Rapid and Sensitive Analytical Method for the Determination of Insulin in Liposomes by Reversed-Phase HPLC

Eliete de Souza Von Zuben, 1,* Josimar Oliveira Eloy, 2 Victor Hugo Sousa Araujo, 1 Maria Palmira Daflon Gremião 1 and Marlus Chorilli 1

1 São Paulo State University (UNESP), School of Pharmaceutical Sciences, Department of Drugs and Medicines, Rodovia Araraquara-Jauí, km1, 14800-903, Araraquara, São Paulo, Brazil.
2 Federal University of Ceará, School of Pharmacy, Dentistry and Nursing, Department of Pharmacy, Fortaleza – CE, Brazil

* Corresponding author: E-mail: eliete.vz@gmail.com
Tel.: +551633016960
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Abstract
Insulin is an important anabolic hormone that regulates the metabolism of carbohydrates, lipids and proteins. In this study, a reverse-phase liquid chromatography (RP-LC) method was successfully validated and tested for the encapsulation efficiency assay of insulin and in vitro release studies. HPLC analyses were carried out using a RP C18-Luna® Phenomenex (4.6 × 250 mm, 5 μm particle size) column maintained at room temperature, using a mobile phase constituted by a mixture of acetonitrile and 0.1% TFA aqueous solution (60:40, v/v), in an isocratic mode with a flow rate of 1.0 mL/min, with ultraviolet detection at 214 nm and 20 μL of injection volume. Method validation was performed according recognized guidelines for system suitability, specificity, linearity, precision, accuracy, LOD, LOQ and robustness. The method was shown to be linear in the range of 0.5–100 μg/mL (r² = 0.9993) selective, precise, robust, accurate with LOD and LOQ values were 0.097 μg/mL and 0.294 μg/mL, respectively. The developed method proved to be adequate to analyze the encapsulation efficiency and the profile of insulin release from liposomes.

Keywords: Insulin; liposome; liquid Chromatography; validation Method; in Vitro Release; encapsulation Efficiency.

1. Introduction
Insulin is a polypeptide with a molecular weight of 5.8 kDa, synthesized in the pancreas by β-cells of the islets of Langerhans, from a larger molecule called proinsulin. Proinsulin is enzymatically cleaved to generate insulin (which contains 51 amino acids arranged in two chains interconnected by disulfide bonds, α being 21 amino acids and β with 30 amino acids) and the C peptide. The insulin and peptide C are secreted in equimolar amounts when beta-pancreatic cells are stimulated, however, peptide C does not play any physiological function.1

Insulin is an important anabolic hormone that regulates the metabolism of carbohydrates, lipids and proteins. The uptake of glucose by organs such as liver, skeletal muscle and adipose tissue is performed by the action of insulin. In the absence of insulin, the uptake of glucose by said organs and tissues is affected leading to hyperglycemia. Other insulin functions are related to ion transport, cell proliferation and differentiation, and nitric oxide synthesis.2

Recent research has triggered the development of new insulin delivery systems that allow the use of alternative routes to parenteral (subcutaneous injection). The injection of insulin preparations sometimes results in subcutaneous adipose tissue hypertrophy, if it will be administering repeatedly in the same place and the other side effect is the risk of hypoglycemia when treatment becomes continuous.3

Liposomes in 1995 were the first nanoscale release system to be used for clinical use. Since then, work based on this type of release systems has grown considerably and brought enormous development with significant clinical implications.4 Several researchers used nanostructured systems such as liposomes for encapsulation and release of insulin.5-7

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2. Experimental

2.1. Chemical and Reagents

The human insulin solution Novolin R® (100IU/mL, Lot GS63F52) was purchased from Novo Nordisk, Kalundborg, Denmark. Soybean phosphatidylcholine (Lipoid S100; PC > 96%) was obtained from Lipoid GmbH, Germany. Cholesterol (CH) was purchased from Sigma-Aldrich, USA. Acetonitrile and methanol were acquired from J.T. Baker®, USA (HPLC grade) and trifluoroacetic acid (TFA) were purchased from Sigma-Aldrich, USA. Amicon®-100 kDa ultrafiltration filter was purchased from Millipore. Water was prepared by Milli-Q reverse osmosis (Millipore, USA). All other reagents were commercially available and of analytical grade. Phosphate buffered saline (PBS) is made of NaCl 137 mmol/L, KCl 2.6 mmol/L, Na₂HPO₄·12H₂O 6.4 mmol/L and NaH₂PO₄ 1.4 mmol/L, pH 7.4.

