from the acid production rate (Figure 4.3). The release of acids can be regarded as drug release from a monolithic solution due to the low concentration of water-soluble acids in the PLGA matrix (i.e. below the solubility of water-soluble acid in polymer phase). In addition, the transport of water-soluble acids in the polymer matrix is diffusioncontrolled before extensive polymer erosion occurs. Therefore, Crank's solution for the drug release from spherical geometry can describe the acid diffusion behavior in microspheres. Fitting the fraction of cumulative acid release vs. release time to eq 17 gave the effective diffusion coefficient of acid (D_i) , and excellent fits was observed for each acid (R2=0.9968, 0.9445 and 0.9916 for glycolic acid, lactic acid and lactoyllactic acid respectively, fitting graph in Figure S4.1). Basically, the glycolic acid diffused faster compared with lactic acid in the PLGA polymer matrix, consistent with results of previous studies [21, 22]. This was possibly ascribed to the higher hydrophobicity of lactic acid relative to glycolic acid, which imposes a stronger interaction between the acid and hydrophobic polymer chains, slowing down its effective diffusivity.

4.5.5 Simulation of $\mu p H$ distribution and kinetics in degrading PLGA microspheres

According to eq 16 and its boundary conditions, the parameters for μpH simulation were summarized in Table 1, in which R, $C_{0,i}$, D_i and k_i were obtained from experiments and P_i and K_{a_i} were acquired from literature [12]. At different time points of the simulation, the corresponding value of ϕ_w was used (Figure 4.5). PLGA density was obtained from the manufacturer (ρ_P =1.34 g/mL) and ρ_w =1.0 g/mL.

Table 4.1 Summary of parameters used for simulation of μpH distribution and kinetics in degrading PLGA microspheres.

Parameters	Glycolic acid	Lactic acid	Lactoyllactic acid
R (cm)	0.0025		
C _{0,i} (mol/L)	2.0*10^(-5)	4.0*10^(-5)	4.3*10^(-5)
D _i (cm ² /s)	9.9*10^(-13)	2.9*10^(-13)	1.7*10^(-13)
k _i (day-1)	0.083	0.013	0.079
P _i	6.3	9.5	21
K _{a,i}	3.84	3.82	3.1

In simulation results, the µpH, particularly at later stages of degradation (e.g. 21 days and 28 days), generally increases along the radius of PLGA microspheres (Figure 4.5A). This agrees with our observation from processed confocal images (Figure 4.6) that µpH was higher at the peripheral regions of microspheres and consistent with the anticipation that acids at the center of microspheres are more likely to be accumulated due to the longer diffusion path to be released out [10, 19]. Also, the simulated µpH decreased with the progress of incubation, in agreement with the kinetic trend of µpH development from experiments. However, a small discrepancy still exists between the experimental and predicted results in that the simulated µpH (Figure 4.5B) has a much narrower distribution compared to the measured one (Figure 4.7). In addition, the mean µpH was 0.5 to 0.8 units higher over 4 weeks erosion suggested from experiments compared to simulated results.

4.5.6 Effect of varying model parameters on simulated µpH

Since the simulated μpH was dependent on several parameters, and the ones acquired from experiments (i.e., R, $C_{0,i}$, D_i and k_i) may associate with errors, the effect of varying model parameters on simulated μpH was analyzed and the results are displayed in Figure 4.9. Because microspheres were pre-screened to a narrow size range (45-63 μm), μpH prediction from R of a variation of \pm 25 % in experimental value was performed. For the other parameters, a variation of \pm 100 % and \pm 50 % in experimental value was employed. These results can also be

interpreted as how changing respective parameters could predicatively impact the μpH distribution. Increasing the mean radius of microspheres (R), the initial concentration of water-soluble acids in polymer ($C_{0,i}$), and the production rate of water-soluble acids (k_i) would decrease the predicted μpH , while raising the diffusion coefficient of water-soluble acids (D_i) would lead to its increase. Ω

4.6 Discussion

A preliminary mathematical model incorporating the kinetics of water-soluble acid production, mass transfer via diffusion and partitioning between polymer and aqueous phase was developed to predict the µpH distribution and kinetics in degrading PLGA microspheres. From this model, the kinetics of µpH development was successfully predicted and were shown to be dependent on a number of factors, including the size of the microspheres, the initial concentration of water-soluble acids in polymer, the production rate, and the diffusion rate of water-soluble acids in the polymer matrix. In Figure 4.9, the impact of varying these parameters on the predicted µpH distribution is described. The result corresponded well with experimental results from previous studies on the effect of these factors on µpH by comparing µpH in PLGA microsphere formulations of different size, polymer composition and addition of polyethylene glycol (PEG) [18, 23]. With this model, it becomes easier and efficient to elucidate the influence of these formulation

variables on µpH in PLGA microspheres, thus facilitating the rational design of PLGA delivery systems for acid-labile therapeutics.

However, some differences between predicted and experimental results in present study should be noted. The relative broader μpH distribution in experimental results can be partially explained by the limitation of confocal imaging technique used in this study. It was suggested that the pH of standard pH solutions measured by this technique associate with deviation of \pm 0.2 pH unit [10]. Despite that, the current model still suffers from inadequacy in providing an accurate prediction of μpH distribution inside degrading PLGA microspheres.

Firstly, the model did not account for the factor of initial distribution of water-soluble acids in polymer matrix when predicting μpH development. The narrower μpH distribution reported from simulation can be ascribed to the assumption of uniform distribution of water-soluble acids throughout the polymer phase at beginning of erosion ($C_{HA_i}^P(r,0) = C_{0,i}$). However, it is more likely that a varied distribution of water-soluble acids in the polymer is present at initial period. Although the differential distribution of acids on the surface vs. body of microspheres was considered and measures were taken by incubating microspheres for four days to remove the excessive acids on surface, other factors may also play roles in determining acids distribution. For example, upon immersing microspheres in the incubation medium, water will penetrate into the polymer matrix and consequently lower the glass transition temperature of the polymer [16]. With the

change of polymer from glassy to rubbery state at incubation temperature of 37°C, the polymer chains would become mobile and flexible, leading to the re-distribution of water-soluble acids.

