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Author(s): Hesham M. Tawfeek, Sayed H. Khidr, Eman M. Samy, Sayed M. Ahmed, Elsie E. Gaskell, and Gillian A. Hutcheon

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1		Hesham M.	Tawfeek	
2		Sayed H.	Khidr	
3		Eman M.	Samy	
4		Sayed M.	Ahmed	
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Evaluation of biodegradable polyester-co-lactone microparticles for protein delivery

Hesham M. Tawfeek^{1,2}, Sayed H. Khidr², Eman M. Samy², Sayed M. Ahmed², Elsie E. Gaskell¹, and Gillian A. Hutcheon¹

¹School of Pharmacy and Biomolecular Sciences, Liverpool John Moores University, Liverpool, UK and ²Department of Industrial Pharmacy, Faculty of Pharmacy, Assiut University, Assiut, Egypt

Abstract

Poly(glycerol adipate-co- ω -pentadecalactone) (PGA-co-PDL) was previously evaluated for the colloidal delivery of α -chymotrypsin. In this article, the effect of varying polymer molecular weight (M_w) and chemistry on particle size and morphology; encapsulation efficiency; *in vitro* release; and the biological activity of α -chymotrypsin (α -CH) and lysozyme (LS) were investigated. Microparticles were prepared using emulsion solvent evaporation and evaluated by various methods. Altering the M_w or monomer ratio of PGA-co-PDL did not significantly affect the encapsulation efficiency and overall poly(1,3-propanediol adipate-co- ω -pentadecalactone) (PPA-co-PDL) demonstrated the highest encapsulation efficiency. *In vitro* release varied between polymers, and the burst release for α -CH-loaded microparticles was lower when a higher M_w PGA-co-PDL or more hydrophobic PPA-co-PDL was used. The results suggest that, although these co-polyesters could be useful for protein delivery, little difference observed between the different PGA-co-PDL polymers and PPA-co-PDL generally provided a higher encapsulation and slower release of enzyme than the other polymers tested.

Keywords

α -chymotrypsin, biodegradable polyesters, lysozyme, microparticles, PGA-co-PDL, protein delivery

History

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Introduction

Numerous protein and peptide pharmaceuticals such as recombinant human growth hormone, goserelin acetate, leuprolide acetate and recombinant bovine somatropin have already received approval from regulating authorities worldwide¹. However, there are many difficulties associated with delivering biopharmaceutical drugs. The oral route of administration of proteins results in substantial degradation and poor bioavailability², therefore, parenteral delivery is usually preferred. However, proteins often exhibit short half-lives in serum, thus requiring frequent administration to maintain their plasma level³. To prolong the therapeutic level of proteins, controlled release is required and this can be achieved using biodegradable polymers⁴. A range of formulation methods have been utilized to encapsulate proteins in polymeric micro- and nanoparticles, but water-in-oil-in-water (w/o/w) emulsion solvent evaporation is the most frequently used method. Difficulties in the encapsulation of proteins are related to their high molecular weight (M_w), high water solubility and instability upon exposure to formulation conditions⁵. An initial burst release followed by slow, incomplete release of the native protein as a result of protein instability and aggregation has also been recognized as a major problem⁶. Interactions between the protein and the polymer also influence the release profile. These interactions are dependent on protein M_w ; isoelectric point;

amino acid composition; and hydrophobicity, as well as polymer M_w and chemistry¹. Polymer properties such as M_w , copolymer composition and crystallinity can also be tailored to alter polymer degradation and subsequent drug release profiles^{7,8}. For example, an increase in the M_w of Poly(lactic-co-glycolic acid) (PLGA) resulted in longer degradation times and slower release of bovine serum albumin and tetanus toxoid^{9,10}. Bovine serum albumin (BSA) and lysozyme (LS) were encapsulated using two different M_w s of PLGA by (w/o/w) solvent extraction and oil-in-oil (o/o) solvent evaporation systems¹¹. BSA was efficiently encapsulated independently of PLGA M_w , whereas the encapsulation of LS was favored with low M_w PLGA.

Although the choice of polymer is critical, few new polymers have been developed for specific drug delivery applications, and mono- and copolymers of poly(lactic acid) (PLA) and poly(glycolic acid) (PGA) are commonly adopted due to their widespread availability and approval for human use. One alternative is to develop new polymeric delivery systems to release the protein and retain bioactivity over the required target period¹².

A family of biodegradable polyesters with backbone functionality, synthesized via the enzyme catalyzed transesterification of a combination of activated diacids, glycerol and lactone monomers has been designed to overcome the lack of chemical functionality of the commonly used polyesters^{13,14}. The free hydroxyl group from the glycerol monomer allows for the attachment of chemical moieties such as pharmaceutically active drugs, hence introducing the potential for the controlled incorporation and release of desired molecules (drugs, proteins and peptides). In addition, the physical characteristics (hydrophilicity and hydrophobicity) of these polymers can easily be manipulated by varying the

Address for correspondence: Dr Gillian A. Hutcheon, School of Pharmacy and Biomolecular Sciences, Liverpool John Moores University, Byrom Street, Liverpool, L3 3AF, UK. Tel: 0151 2312130. E-mail: G.A.Hutcheon@ljmu.ac.uk

backbone chemistry¹⁵. Previously, poly(glycerol adipate) (PGA) and poly(glycerol adipate-co- ω -pentadecalactone) (PGA-co-PDL) have been investigated for the delivery of dexamethasone phosphate¹⁶ and ibuprofen¹⁷. More recently, PGA-co-PDL has shown promise as a sustained release carrier for pulmonary delivery using the model drug, sodium fluorescein¹⁸. PGA-co-PDL (1:1:1, M_w 30.0 KDa) has also previously been used to prepare α -chymotrypsin (α -CH)-loaded microparticles via the double (w/o/w) emulsion solvent evaporation method^{19,20}. In the initial w/o emulsification step, a lipophilic surfactant is incorporated to aid the emulsification of the aqueous drug solution and the organic phase containing the polymer. Gaskell et al. found that on average 22.1 μ g α -CH per 1 mg PGA-co-PDL was encapsulated, and there was a loss of enzyme bioactivity during encapsulation followed by a further gradual loss upon release¹⁹. The low amount of α -CH encapsulated is typical of these systems due to the diffusion of the protein from the inner to outer aqueous phases during particle formation and upon solvent evaporation. These different previous studies have all utilized a 1:1:1 ratio of monomers, and the M_w of the particular polymers used varied depending upon the M_w achieved during synthesis. Which, given the nature of these reactions, can be difficult to precisely control. It is therefore not known whether the copolymer composition or M_w may influence the characteristics of the particles formed.

Polymer properties such as molecular weight M_w , copolymer composition and crystallinity can be tailored to alter polymer degradation and the consequent drug release profiles as well as the microparticles characteristics. The nature of the protein encapsulated can also affect the particle formation, loading, release and bioactivity profiles²¹.

Therefore this study is an extension of the work presented by Gaskell et al., examining the effect of small changes in polymer M_w and copolymer composition on the encapsulation efficiency, loading, particle size, morphology, *in vitro* release and bioactivity of two different proteins, α -CH (25 kDa) and LS (14 kDa). These enzymes differ in size (LS, 14 kDa, α -CH, 25 kDa), isoelectric point (LS, 11.2, α -CH, 9.1) and stability (LS is more stable than α -CH).