2.2. Preparation of Reference Standard Solution

The stock solution of 700 µg/mL was prepared by direct dilution of 2 mL of human insulin Novolin R® (100 IU/mL) in a 10 mL volumetric flask. The volume was completed with phosphate buffer solution (PBS), pH 7.4. From the stock solution, calibration curves for seven human insulin standard solutions were prepared at concentrations of 0.5, 1, 5, 20, 40, 50 and 100 µg/mL in phosphate buffer solution (PBS) pH 7.4. All solutions were filtered through a 0.45 µm PVDF membrane filter (Millipore) prior to injection and assayed by HPLC.

2.3. HPLC System and Chromatographic Conditions

The RP-LC method was performed on an Agilent Technologies LC system, model 1200 DAD/UV–visible detector and was carried out the adaptation (change in the proportion of the mobile phase) of the method described by Sarmento and collaborators. The detector was set to 214 nm and peak areas were integrated automatically using a computer system with ChemStation data acquiring software. RP C18- Luna® Phenomenex (4.6 × 250 mm, 5 µm particle size) column was employed for chromatographic separation with a mobile phase constituted by a mixture of acetonitrile and 0.1% TFA aqueous solution (60:40, v/v), in an isocratic mode with a flow rate of 1.0 mL/min. The HPLC system was performed at room temperature (25 ± 1 °C). The injection volume was 20 µL for both the reference substance solution and the sample solution. Prior to use, acetonitrile and 0.1% TFA aqueous solution prepared with deionized water (Milli-Q®), filtered with a 0.45 µm filter and degassed for 30 min. All samples were filtered through a 0.45 µm PVDF membrane filter from Millipore, USA.

2.4. Validation of HPLC Method

Method validation was performed in agreement with ICH and FDA guidelines together Brazilian National Health Surveillance Agency (ANVISA) resolution for system suitability, specificity, linearity, precision, accuracy, limit of detection (LOD), limit of quantitation (LOQ) and robustness.13–15

2.4.1. System Suitability Test

In order to assure the performance and reproducibility of the chromatographic system before and during the analysis, six injections of insulin standard solution were carried out at a concentration of 50 µg/mL. The parameters measured were retention time, peak area, peak symmetry, tailing factor and number of theoretical plates.14

2.4.2. Selectivity

Chromatograms of mobile-phase components, samples of phosphate buffer solution (PBS), pH 7.4 and samples of blank liposomes prepared without adding insulin were obtained to evaluate the selectivity of the method. All samples were filtered through a 0.45 µm PVDF membrane filter (Millipore) and injected in triplicate.

2.4.3. Linearity

The linearity was evaluated by the construction of three analytical curves, determined by the analysis of seven levels of concentration in three different days. Standard solutions at the concentrations of 0.5, 1, 5, 20, 40, 50 and 100 µg/mL were prepared from the insulin stock solution (700 µg/mL) using phosphate buffer solution (PBS, pH 7.4) as the diluent. For each described concentration, three injections were carried out previously filtered through a 0.45 µm PVDF membrane filter (Millipore). Linearity was determined by the calculation of a linear regression using the least-squares method and the statistical analysis was performed by analysis of variance (ANOVA).
2. 4. 4. Precision

The precision of the analytical method was evaluated by determination of the repeatability and the intermediate precision. The repeatability (intra-day) was determined by injection of six independent preparations of insulin standard solutions at a concentration of 50 µg/mL, on the same day and under the same experimental conditions. The intermediate precision (inter-day) was tested using two different analysts on different days, with the same LC instrument, in the same laboratory. The % RSD of assays was calculated.

2. 4. 5. Accuracy

The accuracy was assessed by standard addition technique where liposomes were spiked with known quantities of insulin standard solution, at three different levels (lower, medium and upper concentration) corresponding to 5, 20 and 50 µg/mL. The samples were prepared in triplicate and the recovery percentage was determined as the percentage of the drug recovered from the sample and expressed as a relative percentage (%).