Secondly, the role of pores within polymer matrix on the distribution of water-soluble acids was neglected. The hydration of pores after microspheres imbibing in water could alter the distribution of acids because of the different partition of water-soluble acids between polymer phase and aqueous phase in pores. And the distribution of pores would consequently influence the distribution of water-soluble acids in polymer. From the processed confocal images (Figure 4.6), the uneven distribution of acids could be noted, as there were some neutral pH regions and some "hot spots" with acidic pH after 7 days incubation. The effect of pores on µpH distribution and development should be considered beyond the initial period. As the porosity, pore size and the opening/closing state constantly change during polymer erosion [24], the acid distribution would be further altered. Since μpH is projected to be higher in the peripheral regions, in the case that pores are predominantly distributed in these regions, the µpH would be higher than the current predicted results. Additionally, acids in the open pores that connect to the surface could be rapidly released out during incubation, giving rise to the higher pH (even some neutral pores) in recorded µpH maps.

Another factor the current model may have overlooked is the heterogeneous bulk degradation in PLGA microspheres. The autocatalytic effect of the accumulated

water-soluble acids would lead to accelerated degradation in regions with higher concentration of acids. In addition, the formation of crystallized domains during PLGA degradation as suggested by literatures [25, 26] would impede polymer degradation in these domains. This different rate of acids production throughout microspheres thus may contribute to the relative broad distribution of µpH observed from CLSM results. Nevertheless, a different degradation rate between the bulk polymer and that locates at the interface between polymer and aqueous pores is not likely, since it was suggested that the bulk water (in aqueous pores) and bound water (associated with pure polymer) could result in similar degradation behavior [16].

In this model, an averaged radius (R) of microspheres was employed to simulate the μpH inside PLGA microspheres. However, it is possible that the size distribution of a microsphere population is not centered on that radius. Therefore, it would be more accurate if the mathematical model takes into account the factor of size distribution of microspheres instead of mean radius for μpH prediction.

4.7 Conclusion

In this study, a mathematical model for predicting the μpH distribution and kinetics in degrading PLGA microspheres was developed, which can be used to evaluate the contribution of several formulation variables to μpH development, including the size of microspheres, the initial concentration of water-soluble acids in

polymer matrix, the production rate and the liberation rate of water-soluble acids from polymer during degradation. To test this model, the production rate constant and the diffusion coefficient of three principal water-soluble acids (i.e., glycolic acid, lactic acid, lactoyllactic acid) in degrading PLGA microspheres were experimentally determined for the first time. This model successfully predicted the μpH development kinetics while showing a small deviation (within 0.5-0.8 pH units) from experimental results in μpH distribution. Future studies could involve incorporating factors such as the initial water-soluble acids distribution, the distribution and kinetics of porosity, the heterogeneous degradation of polymer, and the size distribution of microspheres into the model to improve its accuracy for μpH prediction.

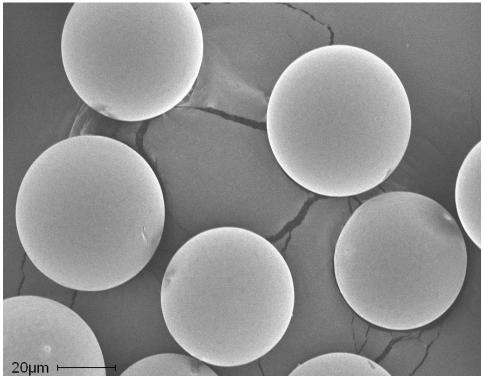


Figure 4.1 Representative SEM micrograph of blank PLGA microspheres that were sieved for the size range of 45 to 63 $\mu m.$

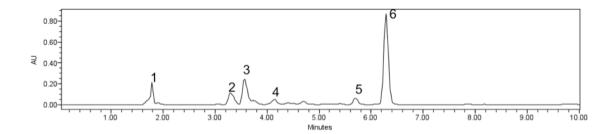


Figure 4.2 HPLC chromatogram of water-soluble PLGA degradation products after conversion to bromophenol esters recovered from microspheres incubated in humid environment at 37°C for 2 weeks. Peak assignment: (1) solvent; (2) glycolic acid; (3) lactic acid; (4) lactoyllactic acid; (5) impurity; and (6) excess reagent of pBPB.

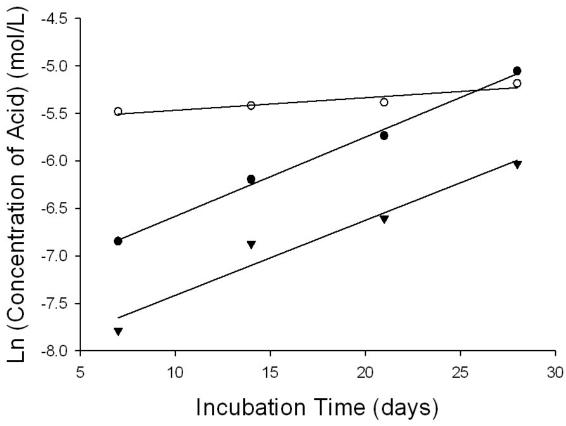


Figure 4.3 The fitted pseudo-first order kinetics of glycolic acid (\bullet), lactic acid (\circ), and lactoyllactic acid (\blacktriangledown) recovered from PLGA microspheres during degradation in humid environment at 37°C for 4 weeks. The fitted equation are y=0.0834x-7.4177, R²=0.995; y=0.0132x-5.6004, R²=0.868 and y=0.0790x-8.2066, R²=0.954, where y is the natural log of the concentration of acid in the polymer and x is the degradation time for glycolic acid, lactic acid and lactoyllactic acid, respectively. Symbols represent mean \pm SD (n=3).

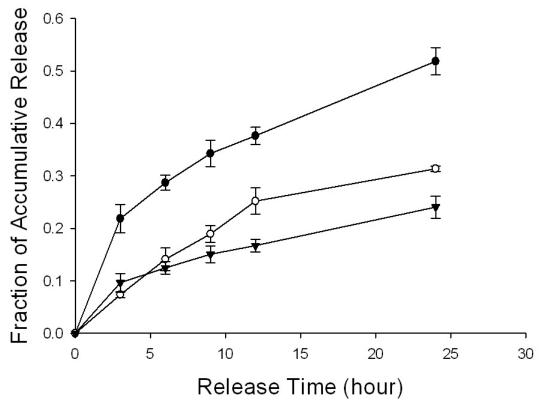


Figure 4.4 Fraction of accumulative release of glycolic acid (\bullet) lactic acid and (\circ) lactoyllactic acid (\blacktriangledown) from pre-degraded PLGA microspheres during 24 hours incubation at 37°C in water. Symbols represent mean \pm SD (n=3).