Materials and methods

Materials

Novozyme 435 (a lipase from *Candida antarctica* immobilized on a microporous acrylic resin) was purchased from Bio Catalytics (USA) and stored over P_2O_5 at 5°C prior to use. Glycerol, 1,3-propanediol, ω -pentadecalactone, α -chymotrypsin (type II from bovine pancreas), lysozyme (from chicken egg white), aerosol OT (dioctyl sodium sulphosuccinate), poly(vinyl alcohol) (PVA, M_w 9–10 KDa, 80% hydrolyzed), azocasein, 4-methylumbelliferyl β -D-N,N',N''-triacetylchitotrioside, citric acid, trichloroacetic acid (TCA) and sodium citrate were all obtained from Sigma-Aldrich Chemicals (UK). Dichloromethane and N-[2-hydroxyethyl]piperazine-N'-[2-ethanesulphonic acid] (HEPES) were purchased from BDH (UK). Tetrahydrofuran (THF) was

purchased from Fisher Scientific. Phosphate buffered saline tablets at pH 7.4 were obtained from Oxoid (UK). Divinyl adipate (DVA) was obtained from Fluorochem (UK). A polystyrene standards kit was purchased from Supelco (USA).

Polymer synthesis

The copolymers PGA-co-PDL and PPA-co-PDL were synthesized, processed and characterized using methods adapted from Thompson et al.²² and further described by Gaskell et al.¹⁹ Polymer M_w was varied by controlling the reaction time. Reaction times of 6, 18 and 24 h were used to prepare PGA-co-PDL (1:1:1) with a M_w of 11.4, 26.0 and 39.2 KDa, respectively. The ratio of divinyl adipate (DVA) and glycerol (1:1) to ω -pentadecalactone was varied to produce polymers theoretically containing 1:1:0.5 and 1:1:1.5 of DVA, glycerol and ω -pentadecalactone, respectively. Using the same reaction conditions, PPA-co-PDL with a M_w of 22.0 KDa was synthesized from a 1:1:1 molar ratio of DVA: 1,3-propanediol: ω -pentadecalactone over 24 h.

The polymers were characterized by gel permeation chromatography, GPC (Viscotek TDA Model 300 ran by OmniSEC3 operating software precalibrated with polystyrene standards) and ¹H-NMR spectroscopy (Bruker AVANCE 300 operated via XWIN-NMR v3.5). ¹H-NMR (δ_H CDCl₃, 300 MHz) PGA-co-PDL (1:1:0.5): 1.34 (s, 11H, H-g), 1.65 (m, 8H, H-e, e', h), 2.32 (m, 6H, H-d, d', i), 4.05 (q)-4.18 (m) (6H, H-a, b, c, f), 5.2 (s, H, H-j), PGA-co-PDL (1:1:1): 1.34 (s, 22H, H-g), 1.65 (m, 8H, H-e, e', h), 2.32 (m, 6H, H-d, d', i), 4.05 (q)-4.18 (m) (6H, H-a, b, c, f), 5.2 (s, H, H-j) and PGA-co-PDL (1:1:1.45) 1.30 (s, 32H, H-g), 1.68 (m, 9H, H-e, e', h), 2.32 (m, 6H, H-d, d', i), 4.05 (q)-4.18 (m) (6H, H-a, b, c, f), 5.2 (s, H, H-j). Protons a to j are illustrated in Figure 1.

Particle preparation

The multiple emulsion-solvent evaporation (w/o/w) technique was employed for the encapsulation of α -CH and LS as reported previously¹⁹. Briefly, a 1% (v/v) solution of protein (100 mg mL⁻¹) in phosphate buffered saline (PBS) pH 7.4 was added dropwise to a homogenizing solution of polymer (30 mg mL⁻¹) and aerosol AOT (2 mM) in dichloromethane (15 ml) and emulsified using a IKA yellowline DI 25 basic at 8000 rpm for 30–40 s. This first emulsion was then gradually added to a mixing 1% (w/v) PVA solution (135 ml). This w/o/w emulsion was left to mix with a Silverson L4 RT mixer at 1000 rpm for 3 h to allow for dichloromethane evaporation at 25 °C. The particles obtained were collected by centrifugation (EBA 20, Hettich) at 6000 g for 6 min at room temperature. The supernatant was labeled “wash 1” and retained for further analysis. The microparticles were re-suspended in 120 ml PBS buffer to remove the residual PVA present on the surface of the particles and centrifuged as before. The collected supernatants were labeled “wash 2”. The microparticles were then filtered, vacuum-dried overnight and stored in the fridge. Three batches of each type of particle were prepared.

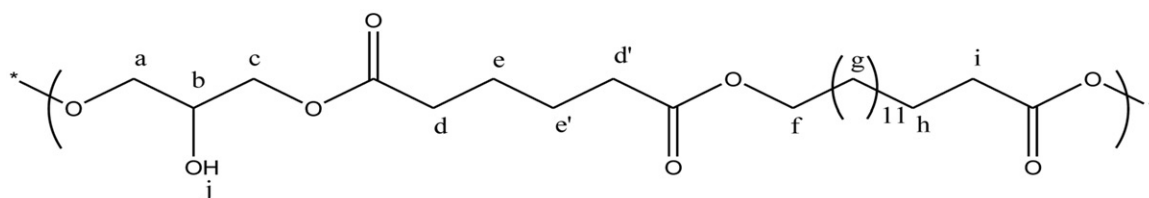


Figure 1. Chemical structure of PGA-co-PDL (1:1:1).

265 Particle characterization

266 The particles were visualized by scanning electron microscopy
267 (FEI – Inspect S Low VAC Scanning Electron Microscope). A
268 suspension of particles in water was deposited on 13 mm
269 aluminum stubs layered with a sticky conductive carbon tab and
270 air dried. An atomic layer of gold was deposited onto the particle
271 containing stubs using an EmiTech K 550X Gold Sputter Coater,
272 25 mA for 3 min.

273 Particle size and size distribution were determined by a laser
274 scattering device (Beckman Coulter LS 13 320, with aqueous
275 liquid module) according to the method described by Pamujula
276 et al.²³ The Fraunhofer method was used to calculate the size
277 distribution of particles in water. The results obtained from
278 measurements of at least three batches of microparticles were
279 described by the volumetric mean diameter of the microparticles
280 (VMD) in micrometers. Equation (1) gives the formula for the
281 span of the volume distribution, which measures the width of the
282 size distribution relative to the median diameter ($d[v,50]$). A more
283 heterogeneous size distribution gives a large span value²⁴.

$$285 \text{Span} = \frac{d[v, 90] - d[v, 10]}{d[v, 50]} \quad (1)$$

288 Powder X-ray diffraction (PXRD) patterns were collected by
289 using a Rigaku Miniflex X-ray diffractometer. Samples were
290 finely ground and packed into an aluminum sample holder.
291 Patterns were collected between 5° and $50^\circ 2\theta$, at increments of
292 $0.02^\circ 2\theta$, scanning speed 2°min^{-1} , voltage 30 KV, current 15 mA
293 using $\text{CuK}\alpha$ (1.54 Å) radiation.

