STUDIA UBB CHEMIA, LXVI, 3, 2021 (p. 209-224) (RECOMMENDED CITATION) DOI:10.24193/subbchem.2021.3.13

CURCUMIN-WHEY PROTEIN SOLID DISPERSION SYSTEM WITH IMPROVED SOLUBILITY AND CANCER CELL INHIBITORY EFFECT

LEVENTE RÁCZ^a, MARIA TOMOAIA-COTIȘEL^a, CSABA-PAL RÁCZ^a, PAULA BULIERIS^b, IOANA GROSU^b, SEBASTIAN PORAV^b, ALEXANDRA CIORÎȚĂ^b, XENIA FILIP^b, FLAVIA MARTIN^b, GEORGETA SERBAN^c, IRINA KACSÓ^{b,*}

ABSTRACT. The solid dispersion system containing a high amount of the natural compound curcumin was prepared with whey protein concentrate by spray-drying method in 5:1 molar ratio. X-ray powder diffraction and DSC techniques show the formation of the solid dispersion system in amorphous state, and the presence of weak hydrogen bond type interactions between the components was established by FTIR analysis. SEM images show highly homogeneous donut-like spherical microparticles morphology for the system. The solubility of curcumin from the system was enhanced compared to practically insoluble raw curcumin, reaching a value of 70 μ g/mL in aqueous buffer solution at pH=8 similar with intestinal environment. The synthesized material had better effects against skin melanoma cells, compared to lung adenocarcinoma cells, but in both cases the effect was promising, and through further and more complex analyses the antitumoral potential of CUC-WPC_SD could be exploited.

Keywords: curcumin, whey protein concentrate, complexation, enhanced solubility, anticarcinogenic effect

INTRODUCTION

Curcumin (CUC) (1E,6E)-1,7-Bis(4-hydroxy-3-methoxyphenyl)-hepta-1,6-diene-3,5-dione), with a diferuloylmethane structure was first isolated from rhizomes of *Curcuma longa* in 1815 as a yellow compound which exhibits

^a Babeş-Bolyai University, Faculty of Chemistry and Chemical Engineering, 11 Arany Janos str., RO-400028, Cluj-Napoca, Romania

^b National Institute of Research&Development for Isotopic and Molecular Technologies, 67-103 Donath str., RO-400293 Cluj-Napoca, Romania

^c University of Oradea, Faculty of Medicine and Pharmacy, 29 Nicolae Jiga str., RO-410028, Oradea, Romania

^{*} Corresponding author: irina.kacso@itim-cj.ro

biological and pharmaceutical activities. It has been used in traditional Chinese and Indian medicines for treatment of various diseases [1]. More and more research results show the efficiency of curcumin in the treatment of many medical conditions, as a result of the antioxidant, anti-inflammatory [2], antimicrobial [3], antiviral [4], neuroprotective [5] and, last but not least, anti-carcinogenic effect [6].

The chemical structure of CUC ($C_{21}H_{20}O_6$) (Figure 1) consists in two aryl rings containing o-methoxy phenolic OH-groups symmetrically linked in conjugation through a β -diketone moiety which gives particular characteristics, such as intramolecular hydrogen atom transfer which determines the existence of keto-enol tautomerism, with important role in its physical-chemical properties [7]. The CUC molecule hydrophobicity is responsible for the very low solubility in aqueous medium, poor absorption, bioavailability and stability [8] that limit its therapeutic effectiveness. Thus, a large number of researches have been carried out and published in the last decade in attempt to improve the solubility and bioavailability of CUC, and to demonstrate its therapeutic effect in treatment of various diseases. Crystal engineering studies have led to the obtaining of polymorphs and co-crystals [9, 10] with higher solubility and dissolution rate compared to pure starting CUC. Inclusion complexes and different incorporation strategies of CUC into various carrier molecules, including but not limited to nanoparticles [11, 12], polymers [13], cyclodextrins [14], liposomes [11] or different proteins [15, 16] have been obtained and characterized, as well as micelles and solid dispersions [10, 11, 17], respectively.