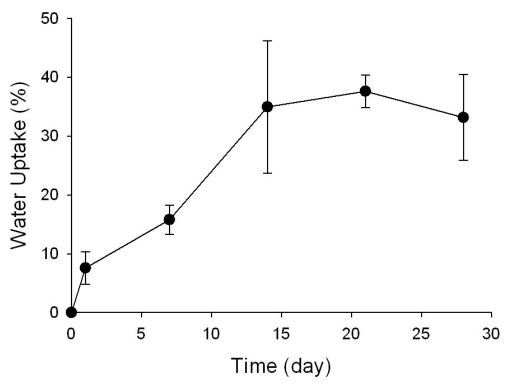
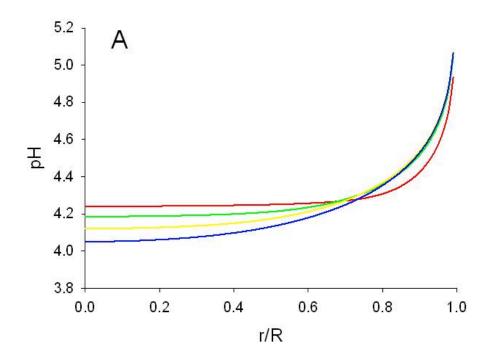


Figure 4.5 Water-uptake kinetics in degrading PLGA microspheres incubated in PBST buffer at 37° C for 4 weeks. Symbols represent mean \pm SD (n=3).



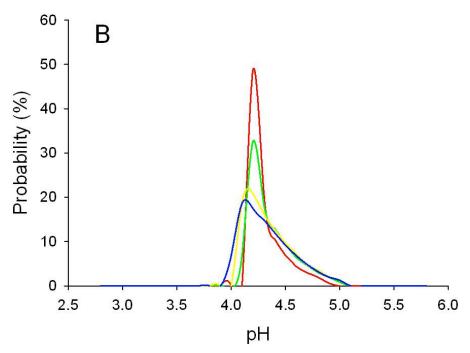


Figure 4.6 Simulated μpH distribution kinetics presented as **(A)** pH *vs.* radial position (r/R); and **(B)** pH *vs.* probability in degrading PLGA microspheres after 7 days (red), 14 days (green), 21 days (yellow) and 28 days (blue) of degradation at 37 °C.

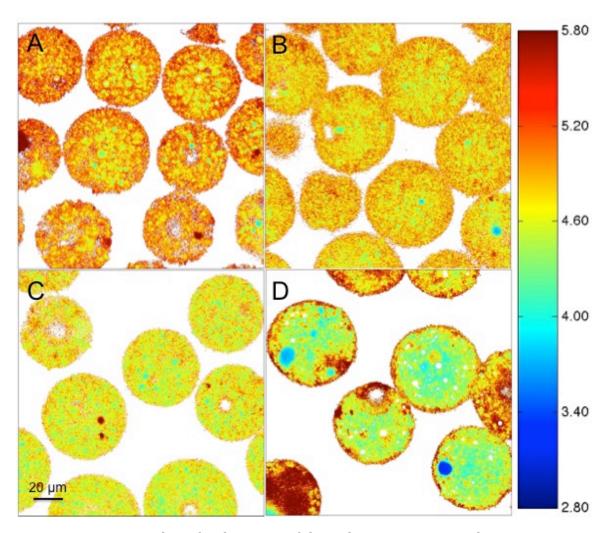


Figure 4.7 Processed confocal images of degrading PLGA microspheres containing fluorescent probe (Lysosensor Yellow/Blue® Dextran) during incubation in PBST buffer at 37 °C for 4 weeks. Images were taken after **(A)** 7; **(B)** 14; **(C)** 21; and **(D)** 28 days incubation.

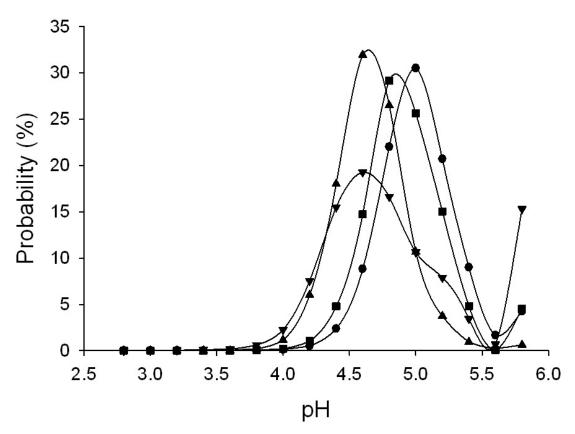


Figure 4.8 µpH distribution kinetics in degrading PLGA microspheres after incubation at 37°C in PBST buffer for 7 days (\bullet), 14 days (\blacksquare), 21 days (\blacktriangle), 28 days (\blacktriangledown) measured by CLSM.

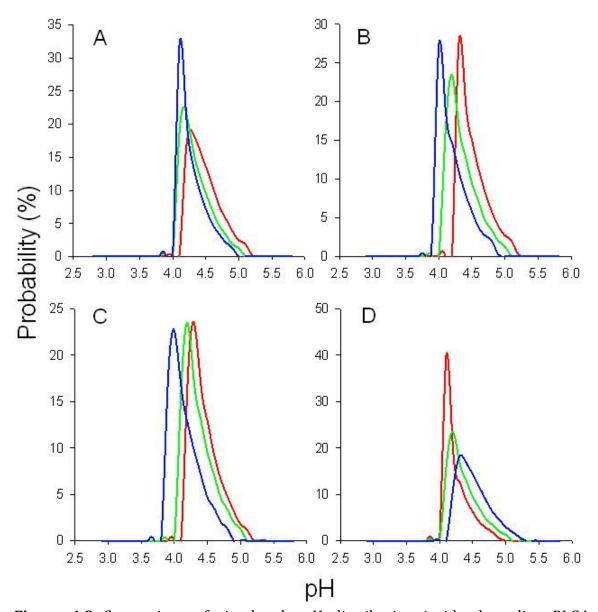


Figure 4.9 Comparison of simulated μpH distribution inside degrading PLGA microspheres after 14 days incubation by varying the value of **(A)** the mean radius of microspheres (R) to 125 % of the experimental value (blue), the experimental value (green) and 75% of the experimental value (red); **(B)** the initial concentration of water-soluble acids ($C_{0,i}$); C) the production rate constant of water-soluble acids (k_i); and D) the diffusion coefficient of water-soluble acids (D_i) to 200 % of the experimental value (blue), the experimental value (green) and 50 % of the experimental value (red).