295 Drug loading and encapsulation efficiency

296 The theoretical encapsulation efficiencies and enzyme loading
297 from three different batches of microparticles were calculated
298 from the measurement of the non-encapsulated protein fraction
299 present in the wash samples (Equation 2) and with the assumption
300 that no protein was lost during the preparation and processing of
301 the particles¹⁹. The enzyme loading ($\mu\text{g}/\text{mg}$) was determined
302 using (Equation 3).

$$304 \text{Encapsulation Efficiency (\%)} \\ 305 = \frac{\text{Protection not washed out (mg)}}{\text{Amount of protetin initially added (mg)}} \times 100 \quad (2)$$

$$310 \text{EnzymeLoading} \\ 311 = \frac{\text{Total amount of encapsulated enzyme } (\mu\text{g})}{\text{Total amount of polymer (mg)}} \quad (3)$$

315 In vitro release of enzyme from microparticles

316 Sacrificial sampling was used to observe the release of the
317 enzyme from the particles. In a clean dry 1.5 ml microtube, 10 mg
318 of vacuum-dried particles and 1 ml of phosphate buffer saline pH
319 7.4 at 37°C were placed under sink conditions. The microtubes
320 were then incubated at 37°C in an orbital shaker (IKA KS 130) at
321 250 rpm. Samples were removed at increasing time points over
322 24 h and centrifuged (5 min at 13 500 rpm (17 000 g), accuSpin
323 Micro 17) to collect the particles. The supernatants were retained
324 for analysis by the protein assays described below.

325 The bioactivity of both enzymes was presented as the bioactive
326 fraction of the released enzyme. This was calculated from the
327 ratio of enzyme concentration determined from enzyme activity
328 and the total enzyme concentration as determined by UV
329 spectroscopy using the methods described below²⁵.

Methods for assessing protein content and activity

331 The encapsulation washes (wash 1 and 2) and supernatants from
332 the release studies were analyzed for protein content¹⁹ and
333 activity using the following methods.

UV spectrophotometry

334 To determine the total protein content in a sample, the absorbance
335 was measured at 282 nm for both α -CH and LS, (UV/VIS
336 spectrophotometer Lambda 40, Perkin Elmer, run via the UV
337 WinLab version 2.80.03 software).

Azocasein assay

338 The proteolytic activity of α -CH following release from particles
339 was determined using a chromogenic-based technique as modified
340 by Gaskell et al.¹⁹ Briefly, 50 μL sample, standard or blank and
341 200 μL of azocasein (10 mg/ml), prepared in 25 mM HEPES
342 buffer were incubated for 3 h at 37°C . The reaction was
343 terminated by addition of 750 μL of 0.3 M trichloroacetic acid
344 to precipitate the undigested protein–chromogenic conjugate and
345 the samples were centrifuged for 5 min at 13 500 rpm (17 000 g)
346 (accuSpin Micro 17) to remove the precipitate. Blank samples
347 were prepared using deionized water to determine the amount
348 of azo-dye released nonenzymatically from the substrate.
349 Absorbance of the samples was recorded at 415 nm compared to
350 blank reagent samples using UV/VIS spectrophotometer Lambda
351 40, Perkin Elmer, using the UV WinLab version 2.80.03 software.
352 Three replicates of each sample were obtained and processed.

Muramidase assay

353 The muramidase activity of LS was determined using the method
354 described by Telkov et al.²⁶ Supernatant (760 μL) was incubated
355 with 8 μM 4-Methylumbelliferyl- β -D-N,N',N''-triacetylchitotri-
356 oside in 50 mM citrate buffer, pH 6.0, in the presence of 5 mM
357 MgSO_4 for 3 h at 37°C . The fluorescence intensity was measured
358 using a fluorescence spectrophotometer (Varian Cary Eclipse,
359 operated via the Cary Eclipse Advanced Reads Application
360 version 1.1 (132) software) at an excitation wavelength of 350 nm
361 and an emission wavelength of 450 nm.

Statistical analysis

362 Statistical analysis was performed using student t-paired test.
363 The F-test was used to test the significance of variance. The
364 statistical significance level was set at $p \leq 0.05$.

Results and discussion

365 The aim of this research was to investigate if changes to the M_w
366 and chemistry of PGA-co-PDL would alter the encapsulation,
367 release and bioactivity of α -CH and LS loaded into microparticles
368 fabricated by a w/o/w double emulsion solvent evaporation
369 technique.

Polymer synthesis and characterization

370 The lipase catalyzed ring opening polymerization of an equimolar
371 quantity of DVA, glycerol and ω -pentadecalactone produced
372 PGA-co-PDL (1:1:1) of different M_w s (11.2, 26.0 and 39.2 KDa)
373 by altering the time in contact with the lipase (6, 18 and 24 h,
374 respectively) (Figure 1). A maximum M_w for this type of polymer
375 is usually obtained around 24 h synthesis followed by a subse-
376 quent decrease in M_w as hydrolytic reactions dominate²⁷.
377 This means that the range and difference in M_w s achievable is
378 small and can be difficult to control. The incorporation of 1,3-
379 propanediol in place of the glycerol produced PPA-co-PDL (1:1:1,
380

397 M_w 22.0 KDa) which is more hydrophobic than PGA-co-PDL as
398 it does not have pendant hydroxyl groups.

399 A different set of polymers with a constant 1:1 ratio of DVA
400 and glycerol, but with either 0.5 or 1.5 equivalents of
401 ω -pentadecalactone, was also prepared (1:1:0.5, M_w 23.0 KDa
402 and 1:1:1.45, M_w 34.0 KDa). These polymers should be more
403 (1:1:1.45) and less (1:1:0.5) hydrophobic than PGA-co-PDL
404 (1:1:1) depending on the relative number of hydroxyl groups. It is
405 difficult to control the M_w of these polymers as an increase in the
406 amount of ω -pentadecalactone increases the polymer M_w
407 obtained. This means it can be difficult to directly compare the
408 effect of monomer ratio on polymer and particle properties as
409 there is also a difference in M_w . $^1\text{H-NMR}$ integration patterns
410 were used to confirm that the monomeric content in the polymers
411 were as expected and comparable to that reported in previous
412 work²². The difference in the number of protons at $\delta 1.34$ is
413 indicative of the different proportions of pentadecalactone within
414 the polymer backbone (1:1:0.5 (11H), 1:1:1 (22H) and 1:1:1.45
415 (32H)).

417 Particle characterization

418 Protein-containing and blank particles containing no protein were
419 prepared from each of the different polymers. The mean median
420 of particle diameters (d_{50}) of three separate batches of α -CH- or
421 LS-loaded microparticles and the span values are presented in
422 Table 1.

423 The particle sizes obtained ranged between 9 and 18 μm . The
424 particles prepared from PGA-co-PDL (1:1:0.5) were aggregated
425 so no size data was obtained for this polymer. There was no
426 significant difference observed between the sizes of most of the
427 α -CH- or LS-loaded particles for the different polymers used
428 except with PGA-co-PDL (1:1:1, 39.2 KDa) where significantly
429 larger LS-loaded particles were obtained ($p < 0.05$). Previously, it
430 was reported that the higher the M_w or concentration of polymer
431 in the emulsion, the larger the diameter of the produced
432 particles²⁸. It was not anticipated that any great differences in
433 particle size would be observed because the polymer M_w range
434 studied was small, and the stirring speed, solution concentrations
435 and the organic phase volume were fixed which are the main
436 contributing factors affecting particle size²⁰. Additionally, analysis
437 of the span values (see Table 1) indicates that all
438 microparticles produced had a large size distribution which
439 made it difficult to draw any real trends from the data obtained.