Figure 1. Chemical structure of curcumin

One of the most used protein for increasing the solubility and bioavailability of CUC is whey protein. This protein possesses important nutritional and biological properties [18], antimicrobial, antiviral, anticarcinogenic activity and other metabolic features that have been associated with it [19]. It's being used in the whey protein concentrate (WPC) or isolate (WPI) form, that contains 65-90% β -lactoglobulin [18]. In addition to the β -lactoglobulin main component the whey protein also contains α -lactalbumin, immunoglobulin, serum albumin, casein and other components. The amino acid sequence and three-dimensional structure confer to the β -lactoglobulin the ability to

bind small hydrophobic ligands and therefore it can act as a specific transporter for them [20]. This ability is also supplemented by its antioxidant potential [21].

The solubility, stability and bioavailability improvement of CUC with whey protein or β -lactoglobulin was performed by different ways. CUC was loaded in WPI obtaining whey protein aggregates in emulsion [16] or whey protein hydrolysate in suspension [22]. Other studies reported the complexation of CUC and WPI in form of micro- or nano-encapsulates [23-25] or nano-emulsions [26], the β -lactoglobulin being also used for the same purpose [27-29]. In addition, the biological activity of CUC, such as the antioxidant [16, 23, 30] and anticancer [25] effect, was enhanced, thus encouraging the use of these complexes in pharmaceutical, nutraceutical or food [16] applications.

Most studies on CUC with whey protein or β -lactoglobulin complexation have been performed on 1:1 molar ratio [16, 27, 28, 31]. Few studies investigate the complexation using an excess of CUC, as in the case of nano-encapsulates [25].

The aim of our study is to prepare an enhanced solubility solid-state curcumin-whey protein concentrate dispersion system (CUC-WPC_SD) with high amount of curcumin content by spray drying (SD) method, whose efficiency is proved by the literature data [15, 24, 30].

Curcumin – whey protein solid dispersion systems were prepared in different molar ratios, from 1:1 to 20:1, whose preliminary analysis regarding the solubility and stability showed that 5:1 stoichiometry (CUC-WPC_SD) is the optimal formula for the proposed purpose. The identification of the system was highlighted by X-ray powder diffraction (XRPD), Fourier transform infrared spectroscopy (FTIR) and differential scanning calorimetry (DSC). The morphological characterization was realized by scanning electron microscopy (SEM) and the dissolution by UV spectrophotometry. The *in vitro* cytostatic activity was tested on several cancer cell types.

RESULTS AND DISCUSSION

CUC-WPC_SD complex characterization

X-ray powder diffraction (XRPD)

In order to prove the solid disperse system CUC-WPC_SD obtained by CUC incorporation into WPC, in the optimized molar ratio 5:1, first it was analyzed by XRPD.



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Figure 2. The compared PXRD patterns of CUC, WPC and CUC-WPC_SD

The diffractograms of raw CUC and WPC were compared with CUC-WPC_SD one (Figure 2). It was observed that the analyzed sample is in an amorphous state, without being able to distinguish the characteristic diffraction lines of CUC, which suggests its loading in WPC and a homogeneous dispersion system formation.

Differential scanning calorimetry (DSC)

The DSC curve of CUC (Figure 3) shows a sharp endothermic melting signal with T_{on} =172.6°C, T_{peak} =177.8°C and Δ H=-194.6 J·g⁻¹, and presents thermal stability up to 200°C. The DSC trace of WPC shows an amorphous material, without significant thermal events, observing only one broad endothermic signal between 40-100°C, corresponding to the unbounded water elimination.

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Figure 3. The compared DSC curves of CUC, WPC and CUC-WPC_SD

Analyzing the DSC curve of CUC-WPC_SD one can observe one low intensity broad endotherm signal between 40 and 90°C due to the residual unbounded water elimination, but the melting signal of CUC does not appears, which indicates that the loading of curcumin in whey protein in 5:1 stoichiometric ratio occurs.