4.8 Supporting Information

4.8.1 Correction of the concentration of the *i*th water-soluble acid in the polymer phase after 4 days incubation

The moles of the *i*th water-soluble acid (n_{HA_i}) recovered from the polymer matrix equals to the sum of moles of HA_i in the polymer phase and HA_i and its conjugate base A_i^- in the aqueous phase, which can be written as a function of the corresponding molar concentrations ($C_{HA_i}^P$, $C_{HA_i}^w$, $C_{A_i}^w$) and volume of the polymer phase (V_P) and aqueous pore phase (V_W), respectively:

$$n_{HA_i} = C_{HA_i}^P V_P + (C_{HA_i}^w + C_{A_i}^w) V_w$$
 (1)

Consider the dissociation constant of *i*th water-soluble acids gives,

$$C_{A_{i}^{-}}^{w} = \frac{C_{HA_{i}}^{w} K_{a_{i}}}{C_{H^{+}}}$$
 (2)

The volume of polymer phase (V_P) and aqueous pore phase (V_w) can be calculated as follows:

$$V_P = \frac{M_P}{\rho_P} \tag{3}$$

$$V_{w} = \frac{M_{P}\phi_{w}}{\rho_{w}} \tag{4}$$

where M_P is the dry weight of microspheres, $\phi_{\scriptscriptstyle W}$ is the water-uptake by microspheres, $\rho_{\scriptscriptstyle P}$ and $\rho_{\scriptscriptstyle W}$ are the density of PLGA polymer and water, respectively.

Consider the definition of polymer/water partition coefficient of the ith water-soluble acid (P_i) gives:

$$C_{HA_i}^{w} = \frac{C_{HA_i}^{P}}{P_i} \tag{5}$$

Inserting eq 2-5 to eq 1 gives

$$n_{HA_i} = C_{HA_i}^P M_P \left(\frac{1}{\rho_P} + \frac{\phi_w}{\rho_w P_i} (1 + \frac{K_{a_i}}{C_{H^+}}) \right)$$
 (6)

Normalize eq 6 for M_P gives the water-soluble acid content per unit mass of microspheres, which corresponds to the value obtained from experiment

$$\frac{n_{HA_i}}{M_P} = C_{HA_i}^P \left(\frac{1}{\rho_P} + \frac{\phi_w}{\rho_w P_i} (1 + 10^{\mu pH - pK_{a_i}}) \right) \tag{7}$$

After 4 days incubation, the ϕ_w was approximately to be 0.1 from experiment (Figure 4.8). From confocal microscopic imaging results after 4 days degradation, the average μpH was estimated to be 5. K_{a_i} and P_i can be obtained from the literature [12]. PLGA density was obtained from the manufacturer (ρ_P =1.37 g/mL) and ρ_w =1.0 g/mL. Hence, according to eq 7, the ith water-soluble acid concentration in polymer phase $C_{HA_i}^P$ in PLGA microspheres after 4 days incubation can be calculated from experimental value $\frac{n_{HA_i}}{M_P}$

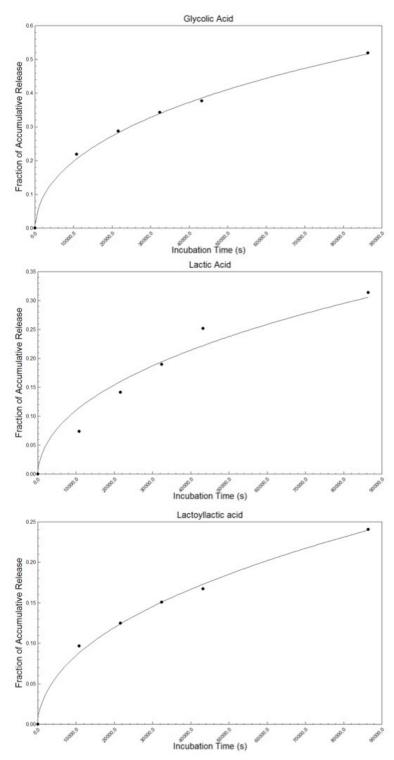


Figure S4.1 Fitting graphs of fraction of accumulative acid release vs. incubation time to eq 17 in pre-degraded PLGA microspheres incubated in water for 24 hours. The correlation coefficient R^2 is 0.9968, 0.9445 and 0.9916 for glycolic acid, lactic acid and lactoyllactic acid, respectively.