440 The morphology of microparticles is very important as it
441 influences particle degradation and hence can affect the protein
442 release²⁹. Moreover, particle morphology is dependent on the
443 nature, composition and M_w of the polymer^{30,31} as well as the
444 particle formulation conditions²⁰.

445 The SEM images of the external structure of α -CH loaded
446 PGA-co-PDL microparticles prepared from PGA-co-PDL (1:1:1)
447 of different M_w are presented in Figure 2(A–C). Almost spherical
448 microparticles with a slightly irregular shape and a rough ridged
449 surface were observed. A high variability in microparticle size
450

was noted during the SEM analysis which supports the span value
data shown in Table 1. A similar morphology was also observed
with LS-loaded microparticles fabricated from the same polymers
(Figure 2G–I). Hence, changing either the polymer M_w or the
type of protein encapsulated did not alter the particle morphology.

Altering the chemistry did, however, have an effect on particle
morphology. PGA-co-PDL (1:1:0.5) produced small, aggregated,
non-uniform particles (Figure 2D), and increasing the lactone
content within the polymer changed the particle morphology
slightly. With both α -CH- and LS-loaded PGA-co-PDL (1:1:1.45)
particles, some of the particles appeared irregular in shape with
rough surfaces, while the others were spherical with a slightly
smoother surface than those prepared from PGA-co-PDL (1:1:1)
(Figure 2E and J). These smooth particles were more similar to
those obtained from PPA-co-PDL (Figure 2F and K). A similar
morphology to α -CH-loaded microparticles was observed with
the LS-loaded microparticles (Figure 2H–K). Thompson et al.
reported similar morphological characteristics for particles
prepared from PGA-co-PDL and PPA-co-PDL²². Drug-free and
ibuprofen-loaded microspheres¹⁷ produced using PGA-co-PDL
were rough with a ridged morphology, whereas the equivalent
PPA-co-PDL microspheres were smooth.

486 Drug loading and encapsulation efficiency

487 Polymer M_w , degree of hydrophilicity, polymer chemistry,
488 volume of organic phase and enzyme and polymer concentration
489 play an important role in determining the amount of enzyme
490 encapsulated. It was reported that increasing the M_w of poly
491 (ϵ -caprolactone), PLA and PLGA increased the encapsulation
492 efficiency and the mean particle size due to the increased
493 viscosity of the organic phase, which reduces protein diffusion
494 into the external aqueous phase before polymer hardening^{8,32}.
495 Partitioning of the drug from the internal to the external aqueous
496 phase limits the encapsulation efficiency and drug loading in
497 particles prepared via the emulsion solvent evaporation technique.
498 During particle formation, solvent removal and polymer precipi-
499 tation can alter the amount of the protein that partitions into the
500 external aqueous phase³³. It was previously determined that 3 h
501 was the optimum time for PGA-co-PDL protein-containing
502 particle formation as this provided enough time for the solvent
503 to evaporate yet minimized enzyme diffusion to the aqueous
504 phase¹⁹.

505 The encapsulation efficiencies and enzyme loading from three
506 different batches of microparticles prepared using different
507 polymers are presented in Table 2.

508 Increasing the M_w of PGA-co-PDL had no significant effect
509 on either the encapsulation efficiency or α -CH loading ($p > 0.05$).
510 However, a shift in PGA-co-PDL M_w from 11.4 to 39.2 KDa
511 might not be large enough to induce a significant increase in the
512 viscosity of the organic phase, leading to a change in enzyme
513 loading. The degree of crystallinity of the polymer is another
514 important factor affecting drug encapsulation as drugs will tend
515 to be encapsulated in the amorphous region of the polymer³⁴.

518 Table 1. The mean median of particle size and the span values for α -CH- and LS-loaded microparticles prepared via the w/
519 o/w double emulsion solvent evaporation technique. The results are the mean of three different prepared batches \pm S.D.

Polymer type	Mean median of particle size (μm)		Span values	
	CH	LS	CH	LS
PGA-co-PDL (1:1:1, M_w 11.4 KDa)	13.6 \pm 1.4	9.3 \pm 1.4	2.2 \pm 0.2	2.1 \pm 0.2
PGA-co-PDL (1:1:1, M_w 26.0 KDa)	14.4 \pm 2.9	12.2 \pm 0.9	1.9 \pm 0.2	2.3 \pm 0.3
PGA-co-PDL (1:1:1, M_w 39.2 KDa)	13.8 \pm 2.9	17.5 \pm 0.6	2.8 \pm 1.2	3.3 \pm 0.6
PGA-co-PDL (1:1:1.45, M_w 34.0 KDa)	9.6 \pm 0.81	15.1 \pm 1.0	1.6 \pm 0.3	2.1 \pm 0.2
PPA-co-PDL (1:1:1, M_w 22.0 KDa)	10.0 \pm 1.2	14.4 \pm 1.5	2.2 \pm 0.5	2.6 \pm 0.4