2. 4. 6. Limit of Detection (LOD) and Quantitation (LOQ)

The LOD and LOQ were calculated, using the mean values of three independent analytical curves, determined by a linear regression model, based on the standard deviation of the response and on the slope of the calibration curve, according to following equations 1 and 2 respectively:

LOD = 3.3 \((\sigma/S)\) (1)

LOQ = 10 \((\sigma/S)\) (2)

where “\(\sigma\)” represents the standard deviation of y-intercept of regression line and “\(S\)” denotes the slope of regression line.

2. 4. 7. Robustness

Robustness was assessed by varying four parameters independently. The flow rate of the mobile phase (0.9 mL/min and 1.1 mL/min), wavelength (213 nm and 215 nm), column temperature (24 °C and 26 °C) and the ratio of mobile phase constituted by a mixture of acetonitrile and 0.1% TFA aqueous solution (61:39 and 59:41). For this study, the analysis was performed using a concentration of 50 µg/mL and all samples were filtered through a 0.45 µm PVDF membrane filter (Millipore) and injected in triplicate.

2. 5. Application of the Method
2. 5. 1. Preparation of Blank Liposome and Insulin-loaded Liposome

Blank liposomes and insulin-loaded liposomes were prepared by the hydration of the thin lipid film as previously described by Bangham et al. Briefly, a phospholipid mixture of 100mg, including soybean phosphatidylcholine (PC) and cholesterol (CH) was weighed (weight ratios of 5:3) into a round-bottom flask and dissolved in a chloroform / methanol mixture (1:2 v/v), evaporated using a rotary evaporator at 40 °C for 30 minutes. Then, to prepare blank liposomes, the lipid film formed was hydrated with 10 mL of phosphate buffer (PBS, pH 7.4) for 30 min at 40 °C / 100 rpm, obtaining the liposomal suspension. To prepare the liposome loaded with insulin, hydration was performed with 10 mL of phosphate buffer (PBS, pH 7.4), containing 4 mL of insulin solution, equivalent to 1400 µg / mL, obtaining the liposomal suspension.

2. 5. 2. Insulin Encapsulation Efficiency (EE)

For the determination of the encapsulation efficiency of insulin, an indirect method was used by determining the free amount of insulin, non-encapsulated. Briefly, 1 mL of the encapsulated insulin from the liposome was separated from the free insulin by ultracentrifugation of liposomes using Amicon®-100 kDa ultrafiltration filter at 13,000 rpm for 12 min. After centrifugation, 250 µl of the filtrate was diluted to 2 mL with phosphate buffer (PBS, pH 7.4) and measured by an HPLC using the validated analytical method described previously. All experiments were performed in triplicate. The EE % of insulin was calculated according to equation 3 below.

\[
\text{EE\% of insulin} = \left(\frac{\text{Total amount of insulin (µg)} - \text{Free amount of insulin}}{\text{Total amount of insulin (µg)}}\right) \times 100
\] (3)

2. 5. 3. In Vitro Release of Insulin Solution and Insulin from Liposomes

In vitro insulin release profile from solution and liposome was investigated using Franz diffusion cells (Hanson MicroettePlus® equipment, Chatsworth, CA, USA). The cells were filled with 7.0 mL of receptor solution composed by phosphate buffer solution (PBS, pH 7.4), stirred at 300 rpm and heated at 32 °C ± 0.5 °C to simulate nasal temperature. Aliquots of samples (2,8 mL) were withdrawn automatically at predetermined intervals (0.08, 0.5, 1, 2, 4, 8, 12, 16 and 24 h) and sample volumes were replenished with fresh receptor solution.
The amount of insulin released from solution and the liposomes was quantified by the HPLC method as described in the section 2.3.

3. Results and Discussion

A reversed-phase HPLC method for the determination of insulin in liposomes has been developed. The analytical parameters of selectivity, linearity, precision, accuracy, LOD, LOQ, and robustness were evaluated to validate the method, according to recognized guidelines (FDA, ICH e Anvisa). During preliminary studies, some chromatographic conditions were tested. It was decided to carry out the adaptation of a method previously reported. However, the proportion of the mobile phase was modified using the isocratic mode as a choice and this modification provided a reduced run time, best chromatographic peak resolution and good symmetry. Moreover, no degradation peak was observed.