Appendix

Matlab code for solving eq 16 to give $C^P_{H\!A_i}$ as a function of r at different time t

```
File 1: RK2.m
```

```
% Compute the concentrations c on a uniform grid with N grid cells r.
% Integrate the reaction-diffusion equations to t=tfinal using RK2 (explicit midpoint
method).
function [r,c]=RK2(N,tfinal)
clear all;
global D k A a
% Define parameters
D=[8.55*10^{-8}) 2.5*10^{-8} 1.47*10^{-8}]; % Diffusion coefficient, cm<sup>2</sup>/day
k=[0.083 0.013 0.079];% production rate constant, day-1
A=[10^{(-3.82)}/47\ 10^{(-3.84)}/70\ 10^{(-3.1)}/154]; \% A=K_a/(\rho_w*P/(\rho_P*\Phi_w)+1)
a=[10^{(-3.82)}/6.3\ 10^{(-3.84)}/9.5\ 10^{(-3.1)}/21]; \% a=K_a/P
% Radius of microsphere (cm)
R=0.0025;
% Grid spacing and grid points
dr=R/N;
r=linspace(0,R,N+1);
% Define vector space for concentrations
c=zeros(3,N+1);
c1=zeros(3,N+1);
% Set initial conditions
for j=1:N
 c(1,j)=.000020;
 c(2,j)=.000040;
 c(3,j)=.000043;
end
```

```
% time step
fudge=.2;
dtmax=fudge*(.5*dr^2/max(D));
nstep=ceil(tfinal/dtmax);
dt=tfinal/nstep;
dt2=dt/2;
% time stepping
for n=1:nstep
 % get c_t
 cprime=getcp(N,dr,r,c);
 % Euler half step
 c1(:,1:N)=c(:,1:N)+dt2*cprime;
 % get c_t at the half step
 cprime=getcp(N,dr,r,c1);
 % advance solution
 c(:,1:N)=c(:,1:N)+dt*cprime;
end
File 2: getcp.m
% compute c_t using the current concentrations given by c
function cprime=getcp(N,dr,r,c)
global D k A a
% matrix and vector space allocation
cprime=zeros(3,N);
B=zeros(3,3);
F=zeros(3,1);
% compute c_t at r=0 (j=1)
C7 = sqrt(a(1)*c(1,j)+a(2)*c(2,j)+a(3)*c(3,j));
```

```
% evaluate diffusion+reaction terms (r=0)
   F(i)=6*D(i)*(c(i,j+1)-c(i,j))/dr^2 + k(i)*c(i,j);
    % row of coefficient matrix
   factor=-A(i)*c(i,j)/(2*C7^3);
    B(i,:)=factor*a;
end
% diagonal contribution to coefficient matrix
B=B+diag(1+A/C7);
% compute c_t
cprime(:,j)=B\setminus F;
% compute c_t for r>0 (j=2:N)
for j=2:N
   C7 = \operatorname{sqrt}(a(1) * c(1,j) + a(2) * c(2,j) + a(3) * c(3,j));
    for i=1:3
        % evaluate diffusion+reaction terms (r>0)
       F(i)=D(i)*((c(i,j+1)-2*c(i,j)+c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1)-c(i,j-1))/dr^2+2*(c(i,j+1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1)-c(i,j-1
1))/(r(j)*dr))+k(i)*c(i,j);
        % row of coefficient matrix
       factor=-A(i)*c(i,j)/(2*C7^3);
        B(i,:)=factor*a;
    end
    % diagonal contribution to coefficient matrix
    B=B+diag(1+A/C7);
    % compute c_t
    cprime(:,j)=B\F;
```

for i=1:3

end

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CHAPTER 5

Future Directions

Confocal laser scanning microscopy proved to be a useful technique to quantitatively investigate the µpH distribution details in biodegradable polymeric microspheres in this dissertation. The µpH mapping was accomplished in the presence of protein, the reduced acidity of a new biodegradable polymer was demonstrated, and the µpH map of degrading microspheres was simulated and found to be close to measured values. However, µpH was only mapped within an acidic range (pH 2.8 to 5.8). For microspheres that developed neutral pH (e.g., PLHMGA microspheres or PLGA microspheres with incorporation of antiacids), employing a fluorescent probe that senses pH changes in the neutral range in the future might provide more detailed information regarding the µpH distribution. For example, it was shown that the ratio of fluorescent intensities of the dextran-SNARF-1® conjugate at emission wavelengths, 580 nm and 640 nm, is responsive to pH change in the neutral range (pH 5.8-8.0) [1].

Future studies might also involve finding specific formulation approaches that control μpH in PLGA microspheres within several specific and narrow pH

ranges. Different formulations strategies could be evaluated for their ability of controlling μpH in degrading PLGA microspheres. Such strategies might include coincorporating basic salts other than magnesium carbonate (e.g., Mg(OH)₂, ZnCO₃), adding plasticizers (e.g., PEG, triethylcitrite) to facilitate the release of water-soluble acids, adding buffer salts (e.g., ammonium acetate) that could control μpH around the maximum buffering capacity of the salts, and encapsulating basic amines (e.g., proton sponge) that are preferentially protonated in the acidic microenvironment.

Mathematical models of higher complexity that address the participation of encapsulated species (excipients and drugs) in the acid-base equilibrium, ionic strength, and water activity in PLGA pores could be developed in order to more accurately predict the µpH in PLGA microsphere formulations containing multiple components. In that way desired function for pH control of future excipients could be simulated. Ultimately, coupling a microclimate pH model with models for protein release would be very powerful, as the addition of excipients to influence both processes can affect each other.

Finally, employing both a) the ability to map μpH in protein-encapsulated biodegradable microspheres according to the method developed in Chapter 2 and b) the acquired knowledge from the μpH simulation, the design of rational PLGA microsphere formulations for clinically relevant therapeutic proteins could be accomplished based on the specific pH requirements of encapsulated proteins.

Reference:

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APPENDIX

Examination of the Influence of Water-soluble Acids in Carboxylic Acid-terminated PLGA on Peptide-PLGA Sorption

A.1 Abstract

The aim of this study is to investigate the effect of water-soluble acids (i.e. monomers and oligomers) present in free acid-terminated poly(D,L-lactic-co-glycolic acid) (PLGA) on peptide sorption. RESOMER® RG502H was incubated as received at 37 °C in PBS or HEPES buffer for 24 hours in the presence or absence of octreotide or leuprolide acetate salts. The kinetics and distribution of water-soluble acids in incubation medium was analyzed by a pre-derivatization HPLC method and PLGA acid number was determined by potentiometric titration with phenolphthalein indicator. Peptide sorption was determined by loss of peptide from solution, monitored by HPLC. The effect of water-soluble acids was determined by comparing the peptide sorption to polymer with and without prior removal of free acids liberated by PLGA. The results showed that in the absence of peptide, the RG502H acid number rapidly decreased after 1 h reaching a quasi-equilibrium (~50% of the initial value) by 3 h irrespective of the buffer used. Over the same

interval, lactic and lactoyllactic acids accumulated in the incubation medium with no detectable glycolic acid, accounting for the acids lost by the polymer. Both peptides sorbed substantially to the RG502H with 25% and 10% loss from octreotide and leuprolide solutions at 1 h, respectively. Interestingly, removal of acids after preincubation of PLGA led to strong inhibition of octreotide sorption whereas weak inhibition for leuprolide solutions. In conclusion, the principal water-soluble acids released initially from RG502H are lactic and lactoyllactic acids, which can have strong or weak effects on peptide sorption. These data may provide further insight into the mechanism of peptide sorption and acylation in carboxylic acid-terminated PLGAs.