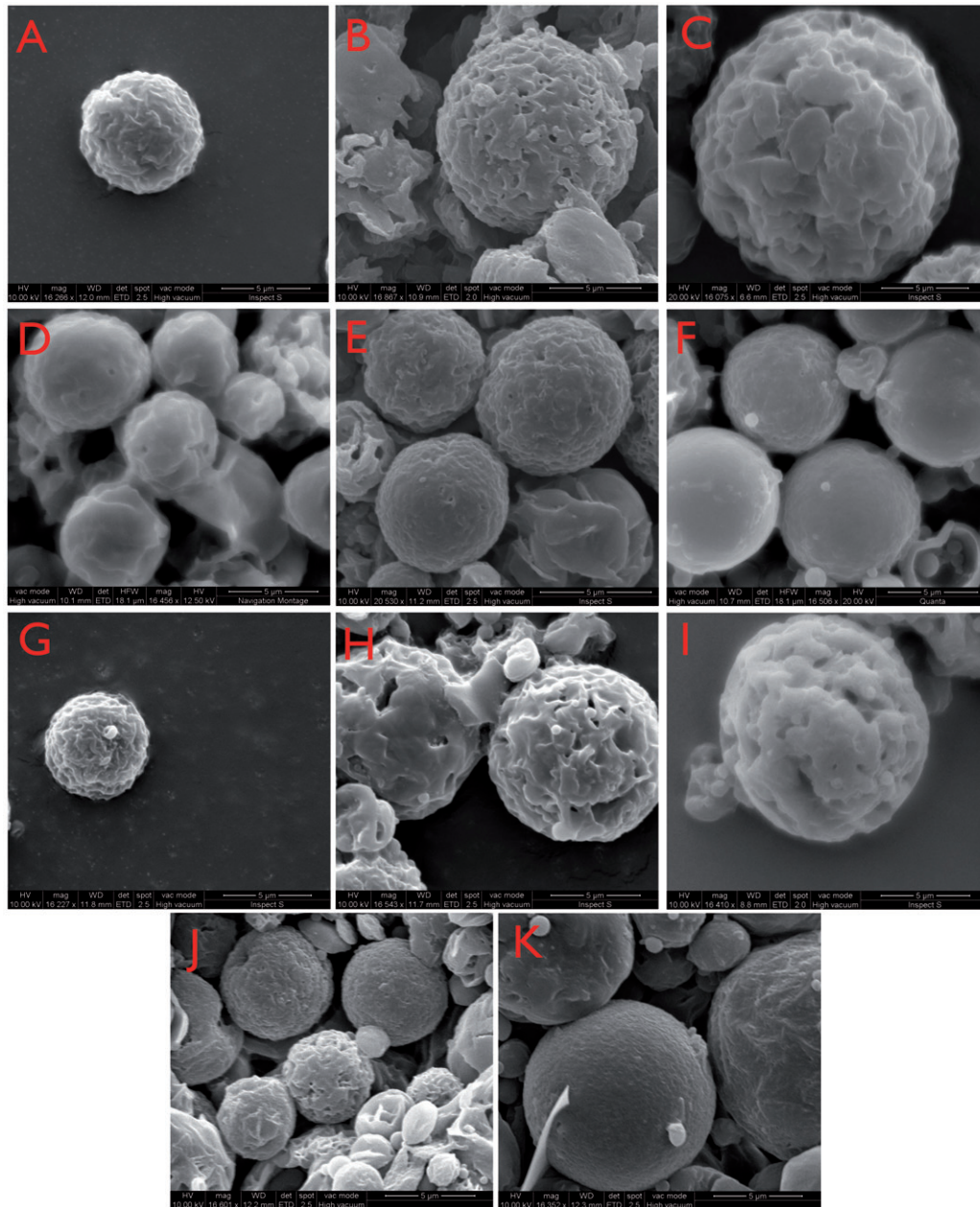
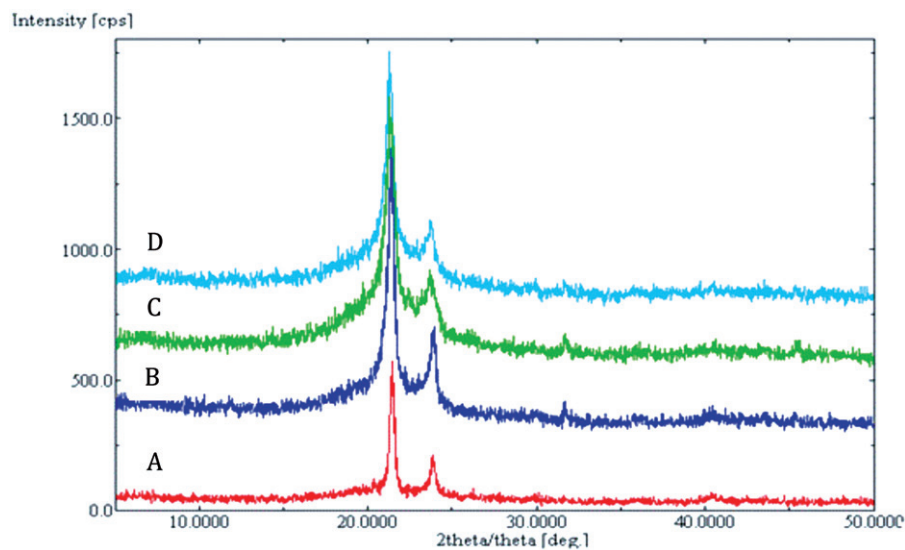


Figure 2. ■■■.

Table 2. Encapsulation efficiencies (%) and enzyme loading ($\mu\text{g}/\text{mg}$ particle) of α -Chymotrypsin (α -CH) and Lysozyme (LS) within polymeric particles formulated over 3 h via the multiple emulsion solvent evaporation technique. The amount of α -CH or LS added into the aqueous phase was 150 mg. The results are the mean of three different prepared batches \pm S.D.

Polymer type	CH		LS	
	EE (%)	Loading ($\mu\text{g}/\text{mg}$ particle)	EE (%)	Loading ($\mu\text{g}/\text{mg}$ particle)
PGA-co-PDL (1:1:1, M_w 11.4 KDa)	14.83 \pm 1.5	49.39 \pm 4.9	32.05 \pm 3.3	108.65 \pm 11.5
PGA-co-PDL (1:1:1, M_w 26.0 KDa)	12.52 \pm 4.4	41.70 \pm 0.01	32.62 \pm 0.5	107.50 \pm 2.1
PGA-co-PDL (1:1:1, M_w 39.2 KDa)	19.40 \pm 2.5	64.60 \pm 8.4	30.11 \pm 4.0	103.60 \pm 10.8
PGA-co-PDL (1:1:0.5, M_w 23.0 KDa)	20.41 \pm 2.5	68.03 \pm 6.4	25.84 \pm 2.6	86.17 \pm 10.6
PGA-co-PDL (1:1:1.45, M_w 34.0 KDa)	23.94 \pm 3.1	79.80 \pm 7.6	**33.93 \pm 1.1	**113.10 \pm 5.8
PPA-co-PDL (1:1:1, M_w 22.0 KDa)	*38.58 \pm 6.4	*128.50 \pm 12.7	36.40 \pm 2.84	121.33 \pm 11.6

**Significant difference PGA-co-PDL (1:1:1.45, M_w 34.0 KDa) versus PGA-co-PDL (1:1:0.5, M_w 23.0 KDa), *significant difference PPA-co-PDL (22.0 KDa) versus PGA-co-PDL (26.0 KDa) at $p < 0.05$.



661 Figure 3. X-ray diffraction pattern of
 662 α -Chymotrypsin-loaded particles formulated
 663 from A, PPA-co-PDL (22.0 KDa); B, PGA-
 664 co-PDL (11.4 KDa); C, PGA-co-PDL
 665 (26.0 KDa); and D, PGA-co-PDL (1:1:1.5,
 666 Mw 34.0 KDa).

682 The PXRD patterns illustrated in Figure 3 indicate that both PGA-
 683 co-PDL and PPA-co-PDL are semicrystalline copolymers. Both
 684 PGA-co-PDL and PPA-co-PDL showed characteristic peaks at
 685 21.5° and 24° 2θ . PGA-co-PDL of different M_w s, have the same
 686 XRD patterns, indicating they have the same level of crystallinity,
 687 and this may explain the similar encapsulation efficiencies
 688 observed. However, the PXRD pattern for PPA-co-PDL has a
 689 flatter baseline between 0° and 20° 2θ , indicating that it is a more
 690 crystalline material. This difference in degree of crystallinity
 691 between PGA-co-PDL and PPA-co-PDL may have influenced
 692 microparticle formation but does not explain the increased
 693 encapsulation efficiency observed with PPA-co-PDL.

694 Furthermore, changing the polymer composition by altering
 695 the pentadecalactone monomeric ratio from 0.5 to 1.5 molar ratio
 696 significantly ($p < 0.05$) increased the encapsulation efficiency of
 697 LS-loaded microparticles. An increase was also observed with
 698 α -CH-loaded particles, but this was not significant ($p > 0.05$).
 699 Compared to PGA-co-PDL, utilizing the more hydrophobic
 700 polymer (PPA-co-PDL) a significant ($p < 0.05$) increase in
 701 encapsulation efficiency and α -CH loading (from 12.52 ± 4.42
 702 to $38.58 \pm 6.48\%$ and 41.70 ± 0.01 to 128.50 ± 12.70 , respect-
 703 ively) was observed. The highest α -CH and LS encapsulation
 704 efficiency and loading were obtained from the most hydrophobic
 705 polymer, PPA-co-PDL. These results suggest that the more
 706 hydrophobic polymers demonstrate better encapsulation effi-
 707 ciency and drug loading of both enzymes compared to the less
 708 hydrophobic variants.

