





Enhanced Solubility and Anticancer Potential of Mansonone G By β-Cyclodextrin-Based Host-Guest Complexation: A Computational and Experimental Study

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Abstract: Mansonone G (MG), a plant-derived compound isolated from the heartwood of *Mansonia gagei*, possesses a potent antitumor effect on several kinds of malignancy. However, its poor solubility limits the use for practical applications. Beta-cyclodextrin (β CD), a cyclic oligosaccharide composed of seven (1-+4)-linked α -D-glucopyranose units, is capable of encapsulating a variety of poorly soluble compounds into its hydrophobic interior. In this work, we aimed to enhance the water solubility and the anticancer activity of MG by complexation with β CD and its derivatives (2,6-di-*O*-methyl- β CD (DM β CD) and hydroxypropyl- β CD). The 90-ns molecular dynamics simulations and MM/GBSA-based binding free energy results suggested that DM β CD was the most preferential host molecule for MG inclusion complexation. The inclusion complex formation between MG and β CD(s) was confirmed by DSC and SEM techniques. Notably, the MG/ β CDs inclusion complexes exerted significantly higher cytotoxic effect (~2-7 fold) on A549 lung cancer cells than the uncomplexed MG.

Keywords: beta-cyclodextrins; inclusion complex; mansonone G; molecular dynamics simulation; lung cancer

1. Introduction

Cancer is a major public health issue and is ranked as the second leading cause of mortality worldwide following cardiovascular disease [1]. Several plant-derived compounds containing naphthoquinone (NQ) moiety such as beta-lapachone, plumbagin, and shikonin have been reported to exert a superior antiproliferative activity [2–4]. Mansonones, group of *ortho*-NQ-containing compounds, are the major bioactive constituents of the diverse plant genera, including *Mansonia*, *Hibiscus*, and *Thespesia* [5]. Previous studies revealed that mansonones D, E, F, and H possess the

anticancer effect on several kinds of solid tumors [6–8]. In addition, the novel derivative of mansonone F exhibited 20-fold stronger DNA topoisomerase II inhibitory activity than the chemotherapeutic drug etoposide [9].

Among the eight different mansonones (mansonones A–H) [10], mansonone G (MG, Figure 1A) is the major product isolated from the heartwood of *Mansonia gagei* Drumm., Sterculiaceae family found in Thailand [11]. Many lines of evidence have shown that MG exhibits a potent anticarcinogenic effect on various types of malignancies, e.g., ovarian (A278, IC₅₀ of 10.2 μ M), colorectal (HCT116, IC₅₀ of 63.4 μ M), liver (HepG2, IC₅₀ of 36.3 μ M and Huh-7, IC₅₀ of 25.9 μ M), breast (MCF-7, IC₅₀ of 23.0 μ M), and cervical (HeLa, IC₅₀ of 18.8 μ M) cancer cell lines [5,12,13], and it has been documented to significantly inhibit the activity of P-glycoprotein efflux pump [5]. Recently, the butoxy MG has shown to potentially induce cell apoptosis in human lung cancer cell lines A549 and H1975 by inhibiting STAT3 and Akt signaling pathways [14]. Even though MG could serve as a promising anticancer agent, its poor water solubility (1.7 mg/L at 30 °C) leads to a limited use for pharmaceutical and medicinal applications.

Cyclodextrin (CD), a naturally occurring cyclic oligosaccharide produced by CD glucanotransferase (CGTase)-catalyzed starch degradation [15], has a unique structure representing a hydrophobic inner cavity and a hydrophilic outer surface. Consequently, CD can potentially enhance the solubility, stability, and pharmacological properties of many lipophilic guest molecules [16,17] through the formation of an inclusion complex driven mainly by van der Waals (vdW) interactions [18]. The common types of CD are alpha (α), beta (β), and gamma (γ) CDs formed by six, seven, and eight α -D-glucopyranose units, respectively. Among the three CDs, β CD (Figure 1B) is the most commonly used in many pharmaceutical purposes due to its low price, easy synthetic access, and structural orientation suitable for inclusion complex generation [19,20]. However, the low water solubility (18.5 mg/mL at 25 °C) and nephrotoxicity of β CD limit its use for practical applications [21].