Fourier transform infrared spectroscopy (FTIR)

The changes in the position, intensity or width of the characteristic vibrational bands of curcumin and whey protein on the CUC-WPC_SD system FTIR spectrum could be explained by the transformations and interactions that occur during the incorporation process. On the CUC-WPC_SD dispersion system FTIR spectrum one can identify the most important absorption bands of CUC [32] and those of WPC with certain modifications (Figure 4).

Regarding the characteristic vibrational bands of CUC in the spectrum of CUC-WPC_SD sample the following changes appear: the C=O stretching from 1627 cm⁻¹ and the aromatic C=C stretching from 1602 cm⁻¹ disappear; the aromatic ring bending vibration from 1509 cm⁻¹ shift to 1516 cm⁻¹; the CH₂ bending from 1429 cm⁻¹ and the C-O stretching from 1207 cm⁻¹ shift to 1431 cm⁻¹ and 1210 cm⁻¹, respectively and both appear as a shoulder; the C-O stretching from 1154 cm⁻¹ and 1121 cm⁻¹ shift to 1163 cm⁻¹ and

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1127 cm⁻¹, respectively; the C-OH stretching from 1028 cm⁻¹ and the benzoate *trans*-CH vibration from 963 cm⁻¹ shift to 1034 cm⁻¹ and at 970 cm⁻¹, respectively.



Figure 4. The compared FTIR spectra of CUC, WPC and CUC-WPC_SD, spectral domain a) 3750-2750 cm⁻¹ and b) 1800-400 cm⁻¹

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As a result of the loading with CUC, the characteristic vibrational bands of WPC have undergone the following changes: the shoulder at 1694 cm⁻¹, assigned to protein aggregates [33], no longer appears; the amide I band, due to the stretching vibrations of C=O in the peptide bonds [34] from 1657 cm⁻¹ shift to 1650 cm⁻¹ and becomes narrower; the vibration bands from 1533 and 1392 cm⁻¹ were shifted to 1517 cm⁻¹ and to 1386 cm⁻¹ respectively; the low intensity vibration band from 1309 cm⁻¹ was shifted and appears as a shoulder at 1314 cm⁻¹, the medium intensities sharp vibration bands from 1237 and 1158 cm⁻¹ shifts to 1240 and 1162 cm⁻¹, respectively. The outlined changes of some vibration bands, especially those characteristics of the OH and COOH groups, suggest the existence of some weak hydrogen bond- and electrostatic-type interactions between the two components of the CUC-WPC_SD dispersion system.

Scanning electron microscopy (SEM)

Morphological analysis of CUC-WPC_SD sample, presented in Figure 5, revealed an amorphous state without any crystalline features. In terms of shape, the sample is highly homogeneous, the particles adopting a donut-like form.



Figure 5. SEM images of the particle size distribution in the CUC-WPC_SD sample

The particle size distribution showed a relatively large interval, varying between 0.17 μ m to 4.9 μ m and having a mean size of 1.5 μ m. Despite the wide range, the size distribution follows a Gaussian curve (Figure 6).

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Figure 6. The Gaussian curve of particle size distribution

Powder dissolution measurements

The solid dispersion system CUC-WPC_SD was subjected to the dissolution experiment in deionized water and in aqueous buffer at 1.8 and 8 physiologically pH values, respectively. The curcumin concentration in the CUC-WPC_SD solution was monitored over time, by recording the absorption intensity for a period of 160 min (Figure 7). On figure 7 the dissolution profile of raw curcumin, practically insoluble in water, was also represented.



Figure 7. Variation of curcumin concentration in aqueous solutions of CUC-WPC_SD compared with the solubility of pure curcumin

The dissolution profile of CUC contained in the CUC-WPC_SD system in water (concentration of CUC (μ g/mL) as a function of time (min)), shows the maximum concentration of ~60 μ g/mL almost instantaneously, after which a linear decreasing trend in the curcumin concentration in solution is manifested, due to CUC precipitation over time.