709 Similarly, McGee et al. showed that ovalbumin-loaded
 710 microparticles prepared with PLGA with higher lactide to
 711 glycolide content (85:15) gave higher protein loading compared
 712 to the more hydrophilic one with 50:50 lactide to glycolide
 713 ratio³⁵. Also, higher amounts of bovine albumin were encapsu-
 714 lated using PLGA (75:25) compared to the more hydrophilic
 715 PEGylated PLGA co-polymer³⁶.

716 Comparing the encapsulation efficiencies and enzyme loading
 717 for both enzymes, it was found that LS showed a higher
 718 encapsulation and loading compared to α -CH with all the PGA-
 719 co-PDL variants assessed (Table 2). LS is a smaller, positively
 720 charged enzyme that has the ability to be adsorbed onto the
 721 surface of polymers and this adsorption will affect its encapsu-
 722 lation and release kinetics³⁷. Furthermore, as previously
 723 reported^{37,38}, the temperature rises during the emulsification
 724 steps and the adjustment of the pH to 7.4 can lead to favorable
 725 conditions for LS adsorbing onto polymers. This could result in
 726 increased amounts of LS being encapsulated within PGA-co-PDL.

748 Also, we cannot neglect that using 1% PVA as an emulsifier
 749 imparts a negative charge to the surface of PGA-co-PDL and
 750 PPA-co-PDL which would support enzyme binding. It was
 751 reported that PVA, which is physically entrapped within the
 752 surface layer of the polymer, imparts a negative surface charge on
 753 the microparticles produced^{39,40}. However, comparable amounts
 754 of 128.5 ± 12.7 and $121.33 \pm 11.6 \mu\text{g}/\text{mg}$ particle of α -CH and LS
 755 were encapsulated, using PPA-co-PDL. This represents a signifi-
 756 cant increase over PGA-co-PDL for α -CH- but not LS-loaded
 757 particles.

759 In vitro release

760 It was anticipated that polymer M_w and polymer backbone
 761 chemistry would be important factors affecting the drug release²¹.
 762 Varying the M_w , varies the degradation rate of the polymer and
 763 release kinetics of the drug can be controlled accordingly⁴¹.
 764 Additionally, the hydrophobicity of the polymer can affect the
 765 drug release by reducing the rate of water penetration into the
 766 microspheres and drug egress to some extent compared to the less
 767 hydrophobic polymers⁴². Furthermore, different particle morphol-
 768 ogies may affect the protein release profile through its effect on
 769 the microspheres porosity and the distribution of the drug within
 770 the matrix^{29,43}.

772 The release profiles of either α -CH or LS under sink conditions
 773 from different batches are shown in Figures 4 and 5. Figure 4
 774 shows the release of α -CH from microparticles prepared using
 775 different polymers over 24 h into PBS buffered saline. Most of the
 776 α -CH-loaded microparticle formulations showed a biphasic
 777 release pattern with an initial high burst release phase followed
 778 by a continuous release phase for the first 5 h which became
 779 constant till the end of the release study. The extent of the burst
 780 release varied between different microparticle formulations,
 781 depending on the polymer used, and a notable difference was
 782 observed between PGA-co-PDL (1:1:1 26.0 KDa or 11 KDa) and
 783 the other polymers.

784 Other research groups^{11,33} have observed that increasing
 785 polymer M_w led to a decrease in the total amount of enzyme
 786 released. In this study, there was no general trend observed
 787 between increasing M_w and decreasing enzyme release, which
 788 may be because the differences in M_w were small, but there was
 789 significantly less release after 24 h with PGA-co-PDL (39 KDa)
 790 particles compared to PGA-co-PDL (26 KDa or 11 KDa) par-
 791 ticles. Varying the proportion of PDL within the polymer from 0.5
 792 to 1.5 mole equivalents did not have any consistent effect on the

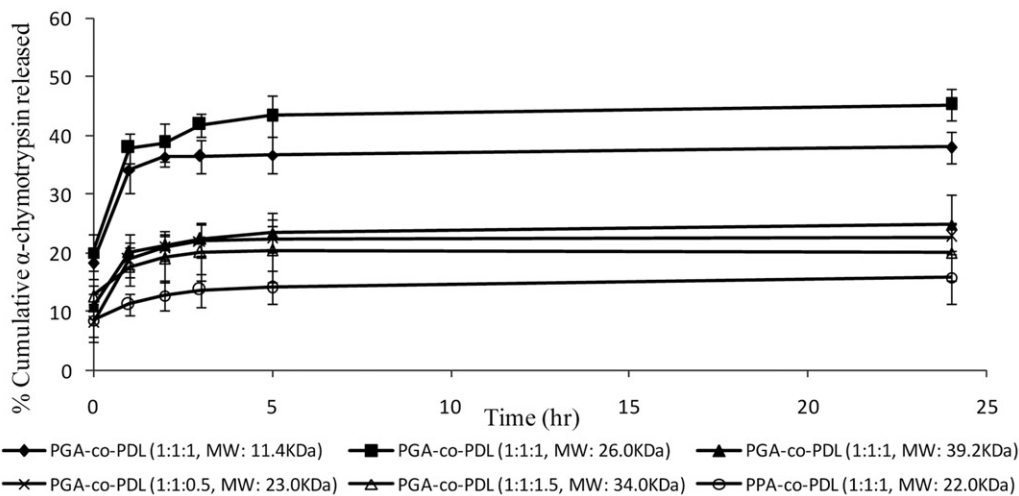


Figure 4. Release profiles of α -Chymotrypsin from polymeric microparticles prepared via the multiple emulsion solvent evaporation technique. The results are the mean of three different prepared batches at each time point \pm S.D.

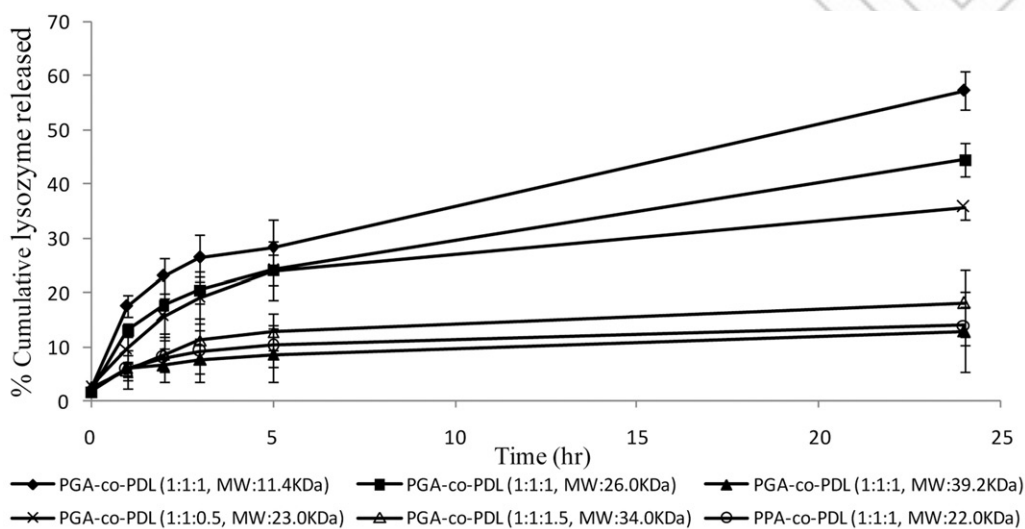


Figure 5. Release profiles of lysozyme from polymeric microparticles prepared via the multiple emulsion solvent evaporation technique. The results are the mean of three different prepared batches at each time point \pm S.D.