3. 1. Validation of HPLC Method

Validation of the method was performed according to ICH and FDA guidelines together Brazilian National Health Surveillance Agency (ANVISA) resolution. Table 1 represents the HPLC conditions for the determination of insulin in liposome.

The system suitability is used to check if the chromatographic system is suitable for the intended analysis that will be executed. Figure 1 shows two peaks (between 1.6 minutes and 3 minutes) from the mobile phase and the phosphate buffer (PBS, pH 7.4) and a representative peak of insulin (4.5 minutes). Besides, no peak of possible degradation products was observed in the chromatograms, as verified in the selectivity tests. The following results were obtained: tailing factor of 1.41, peak asymmetry of 0.64 and theoretical plates of 5089. All parameters found are in agreement with the FDA and ICH guidelines. The average retention time was 4.5 minutes and the % RSD of peak area was 0.22%.

In this selectivity parameter mobile-phase components, samples of phosphate buffer solution (PBS), pH 7.4 and the samples of blank liposomes composed by soybean phosphatidylycholine and cholesterol prepared without adding insulin were previously diluted with acetonitrile, sonicated for 10 minutes, filtered (0.45 μm) and then analyzed by HPLC. All the samples were injected separately and the assay was performed in triplicate and the peaks obtained showed that there was no interference of the samples injected at the peak of insulin.

Table 1. HPLC conditions for determination of insulin in liposomes.

<table>
<thead>
<tr>
<th>System Parameters</th>
<th>Parameters</th>
</tr>
</thead>
<tbody>
<tr>
<td>HPLC equipment</td>
<td>HPLC AGILENT, model 1200</td>
</tr>
<tr>
<td>Mobile phase</td>
<td>Acetonitrile : 0.1% TFA aqueous solution (60:40,v/v)</td>
</tr>
<tr>
<td></td>
<td>RP C18- Luna® Phenomenex (4.6 × 250 mm, 5 μm particle size)</td>
</tr>
<tr>
<td>Column</td>
<td>214 nm</td>
</tr>
<tr>
<td>Wavelength</td>
<td>1.0 mL/min</td>
</tr>
<tr>
<td>Flow rate</td>
<td>20 μL</td>
</tr>
<tr>
<td>Volume of injection</td>
<td>25 ± 1 °C</td>
</tr>
<tr>
<td>Retention time</td>
<td>4.5 min</td>
</tr>
</tbody>
</table>

Figure 1. Representative chromatogram of insulin reference substance. Conditions: mixture of acetonitrile and 0.1% TFA aqueous solution (60:40,v/v) as mobile phase, flow rate at 1.0 mL/min, column temperature 25 ± 1 °C, RP C18- Luna® Phenomenex (4.6 × 250 mm, 5 μm particle size) column, UV detection wavelength at 214 with 20 μL of injection volume.
The linearity of the method was assessed by plotting peak area against seven concentrations. For each concentration, three injections were performed and the average results of the chromatographic peak areas obtained were used for the study of linear regression using the least-squares method. The analytical curve of insulin based on three calibration curves was shown to be linear over the proposed range (0.5–100 µg/mL) resulting in linear regression $y = 55.01775x + 36.56067$, with $r^2 = 0.9993$, as advised and shown in Figure 2. The statistical analysis was evaluated by ANOVA (Table 2) and data no presented significant linearity deviation at a 5% level of variance, exhibiting $F$ calculated (2.80) lower than the $F$ critical (2.96) demonstrating that the method used is linear in the range of 0.5 to 100 µg/mL.

The precision was evaluated by calculating of relative standard deviation (RSD) of the samples, submitted to the repeatability (intra-day) and intermediate precision (inter-day). According to Table 3, the precision values were less than 5% as recommended, where 0.77% was found for repeatability and 1.18% was found for intermediate precision. These results indicate that the analytical method presented has good precision.

The accuracy of the method was determined by recovery studies, where the method of standard addition was used at three different levels (5, 20 and 50 µg/mL). The results are shown in Table 4 and the average recoveries for insulin were 97.58% within the range of 80–120% with an RSD of 1.27%, evidencing the method is appropriate in quantify concentrations of insulin in liposome with accuracy.