In aqueous buffer with pH=1.8 (similar with gastric liquid) the maximum concentration of CUC reaches a value of ~55 μ g/mL. In the first 60 minutes of the dissolution experiment a linear decreasing tendency to a value of 45 μ g/mL was observed, after which there is a sudden decrease of the amount of dissolved curcumin.

The dissolution profile in buffer with pH=8 (similar to the pH of the intestinal environment) shows instantaneously dissolution of CUC to the maximum concentration of ~70 μ g/mL, after which there is a linear decreasing tendency due to CUC precipitation. After 160 min, the final CUC concentration was ~35 μ g/mL.

No decomposition of curcumin at all the pH values in the performed dissolution tests was observed.

Cytotoxicity tests

The obtained CUC-WPC_SD dispersion system was tested *in vitro* against skin melanoma (A375) and lung adenocarcinoma (A549) cells. The A375 cells were affected to a higher extent compared to A549 cells (p<0.0001, Figure 8a), aspect indicated also by the IC₅₀ concentration. The results showed that CUC-WPC_SD has an IC₅₀ value of 45.5 µg/mL against A375 cells and 203.6 µg/mL against A549. Compared to the untreated control, A375 cells were significantly more affected (p<0.0001) starting with the concentration of 30 µg/mL, whereas for A549 the concentrations of 200 µg/mL and 300 µg/mL, were the most effective. However, at the lowest concentration tested (10 µg/mL), both cell lines had a proliferative response, with registered values above those of the untreated controls (119% for A549, and 120% for A375, respectively).

The LDH values calculated herein, show no significant difference between the two cell lines (Figure 8b). Compared to the cells treated with Tween 20, no necrosis was observed, however, the negative values registered could indicate a membrane blockage [35].

A direct dose dependent reaction was observed in A375 cells that registered high NO values at concentrations above 30 μ g/mL. The results are consistent with the observed effects on the MTT assay. A dose dependent reaction was observed for A549 cells as well, however the trend was more chaotic (Figure 8c), compared to the skin melanoma cells. Nevertheless, the highest value registered for the NO concentration, was associated with the lowest value registered for the MTT assay, in both cell types.

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Figure 8. Cytotoxicity assays performed on A375 and A549 cells, treated with CUC-WPC_SD. a) MTT assay showing the extend of damages induced in the cells' viability in 24 h. b) LDH assay showing the amount of LDH released in the culture media of the cells treated with the drug for 24 h. c) NO assay revealing the amount of nitric oxide released in the culture media of the treated cells.

The NO concentration has a controversial background, as it can indicate both apoptotic and proliferative effects. This means that at low concentrations detected, the apoptosis could be rescued, but tumor inhibition through apoptosis has also been reported for the same scenario [36]. Therefore, the Griess assay is associated with other biochemical analyses such as LDH and MTT, considered for this study, to be able to correlate the events observed. Herein, for both A375 and A549 cells a moderate negative correlation (r = -0.7) was observed between the viability and NO assay test results. This indicates that at high concentrations (starting with 30 μ g/mL drug concentration), the inhibition of the cancer cells tested was achieved through apoptosis, rather than necrosis.

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Similar results were obtained in other previous studies. By testing nano-encapsulated curcumin with whey protein against different cancer cell lines (colon, prostate, and pancreatic cancers), a better inhibitory capacity was observed compared to pure curcumin [25]. When polyethylene glycol was used to enhance the solubility of curcumin, a strong inhibitory capacity against multiple pancreatic cancer cells was obtained [37]. The IC₅₀ values of curcumin alone ranged from 7 to 18 μ M, whereas for the PEG-loaded curcumin, the values decreased to more than half. Also, the incorporated curcumin in catanionic lipid nano-systems was tested for its effect against Lewis lung cancer, the IC₅₀ value of curcumin nano-systems (20.2 μ M) was almost half of that of curcumin used alone (39.7 μ M) [38].

CONCLUSIONS

The CUC-WPC_SD solid dispersion system was prepared by spraydrying in 5:1 molar ratio. Combining the XRPD, DSC, FTIR and SEM analysis results highlight the formation of the amorphous material whose components are joined by weak electrostatic interactions, being highly homogeneous, and adopting a donut-like spherical form.