α -CH release from microparticles. The biggest difference in release was found when comparing PGA-co-PDL (26 KDa) with the more hydrophobic polymer of a comparable M_w , PPA-co-PDL (22 KDa). Compared to PPA-co-PDL, PGA-co-PDL showed a significantly ($p < 0.05$) higher burst release of α -CH ($20.13 \pm 3.0\%$ compared to $8.54 \pm 2.7\%$) and a greater amount of release after 24 h in PBS buffer ($45.28 \pm 2.7\%$ compared to $15.84 \pm 4.5\%$). Furthermore, PPA-co-PDL demonstrated the lowest burst and total release of α -CH of all the prepared microparticles.

The initial burst release phase of α -CH from these microparticles could be due to the rapid release of protein near to the surface of microparticles which accumulates at the water/oil interface during the solvent evaporation process. The release of the protein entrapped within the polymeric matrix causes a continuous release of α -CH during the first 5 h. Furthermore, the constant release phase could be attributed to the protein aggregation and degradation that occurs during the release process. Despite the higher encapsulation efficiency gained from PPA-co-PDL, these particles demonstrated a slower burst and continuous release rate compared with PGA-co-PDL with comparable M_w . This might be due to the higher hydrophobicity

and slower rate of degradation of this polymer (unpublished data). The lower surface area available for contact with the dissolution medium and the large particle size could be other contributing factors toward this slow release as denser microparticles with smooth surfaces will usually produce a lower rate of initial release compared with rough, porous microparticles. This is in agreement with Thompson et al. who observed a similar effect for ibuprofen release from PGA-co-PDL and PPA-co-PDL microparticles¹⁷.

The release profiles of LS from the different polymeric microparticles are shown in Figure 5. In this case, the LS-loaded microparticle formulations showed a very small initial burst phase followed by continuous release until the end of the release study at 24 h. With LS there was a general trend of increasing PGA-co-PDL (1:1:1) M_w and decreasing enzyme release. The release of LS from the 39 KDa polymer was significantly lower, and there was less difference observed between the 26 KDa and 11 KDa variants. Although, as with α -CH, there was a difference in the release of LS from PPA-co-PDL (22 KDa) and PGA-co-PDL (26 KDa) of a comparable M_w , with LS the release profile of the PPA-co-PDL particles was virtually the same as that of PGA-co-PDL (39 KDa).

925 It was observed that the pattern of LS release was different
 926 from that obtained with α -CH. α -CH release was characterized by
 927 an initial burst followed by a slow continuous release phase for the
 928 first 5 h then a plateau was reached. On the other hand, LS showed
 929 a lower burst release followed by a higher continuous release
 930 phase. This was especially evident with the lower M_w PGA-co-
 931 PDL. The lower burst release could be attributed to the more
 932 efficient encapsulation of LS inside the microparticles with
 933 minimum amounts remaining adsorbed on the surface. Stronger
 934 binding of LS to these polymers could be another reason for this
 935 as LS is cationic and these particles have a slightly anionic surface
 936 from incomplete removal of PVA.

937 With all the microparticles studied, an incomplete release of
 938 enzyme from these was observed even after 3 weeks. This has
 939 been observed by many researchers, and it might be due to
 940 degradation of the protein during the manufacturing of the
 941 microparticles⁴⁴. Formation of intermolecular linkages, hydrolysis
 942 of the protein molecule and the nonspecific adsorption
 943 between polymer and protein either physically or chemically can
 944 lead to protein degradation⁴⁵.

946 Enzyme bioactivity

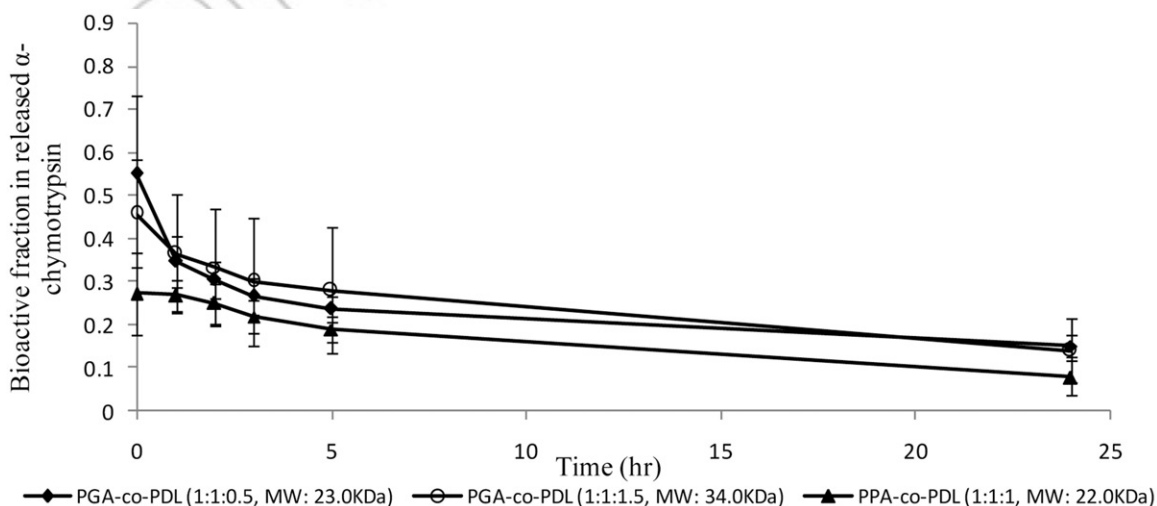
948 Retaining biological activity is crucial for the delivery of enzymes
 949 and peptides, and preservation of the tertiary structure is required
 950 to maintain activity. Enzyme activity before and after encapsu-
 951 lation and upon release can be monitored to investigate the effect
 952 of these processes on biological activity. Many researchers have
 953 estimated the bioactivity of LS by measuring the rate of
 954 degradation of *Micrococcus luteus* cells^{25,46}. However, this
 955 method is not always reproducible because of the dependence
 956 on the ionic strength of the medium^{47,48}. Different methods using
 957 small synthetic substrates have been developed, investigated and
 958 recommended for accurate determination of LS^{49–51}.

959 Observation of the bioactive fraction of α -CH released from
 960 microparticles prepared using PGA-co-PDL and PPA-co-PDL
 961 (Figure 6) indicates that the maximum bioactivity was observed at
 962 zero hours and ranged between 27% and 60%. This was followed
 963 by a sharp decrease in activity during release into PBS buffer (pH
 964 7.4). It was noticed that α -CH released from PGA-co-PDL
 965 exhibited a maximum activity of between 40% and 60%, and PPA-
 966 co-PDL showed the lowest activity of \sim 27% at zero hour.
 967 Furthermore, a gradual loss in bioactivity was recorded for all the
 968 α -CH-loaded microparticles investigated. The reduction in
 969

activity of α -CH could be attributed to conformational changes
 in the α -CH active site during emulsification. The homogeniza-
 tion and use of organic solvents are considered important steps in
 causing protein deactivation and aggregation resulting in a low
 bioactive fraction at zero hour^{52–54}. The gradual loss in activity
 during *in vitro* release was most likely due to autolysis and protein
 fragmentation⁵³. This finding is similar to what was already
 reported by Gaskell et al. where they found that α -CH released
 from PGA-co-PDL-loaded microparticles lost its bioactivity
 gradually with an onset of loss due to proteolysis upon 2 h
 release¹⁹.