KEY WORDS: water-soluble acids; peptide sorption; free acid-terminated PLGA; acylation

A.2 Introduction

Injectable microspheres and implants made of biodegradable poly(D,L-lactic acid) (PLA) and poly (D,L-lactic-co-glycolic acid) (PLGA) are among the most investigated delivery systems for controlled-release of peptides and proteins [1-6]. However, a significant challenge hindering the successful development of such systems is the instability of encapsulated species [7, 8]. The microenvironment inside PLGA aqueous pores where peptide/protein reside often becomes acidic due

to the presence of acidic impurities and the accumulation of degradation products [9-11], which can induce aggregation of encapsulated proteins or trigger unfavorable chemical reactions of peptide/proteins [12, 13]. Apart from that, the presence of moisture inside PLGA matrix [14], and the hydrophobic polymer surface [15] are also recognized deleterious stresses for encapsulated bioactive substances during release and storage.

Recently, acylation has been proposed and proven as an instability mechanism for peptides encapsulated in PLGAs [16-19]. Primary amine groups such as N-terminus and lysine side chain on the peptide can interact with carboxyl groups of PLGA to form acylated peptide impurities, which can potentially result in loss of activity, change of immunogenicity and toxicity. The electrostatically-driven peptide sorption to PLGA is believed to be a critical precursor of acylation [20]. PEGylation of N-terminal of peptide [20] and incorporation of divalent cation salts into PLGA delivery systems [21-23] both demonstrated significant inhibition of peptide acylation via disrupting peptide sorption to PLGA.

Typically, PLGAs from commercial suppliers possess a certain level of residual products from polymer synthesis [24], namely water-soluble monomers and oligomers of glycolic acid and lactic acid, which can also be generated from the degradation of polymer. This study focused on the mechanistic study of the effect of water-soluble acids present in free-acid terminated PLGA on peptide sorption to the polymer. An understanding of the role of water-soluble acids on peptide-PLGA

interaction will shed light on the understanding of peptide sorption mechanism and the design of formulations with stabilized therapeutic peptide against acylation.

A.3 Materials and Methods

A.3.1 Materials

Octreotide acetate was provided by Novatis Pharmaceutical Corp. (Basel, Switzerland). Leuprolide acetate was purchased from Shanghai Shjnj Modern Pharmaceutical Technology Co., Ltd (Shanghai, China). PLGA 50/50 (Resomer® RG502H, i.v. 0.2 dl/g) was purchased from Boehringer Ingelheim (Ingelheim, Germany). (Hydroxyethyl)-piperazine-(ethanesulfonic acid) (HEPES) and sodium phosphate was purchased from Sigma-Aldrich Chemical Co. (St. Louis, MO). All other reagents were of analytical grade or higher and obtained from commercial suppliers.

A.3.2 Kinetics of acid content of RG502H during incubation

100 mg of free-acid terminated RG502H was incubated as received in 10 ml phosphate buffer saline (7.74 mM Na₂HPO₄, 2.26 mM NaH₂PO₄, 137 mM NaCl and 3 mM KCl) (PBS, 10mM, pH 7.4) and HEPES buffer (10mM, pH 7,4) respectively at 37 °C under continuous agitation at 320 rpm by a KS 130 basic shaker (IKA® Works Inc., Wilmington, NC). At pre-determined times, the sample was centrifuged and the

supernatant was removed. The remaining polymer was washed with double distilled water for three times, before freeze-dried on a FreeZone 2.5 Liter Benchtop freeze dry system (Labconco, Kansas City, MO) for 24 hours.

The acid content of the polymer was determined using a non-aqueous titration method as reported by Zhang *et al.* [22]. Briefly, the dried polymer was dissolved in 20 ml organic solvent (acetone/THF=1:1), followed by titration with 0.01M potassium hydroxide in methanol using phenolphthalein methanol solution (0.1 wt %) as an indicator to a stable pink end point. 20 ml of actetone/THF (1:1) mixture without polymer was used as a control.

A.3.3 Kinetics of water-soluble acids in incubation medium

300 mg RG502H polymer was incubated in 30 ml PBS buffer (10 mg/ml) at 37°C under continuous agitation at 320 rpm. At pre-determined times, the incubation medium was separated by centrifugation followed by freeze-drying. The amount of water-soluble acids in the incubation medium after lyophilization was determined using a pre-derivatization method as described by Ding *et al.* [25]. Briefly, the dried acids were dissolved in dimethyl sulfoxide (DMSO) followed by adding 2-fold or greater mole excess of triethylamine (TEA) and bromophenacyl bromide (pBPB) solution in acetonitrile. The reaction was carried out at 50°C in amber glass threaded vials in an oven for 5 h to convert the acids to bromophenacyl esters, which was then quantified by reversed phase high performance liquid

chromatography (RP-HPLC, Waters Alliance, Midford, MA). The resulting solution was loaded to a 5 μ m Symmetry® C-18 column (2.5 cm×4.6 mm i.d.; Waters) with mixture of acetonitrile and water as mobile phase at the flow rate 1.0 ml/min. A linear gradient of 60% to 70% of acetonitrile in 15 min was used, and the eluent absorbance was detected by UV detector at 254 nm.

A.3.4 Peptide sorption kinetics to RG502H

1 ml of peptide solutions (octreotide acetate or leuprolide acetate) in PBS of concentration of 1 mg/ml was added to 10 mg RG502H and incubated at 37°C under constant agitation at 320 rpm. At predetermined times, the samples were removed from the incubator, centrifuged and the amount of peptide sorbed to polymer was determined by the loss of peptide from solution. The supernatant was analyzed using RP-HPLC for peptide quantification. Specifically, a Nova Pak C-18 column (3.9×150 mm, Waters) was used with mixture of 0.1% TFA in acetonitrile and 0.1% TFA in water as mobile phase. The samples were eluted at a linear gradient of 25% to 35% of acetonitrile in 10 min with the flow rate is 1.0 ml/min and the detection was carried out by UV detector at 280 nm.