The LOD and LOQ are used to measure the sensitivity of the method. DL represents the smallest amount of drug present in the sample that can be detected, but not necessarily quantified while QL represents the smallest amount of drug in the sample that can be measured with precision and accuracy by the method developed. The calculated LOQ and LOQ values were 0.097 µg/mL and 0.294 µg/mL, respectively, demonstrating that the proposed method is adequate and safe for the quantification and detection of low insulin concentrations.

Robustness is a parameter typically performed in the development of the analytical method that indicates its ability to withstand small and deliberate variations in analytical conditions. It was evaluated by small modifications in four pre-established parameters, such as flow rate of the mobile phase, ratio of mobile phase, column tem-

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**Table 2.** Analysis of variance (ANOVA) for linearity

<table>
<thead>
<tr>
<th>Source of variation</th>
<th>DF</th>
<th>SS</th>
<th>MS</th>
<th>F calculated</th>
<th>F critical</th>
</tr>
</thead>
<tbody>
<tr>
<td>Between groups</td>
<td>6</td>
<td>71158775</td>
<td>11859796</td>
<td>2.80</td>
<td>2.96</td>
</tr>
<tr>
<td>Within groups</td>
<td>14</td>
<td>2222</td>
<td>159</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Total</td>
<td>20</td>
<td>71160997</td>
<td></td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

DF—degrees of freedom; SS—Sum of squares; MS—Mean square

**Table 3.** Results of intra and inter-day precision for determination of insulin in standard solutions.

<table>
<thead>
<tr>
<th>Level</th>
<th>Concentration µg/mL</th>
<th>RSD* (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Repeatability (intra-day)</td>
<td>50µg/mL</td>
<td>0.77 (n = 6)</td>
</tr>
<tr>
<td>Intermediate precision (inter-day)</td>
<td>50µg/mL</td>
<td>1.18 (n = 12)</td>
</tr>
</tbody>
</table>

*RSD: relative standard deviation.

**Table 4.** Determination of the recovery of the analytical method for insulin in liposomes

<table>
<thead>
<tr>
<th>Insulin standard solution added (µg/mL)</th>
<th>Insulin standard solution found * (µg/mL)</th>
<th>Recovery (%)</th>
<th>RSD b (%)</th>
<th>Average Recovery (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>R1</td>
<td>5</td>
<td>4.88</td>
<td>93.88</td>
<td>2.26</td>
</tr>
<tr>
<td>R2</td>
<td>20</td>
<td>19.85</td>
<td>99.26</td>
<td>0.27</td>
</tr>
<tr>
<td>R3</td>
<td>50</td>
<td>49.74</td>
<td>99.61</td>
<td>1.28</td>
</tr>
</tbody>
</table>

* Mean of three replicates analysis.  
  b RSD: relative standard deviation.
perature and wavelength (nm), using a concentration of 50 µg/mL. According to Table 5, it was possible to verify that the method is robust since even with the variations made in the analytical conditions, the recovery results are within a previously specified range of 80–120% and RSD values showed less than 2%.

**Table 5. Robustness of the analytical method for insulin quantification**

<table>
<thead>
<tr>
<th>Chromatographic conditions</th>
<th>Recovery(%)</th>
<th>RSD a (%)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Flow rate of the mobile phase</td>
<td></td>
<td></td>
</tr>
<tr>
<td>0.9 mL/min</td>
<td>98.8</td>
<td>0.60</td>
</tr>
<tr>
<td>1.1 mL/min</td>
<td>104.4</td>
<td>0.18</td>
</tr>
<tr>
<td>Ratio of mobile phase</td>
<td></td>
<td></td>
</tr>
<tr>
<td>61:39</td>
<td>99.3</td>
<td>0.05</td>
</tr>
<tr>
<td>59:41</td>
<td>98.0</td>
<td>0.20</td>
</tr>
<tr>
<td>Column temperature (°C)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>24°C</td>
<td>101.3</td>
<td>0.58</td>
</tr>
<tr>
<td>26°C</td>
<td>99.2</td>
<td>0.49</td>
</tr>
<tr>
<td>Wavelength (nm)</td>
<td></td>
<td></td>
</tr>
<tr>
<td>213 nm</td>
<td>100.8</td>
<td>0.83</td>
</tr>
<tr>
<td>215 nm</td>
<td>100.3</td>
<td>0.34</td>
</tr>
</tbody>
</table>

a RSD: relative standard deviation.