In recent years, the use of β CD derivatives (e.g., methylated (M) and 2-hydroxypropylated (HP) β CDs) for host-guest encapsulation has gained much attention in an attempt to improve the water solubility and to reduce the limitations of natural β CD. The M β CD (e.g., 2,6-di-O-methyl- β CD; DM_βCD) and the HP_βCD derivatives exhibit higher aqueous solubility and lower toxicity than the unmodified β CD [21–23]. Several experimental and theoretical reports (both classical molecular dynamics (MD) and density functional theory (DFT)) have demonstrated that the water solubility, chemical stability, and anticancer activity of poorly soluble molecules are significantly improved by complexation with DMBCD and HPBCD derivatives [24-28]. However, the information on the inclusion complexes of MG with β CDs has never been previously reported. Therefore, the present study aimed to enhance the physical and biological properties of MG using β CD and its derivatives $(DM\beta CD and HP\beta CD)$ based on working hypothesis that the anticancer activity of MG may be positively affected by β CDs. The all-atom MD simulations and binding free energy calculations were firstly applied (i) to search for the suitable β CD encapsulating molecule for MG and (ii) to investigate the atomistic insights into host-guest complexation and its underlying interactions. Subsequently, the phase solubility study between MG and β CDs was experimentally verified, and the physical and chemical characterization techniques were used to confirm the inclusion complex formation. Finally, the resulting MG/ β CDs complexes were subjected to evaluate the antitumor activity toward A549 human lung cancer cell line in comparison to the MG alone.

2. Materials and Methods

2.1. Computational Details

The 3D structures of MG and all the investigated β CD analogs (β CD, DM β CD, 2HP β CD, 6HP β CD, and DHP β CD) were taken from previous studies [29–31]. It should be noted that (*i*) the hydroxyl group of MG is protonated (neutral form) based on NMR data from previous studies [11,32] and (*ii*) the commercially available HP β CD is a mixture of various degree of substitution (DS), while the models we built in this study were fully (DHP β CD; DS = 14) and partially (2HP β CD and 6HP β CD; DS = 7) substituted HP β CDs. The inclusion complexes were generated in Accelrys

Discovery Studio 2.5 (Accelrys Software Inc., San Diego, CA, USA) by docking protocols using CDOCKER module. The partial charges of MG molecule were created as per the standard procedures [33–35]. The Glycam-06 [36] and general AMBER force fields [37] were applied on β CDs and MG, respectively. Using a truncated octahedral box, the TIP3P water molecules were used to solvate around MG/ β CDs complexes with a spacing distance of 15 Å. After that, the added water molecules were minimized using 1000 steps of steepest descent and continued by 3000 steps of conjugated gradient. Lastly, the whole model was optimized as per the same methods.

Under periodic boundary condition with a time step of 2 fs, each solvated inclusion complex was heated up from 10 K to 303 K for 60 ps, and followed by all-atom MD simulations (*NPT* ensemble, temperature of 303 K and pressure of 1 atm) using AMBER16 for 90 ns in triplicate. All chemical bonds involving hydrogen were constrained using the SHAKE algorithm [38]. The particle mesh Ewald method [39] was employed to treat charge-charge interactions with a cutoff of 12 Å. The molecular mechanics/generalized Born surface area (MM/GBSA)-based binding free energy (ΔG_{bind} , MM/GBSA) calculation [40] was used to estimate the binding affinity of all the studied inclusion complexes.

2.2. Experimental Part

2.2.1. Chemical Reagents

MG was extracted from *Mansonia gagei* Drumm according to the previous study [11]. βCD (MW 1134.98) was obtained from Acros Organics (Morris Plains, NJ, USA), whereas the Heptakis(2,6-di-O-methyl)-β-cyclodextrin (DMβCD; MW 1331.36), (2-Hydroxypropyl)-β-cyclodextrin (HPβCD; MW 1396; average DS is 0.5–1.3 unit of 2-hydroxypropyl group per glucose unit), dimethyl sulfoxide (DMSO), and 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT) were purchased from Sigma-Aldrich (St. Louis, MO, USA).

2.2.2. Cell lines and Culture

A549 human lung cancer cell line was obtained from Dr. Apiwat Mutirangura (Chulalongkorn University, Thailand). A549 cells were cultured in Dulbecco's modified Eagle's medium (DMEM; Gibco, Grand Island, NY, USA) containing 10% fetal bovine serum (FBS; Gibco), 100 U/mL penicillin, and 100 μ g/mL streptomycin (Gibco) and were maintained at 37 °C in a humidified 5% CO₂ atmosphere.