A.4 Results and Discussion

A.4.1 Evidence of free water-soluble acids in PLGA RG502H

When raw PLGA RG502H particles were placed in the incubation medium, free water-soluble acid monomers and oligomers would be expected to diffuse out of the polymer and into the release medium. PLGA acid number, as a measure of the acid content directly related to the number of free carboxylic acid functionalities in the polymer, was determined by non-aqueous potentiometric titration of the incubated polymer. As shown in Figure A.1, the PLGA acid number (11.7 \pm 0.1, mean \pm SD, n=3) rapidly decreased after 1 h reaching a quasi-equilibrium (\sim 50% of the initial value) by 3 h irrespective of the buffer used. Since polymer degradation is not expected at such early polymer hydration phase, the results suggest the presence of free water-soluble acids in the polymer. However, it should be noted that the amount of water-soluble acids may be varied depending on the batch of polymer tested due to the discrepancies in manufacturing process and storage conditions. From the same study of another batch of RG502H in phosphate buffer, the acid number only dropped to 96% of the original value after 1 h incubation, exhibiting very small amount free acid residuals. (Supplemental Figure SA.1)

A.4.2 Composition of water-soluble acids in release medium

The composition of free acids released in the incubation medium over the same time interval was analyzed and quantified using a previous reported prederiverization HPLC method. In Figure A.2, only lactic acid and lactoyllactic acid were accumulated in the incubation medium with no detectable glycolic acid during

the first 3 h incubation, consistent with the fact that lactic acid monomers and dimers consisting the major impurities found in raw RG502H originating from synthesizing residuals [24]. Notably, the amount of acids in the release medium during the first 3 h incubation accounted for the lost of acids from the polymer, as seen from the comparable value between the total amount of acids from the polymer (as calculated from the acid number of polymer) and the acids released in the medium and the amount of acids in the raw polymer without incubation (i.e., 107.1% and 107.2% mass balance at 1 and 3 h) (Figure A.3). However, significant glycolic acid appeared after 24 h, and the amount of lactic acid continued to rise, helping to increase the total number of acids from the polymer and release media by $\sim 68\%$ of original. This can be possibly due to the onset of polymer degradation after initial hydration, which produces both lactic acid and glycolic acid.

A.4.3 Sorption kinetics of peptides to RG502H

The sorption kinetics of two model peptides, leuprolide and octreotide, which have different primary structures, net charges and conformational flexibility, to RG502H was firstly investigated. Peptide sorption was determined by the loss of peptide from solution, which was validated by previous study [21]. As shown in Figure A.4, both peptides sorbed substantially to the RG502H with 25% and 10% loss from octreotide and leuprolide solutions at 1 h, respectively. And the sorption continued to increase with respect to increasing incubation time. However, in a

sorption kinetics study of another batch of RG502H, only negligible peptide loss (2%) were recorded from both octrotide and leuprolide solutions after 1 h incubation (Supplemental Figure SA.2).

A.4.4 Effect of water-soluble acids on peptide-PLGA sorption

To study the effect of water-soluble acids in RG502H on peptide sorption, peptides were incubated with polymer for 1 h that underwent pre-incubation and removal of acids released in the incubation medium and compared with that with polymer without pre-incubation. The sorption of octreotide was found to be greatly inhibited to polymer with 1 and 3 h pre-incubation compared with control (e.g., $7.8\%\pm1.8\%$ and $6.2\%\pm0.3\%$ loss vs. $25.0\%\pm2.3\%$ control), while far less significant change was observed for the sorption behavior of leuprolide (e.g., $8.8\%\pm0.6\%$ and $7.4\%\pm0.1\%$ loss vs. $10.0\%\pm0.8\%$ control) (Figure A.5).

The different effect of water-soluble acids on the sorption behavior of different peptides is still unclear. It is hypothesized that peptide sorption to free carboxylic acids in PLGA could occur either to the free water-soluble acids or to carboxylic acid end groups on the backbone of PLGA chains. In the case of octreotide, strong inhibition of peptide sorption by removal of water-soluble acids in polymer suggests the majority of electrostatically interaction occurs between peptide and free water-soluble acids present in RG502H. In contrast, leuprolide mainly interacts with acids on the backbone of polymer as indicated by the weak

inhibition of removal of water-soluble acids. From peptide sorption study of another batch of RG502H, which does not possess significant level of water-soluble acids, the loss of octreotide from solution greatly decreased to only $2\% \pm 0.1\%$ (vs. $25.0\%\pm2.3\%$ control) after 1 h incubation, supporting the hypothesis that free water-soluble acids play an major role in the sorption of octreotide to PLGA.

The peptide-PLGA interaction is a complicated process and not fully understood yet. The binding affinity and kinetics between peptide and PLGA surface depends on a number of factors, including the surface charge, conformation and orientation of peptides and mass-transport rate of peptides to polymer surface, etc [21]. Further studies are needed to elucidate more on the mechanism of peptide interaction with carboxylic acid groups of PLGA.

Due to the crucial role of peptide sorption to PLGA as the initial step in peptide acylation pathway, the study of the mechanism of peptide sorption, specifically the role of free water-soluble acids on peptide-PLGA interaction, may facilitate the design of rational formulation strategies that minimize peptide (e.g., octreotide) acylation in PLGA delivery systems. For instance, by controlling the level of water-soluble acid residuals in PLGA before formulation, by adding excipients that promote the diffusion of detrimental acids (e.g., PEG) [26, 27] or by incorporation antacids reagents (e.g., Mg(OH)₂, MgCO₃) to counteract acids produced during degradation [28, 29], the sorption of peptide to the polymer could

be significantly inhibited, thereby peptides could be stabilized against acylation during release.

A.5 Conclusions

Depending on manufacture and storage, the amount of water-soluble acids in carboxylic acid-terminated PLGAs may vary. The principal water-soluble acids released initially from RG502H are lactic and lactoyllactic acids, which can have strong or weak effects on peptide sorption. These data may provide further insight into the mechanism of peptide sorption and acylation in carboxylic acid-terminated PLGA carriers and promote the development of formulation strategies for delivery of stabilized peptides.