The dissolution process of curcumin from this system shows a maximum solubility value of 70 μ g/mL in aqueous buffer solution at pH=8, similar of intestinal fluid, and the active component being stable in this environment.

The obtained dispersion system manifested better effects against skin melanoma cells, compared to lung adenocarcinoma cells, but in both cases the effect was promising, and through further and more complex analyses the antitumoral potential of CUC-WPC_SD could be exploited. The inhibition of skin melanoma cells (A375) is dose dependent in 24 h. However, at low concentrations (10 μ g/mL) the effect was proliferative, with values above the untreated control. The LDH release was constant at all tested concentrations. The nitric oxide (NO) concentration increased in a dose dependent manner, with values above the negative control (cells treated with Tween 20).

EXPERIMENTAL SECTION

Materials

All chemicals used for CUC-WPC_SD preparation and testing: curcumin, ethanol, propanol, DMEM F-12, L-glutamine, penicillin-streptomycin solution, Tris base, lithium lactate, NAD solution, sulfanilamide, formazan

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salt, and *N*-(1-naphthyl) ethylene-diamine were purchased from Sigma-Aldrich, Germany; the weight protein concentrate (WPC) was from Merck, Germany, and the fetal bovine serum from HyClone, Thermo Fisher Scientific, UK. All these chemicals were used without further purification. The cells used in cytotoxicity studies, A549 (ATCC CCL-185) and A375 (ATCC CRL-1619), were purchased from Lomianki, Poland.

Preparation of CUC-WPC_SD solid dispersion system

The solid starting compounds, CUC (M_W=368.38 g/mol) and WPC (M_W=~18400 g/mol), were weighed for a molar ratio of CUC:WPC of 5:1 (1:10 w/w). Curcumin was dissolved in ethanol and the WPC in distilled water. Before mixing the solutions, the WPC aqueous solution was filtered, the volume was corrected with distilled water for a concentration of \leq 4% in ethanol. The obtained yellow, slightly milky solution was processed using a SF-1500LAB spray dryer, applying the following parameters: T_{in}=165°C and T_{out}=49°C, ventilator frequency 58 Hz, solution flow rate 5 mL/min. In the end a fine, homogeneous yellow-orange powder was obtained.

X-ray powder diffraction (XRPD)

Data collection was acquired with the DIFFRAC plus XRD Commander using a Brucker D8 Advance Diffractometer with the tube set at 40 kV and 40 mA, equipped with a germanium (1 1 1) monochromator, used to obtain Cu K α_1 radiation and a LYNKSEYE detector. The measurements were recorded in Bragg-Brentano geometries – variant in reflections- in 3-35° 2 θ range, employing a scan rate of 0.02°/s.

Differential scanning calorimetry (DSC)

Thermal measurements were performed with a DSC-60 Shimadzu differential scanning calorimeter in standard aluminum crimped pans as sample holders and alumina as reference sample. Samples have been analyzed in the 20-350°C temperature range under dry nitrogen flow (3.5 L·h⁻¹) with a 10°C·min⁻¹ heating rate. For data collection and analysis, the Shimadzu TA-WS60 and TA60 2.1 software were employed. The DSC calorimeter was calibrated with reference standards of zinc and indium.

Fourier-Transform Infrared Spectroscopy (FTIR)

FT-IR spectra were obtained using a JASCO 6100 FTIR spectrometer in the 4000 to 400 cm⁻¹ spectral domain, with a resolution of 4 cm⁻¹ by employing the KBr pellet technique. Each sample has been dispersed in about 300 mg of anhydrous KBr and the resulting powder was ground in an agate mortar. The pellet was obtained by pressing the ground mixture into an evacuated die. The spectra were collected and analyzed with Jasco Spectra Manager v.2 software.