**3.2. Application of the Validated Method**

Liposomes constituting important systems for the encapsulation, transport and sustained release of drugs, because their morphology is similar to that cellular membranes, besides protecting the drug against degradation.25–27 The validated method was used for the determination of insulin in liposomes composed by soybean phosphatidylcholine and cholesterol prepared by the thin film-hydration technique, through the encapsulation efficiency assay and the *in vitro* release profile of insulin from the liposomes. The encapsulation efficiency of insulin from liposome was found to be 67.19% ± 2.4 (SD, n = 3), which shows a result superior to that found in similar papers.6,7,28 However, as far as we know, no study reported in the literature has addressed the encapsulation efficiency of insulin from liposomes using Amicon ultrafiltration filter extraction with consequent HPLC quantification. A study by Wallace and colleagues showed that ultrafiltration has several advantages over ultracentrifugation, such as rapid separation with low centrifugal forces, suggesting less potential for particle deformation and therefore less impairment of particle integrity.29 Figure 3 shows the release profiles of insulin in solution (PBS, pH 7.4) and insulin from the liposome. Both profiles have a biphasic release behavior, that is, it was observed an initial burst effect (fast release) in the first 2 hours, followed slower and sustained release delivery of insulin in the next hours. However, comparing the two release profiles it is noted that the release of insulin from the liposome was slower and sustained, suggest it is originated from the drug encapsulated in the aqueous liposome compartment and due to the process of diffusion of insulin through the bilayers. Pardakhty and collaborators obtained similar results in their studies as well as Zhang and collaborators.30,31

**Figure 3.** Release profiles of insulin in solution (open circles) and insulin from the liposome (solid circles). Each value express mean± standard deviation, n = 5.

**4. Conclusion**

The RP-HPLC method was developed and validated according to recognized guidelines (FDA, ICH and Anvisa) for the measurement of insulin in liposomes composed by soybean phosphatidylcholine and cholesterol and prepared using the hydration technique. The insulin encapsulation efficiency assay in liposomes and the *in vitro* release study using Franz diffusion cells were evaluated. The results indicated that the analytical method was linear, selective, precise, accurate and robust with low detection and quantification limits in a range from 0.5 to 100 µg/mL and are within the limits proposed by those guidelines. The method is simple, rapid and sensitive and may be used to determine insulin in liposomes in a short analysis time (about 4.5 minutes), which shows an advantage of the proposed method over those described in the literature. The suggested method can be used to analyze the encapsulation efficiency with adequate reliability. Regarding release profile of insulin from liposomes the data demonstrated a biphasic release behavior characterized by burst effect followed slower and sustained release.

**Acknowledgements**

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Conflict of Interest
There are no conflicts to declare.

5. References
Povzetek

Inzulin je pomemben anabolični hormon, ki uravnava presnovo ogljikovih hidratov, lipidov in beljakovin. V tej raziskavi je bila metoda reverzno fazne tekočinske kromatografije (RP-LC) uspešno validirana in preizkušena na učinkovitost enkapsulacije inzulina in in vitro študij sproščanja. HPLC analize smo izvedli s kolono RP C18-Luna® Phenomenex (4,6 × 250 mm, velikost delcev 5 μm), pri sobni temperaturi, z uporabo mobilne faze sestavljene iz mešanice acetonitrila in 0,1 % vodne raztopine TFA (60:40 , v/v), v izokratskem načinu s pretokom 1,0 mL/min, z ultravijolično detekcijo pri 214 nm, volumen injicirane raztopine pa je znašal 20 μL. Validacija metode je bila izvedena v skladu s priznanimi smernicami glede ustreznosti sistema, specifičnosti, linearnosti, natančnosti, točnosti, meje detekcije (LOD), meje kvantizacije (LOQ) in robustnosti. Pokazalo se je, da je metoda linearna v območju od 0,5-100 μg/mL (r^2 = 0,9993) selektivna, natančna, robustna, točna z vrednostmi LOD in LOQ 0,07 μg/mL oziroma 0,294 μg/mL. Razvita metoda se je izkazala za ustrezno za analizo učinkovitosti enkapsulacije in analizo profila sproščanja inzulina iz liposomov.