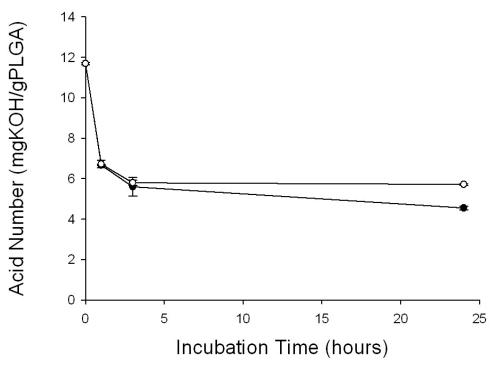


Figure A.1 Kinetics of acid number of RG502H during incubation in PBS (\circ) (10 mM, pH 7.4) and HEPES (\bullet) buffer (10mM, pH 7.4) at 37°C for 24 hours.

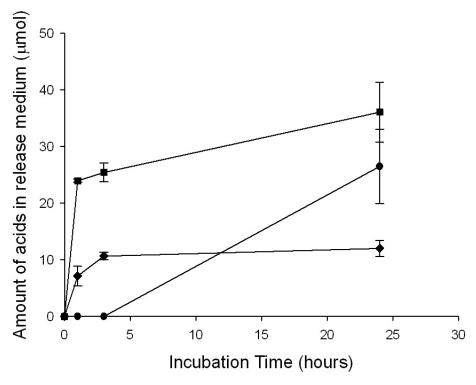


Figure A.2 Kinetics of glycolic acid (●), lactic acid (■), and lactoyllactic acid (♦) in incubation medium (PBS, 10 mM, pH=7.4) released from 300mg RG502H under incubation at 37 °C for 24 hours.

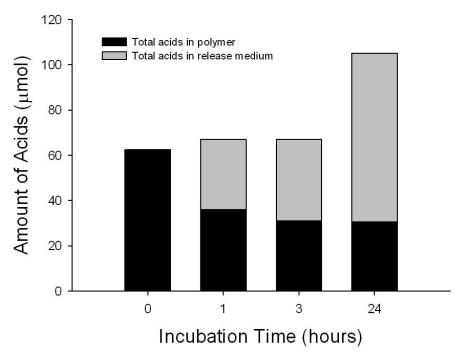


Figure A.3 Kinetics of total carboxylic acids from 300mg RG502H incubated in PBS buffer (10 mM, pH=7.4) at 37° C for 24 hours calculated from acid number in polymer and the amount of acids in incubation medium quantified by HPLC.

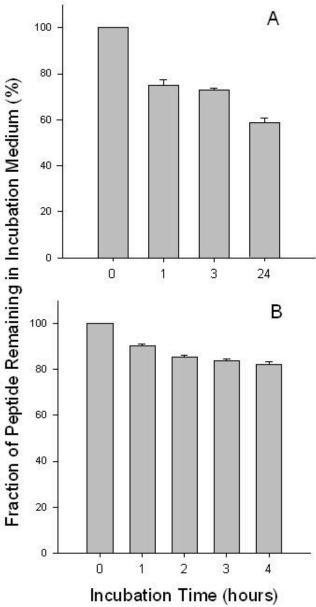


Figure A.4 Sorption kinetics of **(A)** octreotide and **(B)** leuprolide to RG502H incubated in PBS (10 mM, pH=7.4) buffer at $37 \, ^{\circ}$ C.

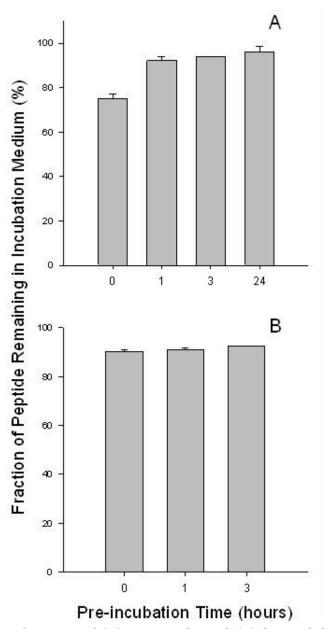


Figure A.5 Sorption kinetics of **(A)** octreotide and **(B)** leuprolide to RG502H after incubation in PBS (10 mM, pH=7,4) buffer for 1 h with and without prior removal of free water-soluble acids liberated by RG502H that were pre-incubated in PBS at 37 $^{\circ}$ C.

Supplemental Figures

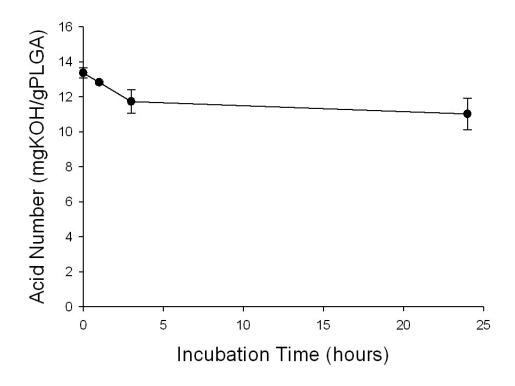


Figure SA.1 Kinetics of acid number of RG502H during incubation in PBS (\circ) (10 mM, pH 7.4) buffer at 37°C for 24 hours.

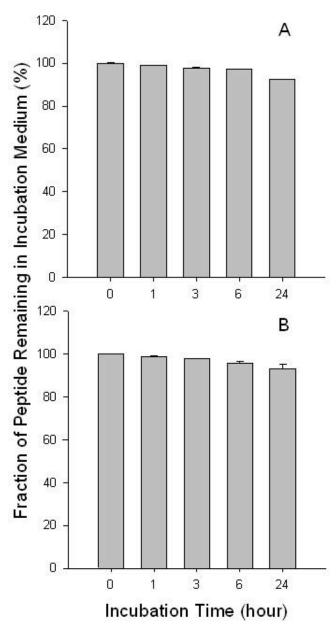


Figure SA.2 Sorption kinetics of (A) octreotide and (B) leuprolide to RG502H incubated in PBS ($10\,\text{mM}$, pH=7.4) buffer at 37 °C for 24 hours.

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