At zero hour of release, LS retained almost 100% of its initial
 bioactivity within all the particles investigated. Then, with time it
 began to gradually lose its bioactivity (Figure 7). The higher M_w
 polymer, PGA-co-PDL (1:1:1, 39.0 KDa), and the more hydro-
 phobic polymers, PGA-co-PDL (1:1:1.45) and PPA-co-PDL,
 showed a significantly ($p < 0.05$) higher bioactive fraction, after
 5 and 24 h release, compared to the other co-polymers. The
 maximum LS bioactive fraction was found using PGA-co-PDL
 (1:1:1.45, M_w 34 KDa) and PGA-co-PDL (1:1:1, M_w 39.2 KDa)
 0.78 ± 0.08 and 0.42 ± 0.02 , respectively, after incubation in
 PBS for 24 h.

LS is a relatively stable enzyme⁵⁵ which can better withstand
 the harsh condition of the emulsification process and this was
 confirmed by the retention of its bioactivity at zero time of release
 (bioactive fraction ranged from 0.9 to 1.03 for all the investigated
 polymers, Figure 7). Similarly, it was reported by Giteau and
 coworkers that the LS released from PLGA microspheres was still
 biologically active compared to α -CH, peroxidase and β -galactosidase-
 loaded PLGA microspheres⁵⁷. However, during *in vitro*
 release there was a gradual decrease in the bioactive fraction
 which could be attributed to the effect of PBS buffer on the
 released LS. So, the nature of the release medium on the enzyme
 activity is very important, as many proteins are not stable in buffer
 media at 37 °C. However, for most studies the choice of release
 medium is dictated by the *in vivo* target for delivery of the
 enzyme. Jiang et al. investigated protein stability and protein-
 polymer interactions in different release media and their effect on
 protein release profiles from PLGA microspheres using LS as a
 model protein³⁷. They found that LS showed a higher stability at
 pH 4.0 acetate buffer and pH 2.5 glycine buffer, whereas at pH 7.4
 PBS, the stability was low and significant protein adsorption was
 evident. Furthermore, the higher bioactive fraction of LS in PGA-
 co-PDL (1:1:1, M_w , 39.2 KDa) and PGA-co-PDL (1:1:1.45) could



988 Figure 6. Bioactive fraction of released α -Chymotrypsin from (A) PGA-co-PDL (1:1:1, MW 11.4, 26.0 and 39.2 KDa) and (B) PGA-co-PDL (1:1:0.5,
 989 MW 23.0 KDa, 1:1:1.5, MW 34.0 KDa) and PPA-co-PDL (1:1:1, MW 22.0 KDa) in PBS buffer, pH 7.4. Triplicate samples were used from two
 990 different prepared batches at each time point \pm S.D.

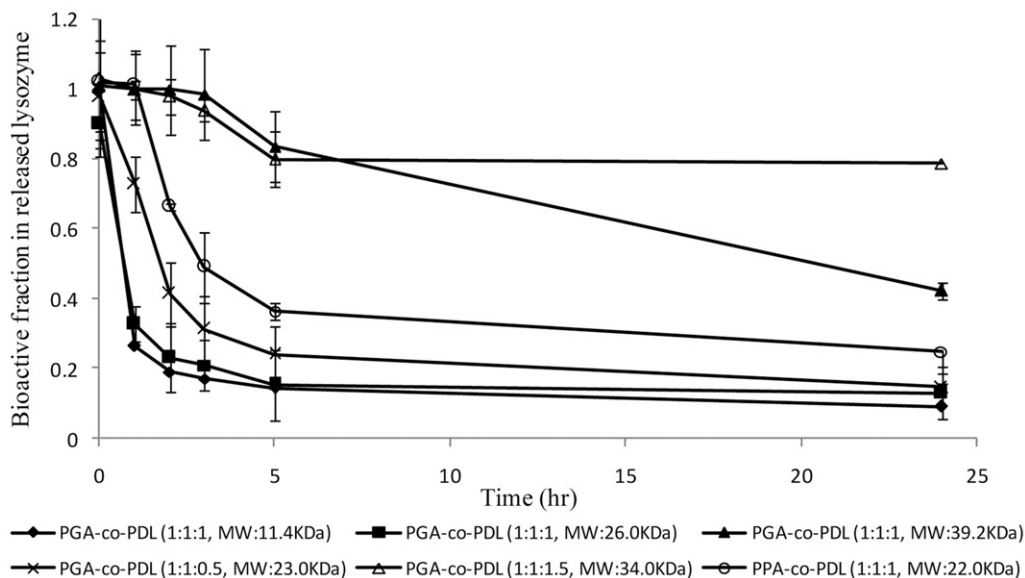


Figure 7. Bioactive fraction of released lysozyme from the different investigated polymers in PBS buffer, pH 7.4. Triplicate samples were used from two different prepared batches at each time \pm S.D.

possibly be attributed to the higher solubility of these polymers in DCM compared with PPA-co-PDL and the lower M_w PGA-co-PDL polymers. Additionally, the longer the contact time of the enzyme in the organic phase, the more enzyme activity would be lost. Thus, a higher solidification rate would be beneficial in retaining the LS biological activity. Similar results were reported by Ghaderi and Carlfors regarding stability of LS during emulsification process within PLGA⁴⁸. Future work will focus on enhancing macromolecule encapsulation efficiency as well as maintaining stability during the manufacturing process. For example, the use of additives to protect the protein structure or the application of alternative formulation methods such as spray drying or *s/o/w* emulsions may substantially reduce the loss in bioactivity during encapsulation.

Conclusion

This research has shown that altering the M_w of PGA-co-PDL from 11.2 to 39.2 KDa had little impact on particle morphology, size, encapsulation efficiency or bioactivity of α -CH- and LS-loaded microparticles. Altering the polymer chemistry had a greater effect, as a higher encapsulation efficiency and drug loading of both α -CH and LS were obtained with PPA-co-PDL compared to PGA-co-PDL particles. A biphasic release pattern was obtained with all microparticles studied, and the release profiles varied according to the polymer used. A lower burst and continuous release was obtained for both enzymes with the more hydrophobic polymers, PPA-co-PDL and PGA-co-PDL (1:1:1.45) and with the higher M_w PGA-co-PDL (39.2 KDa). Furthermore, a very low burst release was recorded with LS compared to α -CH with all the investigated polymers.

One benefit of the low impact of small changes in M_w or PDL content on encapsulation and release is that batch-to-batch variations in the polymers should not have a demonstrable effect on either the properties of particles formed or the encapsulation and release data obtained. These findings suggest that more substantial changes to polymer properties are required to significantly influence the encapsulation and release of proteins. The nature of this type of polymerization reaction means that it is difficult to achieve higher M_w materials and extend the range of M_w s studied. Small changes to the polymer chemistry has been shown to have a greater effect, hence future

studies will focus on further modifying the polymer chemistry either by incorporating different monomers into the backbone or via modification of the pendant hydroxyl groups.

Declaration of interest

The authors report no conflicts of interest.

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