Scanning electron microscopy (SEM)

Samples provided for SEM analysis were prepared in the same fashion, briefly. The spray-dried samples were spread over the double-sided conductive tape (12 mm) fixed on aluminum stubs and coated with a 10 nm layer of gold. Image acquisition was conducted on a Hitachi SU8230 High Resolution Scanning Electron Microscope equipped with a cold field emission gun and a 80 X-Max system from Oxford Ins. for EDS analysis. For this analysis the microscope was operated at 30 kV in high mag. mode. Approximately 85% of the carbon disk was scanned to give a realistic overview of the sample and only a few representative areas were captured.

Powder Dissolution Experiments

The absorbance values for curcumin and CUC-WPC SD system in deionized water and buffer solutions were detected by a pION µDISS Profiler apparatus, consisting of an integrated diode array spectrophotometer connected to a fiber optic UV probe located directly in the reaction vessel and is able of measuring the concentration as a function of time without having to filter the solution. Measurement of dissolution kinetics and equilibrium solubility was carried out at 470 nm, where the used protein has no absorption, and the concentration of curcumin was calculated by means of a standard curve. In a typical experiment, 10 mL of solvent (deionized water, pH 5.8 or buffer solutions with pH 1.8 and 8.0) was added to a flask containing 10 mg of sample, and the resulting mixture was stirred at 25 °C and 600 rpm. The dissolution experiment with 160 min monitoring time was carried out in triplicate. The calibration curve was performed at 470 nm, a value at which WPC does not show an absorption signal, using a solution of curcumin in ethanol (Figure 9). The linear equation (y = 0.0331x + 0.0149) derived from the calibration curve was used to calculate the amount of dissolved CUC during the dissolution process.

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Figure 9. Calibration curve of curcumin concentration in ethanol

Cytotoxicity experiments

Cell culture. CUC-WPC_SD was tested in vitro against lung adenocarcinoma A549 (ATCC CCL-185) and skin melanoma A375 (ATCC CRL-1619) cells. A549 cells were maintained in DMEM F-12 media, supplemented with 10% fetal bovine serum, and A375 cells was kept in DMEM F-12 media supplemented with 10% fetal bovine serum, 1% L-glutamine and 1% penicillin-streptomycin solution. Both cell lines were grown in 25 mL flasks, at 37°C and 5% CO₂ in humified atmosphere.

Treatment. Three types of biochemical analyses were performed according to the ISO 10993-5 standard. Briefly, at an 80% confluence, the cells were transferred in 96 well plates and left for 24 h to attach. After this period, the treatment with CUC-WPC_SD was applied in increasing concentrations (10 – 300 μ g/mL) for an additional 24 h.

Cell viability was assessed through the MTT assay and the value at which 50% of the cells is affected (IC₅₀) was calculated. The media from the treated cells was analysed for lactate dehydrogenase release (LDH assay) and for nitric oxide production (NO Griess assay). The work flow was as follows: 150 μ L of LDH regent (Tris base 20 mM, lithium lactate, and NAD solution) was mixed with 50 μ L of media, and 100 μ L of NO regent (sulphanilamide and *N*-1-napthylethylenediamine) was mixed with 50 μ L of media. The reagents were left with the media for 10 min in dark, at room temperature, after which the absorbances were read at 630 nm and 490 nm for LDH, and 548 nm for NO.

The cells remaining in the wells were incubated for an additional 1.5 h with 100 μ L MTT solution. Forward, the formazan salt was dissolved with acidified propanol for 5 min, and the absorbance was read at 630 nm and 550 nm. The BioTek plate reader, coupled with Gen5 Software was used (Epoch, Germany), and each plate contained a positive untreated control, a negative control (cells treated with 2% Tween 20), and vehicle controls (media mixed with reagents and CUC-WPC_SD, without cells).

Statistical analyses. Each experiment had six replicates and the mean was calculated from at least three independent experiments. One-way ANOVA and Students test were performed using the OriginPro 9.3 Software, and P values of ≤ 0.05 were considered significant.

ACKNOWLEDGMENTS

The authors acknowledge financial support from the Ministry of Research and Innovation–MCI, Operational Program Competitiveness, POC Project 18/01.09.16, SMIS Code 105533.

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