2.2.3. Phase Solubility Study and Evaluation of Thermodynamic Parameters

Phase solubility study was conducted by following the method of Higushi and Connors [41]. Briefly, an excess amount of MG was added to β CD(s) solutions (0–10 mM). The mixtures were shaken (250 rpm) at 30 °C, 37 °C, and 45 °C for 72 h. Subsequently, the suspensions were centrifuged at 12,000 rpm for 15 min. The amount of MG presenting in solution was measured by UV-Vis spectroscopy at 415 nm. The Equation (1) was used to calculate the apparent stability constant (*K*_c), where *S*₀ is y-intercept.

$$K_c = \frac{Slope}{[S_o(1-slope)]} \tag{1}$$

The Van't Hoff plot based on Equation (2) [17] was used to identify the thermodynamic properties, including enthalpy (ΔH) and entropy (ΔS) changes of inclusion complexation, whilst the experimental Gibbs free energy ($\Delta G_{\text{bind, exp}}$) was obtained from Equation (3).

$$\ln K_c = -\frac{\Delta H}{RT} + \frac{\Delta S}{R} \tag{2}$$

$$\Delta G_{\text{bind, exp}} = \Delta H - T \Delta S \tag{3}$$

A 1:1 stoichiometric ratio of MG/ β CD(s) inclusion complex was prepared by the freeze-drying method. Each accurately weighed compound (73.28 mg MG, 340.49 mg β CD, 399.41 mg DM β CD, and 418.80 mg HP β CD) was dissolved in deionized water (30 mL), then the mixture was magnetically stirred at room temperature for 24 h. After that, the suspension was centrifuged (12,000 rpm for 15 min) and filtered through a 0.45 µm filter. The obtained solution was then frozen overnight at -80 °C and subsequently lyophilized using LYO-LAB (Lyophilization Systems Inc., USA) for 3 days. In addition, a 1:1 molar ratio of MG and β CD(s) was physically mixed at room temperature to obtain the physical mixtures, which were used for comparison. The resulting freeze-dried powders and physical mixtures were stored in desiccator for further use.

2.2.5. Determination of Solubility

Excess quantity of pure MG and its inclusion complexes was added to 1 mL of deionized water. The mixture was continuously stirred at 30 °C for 24 h. Subsequently, the suspension was filtered through a 0.45 μ m filter, and the solubility of MG was characterized by measuring the absorbance at a wavelength of 415 nm.

2.2.6. Inclusion Complex Characterization

Differential Scanning Calorimetry (DSC)

The thermal behavior of the MG, β CD, DM β CD, MG/ β CD, and MG/DM β CD was characterized using NETZSCH DSC 204F1 Phoenix. Each solid sample (~1–2 mg) was heated from 25 °C to 300 °C in aluminum pans at a rate of 10 °C/min.

Scanning Electron Microscope (SEM)

The surface morphology of MG, β CD, DM β CD, the freeze-dried inclusion complexes, and the physical mixtures was analyzed using a Scanning Electron Microscope (JEOL JSM-IT500HR). Samples were coated with a thin layer of gold in vacuum before viewing under 300 times magnification. Observations were performed using an accelerating voltage of 10 kV.

2.2.7. Cytotoxicity of MG toward Lung Cancer Cells

Cell viability was evaluated using MTT assay. A549 cells were seeded into 96-well plates at a density of 3×10^3 cells/well. After overnight incubation, cells were treated with logarithmic concentrations (1, 3, 10, 30, and 100 μ M) of MG, MG/ β CD, MG/DM β CD, and MG/HP β CD for 48 h. Note that the amount of treated MG in free form and in complexes is equivalent. After that, MTT solution (5 mg/mL) was added to each well and then incubated at 37 °C for another 4 h. Subsequently, the culture medium was withdrawn and 150 μ L of DMSO solution was added to dissolve formazan crystals. Finally, the absorbance was measured at 570 nm using UV-Vis spectrophotometer (Thermo Scientific, Vantaa, Finland). The half-maximal inhibitory concentration (IC₅₀) was calculated using GraphPad Prism 7 software.

2.2.8. Statistical Analysis

The quantitative data are expressed as mean \pm standard error of mean (SEM) of three independent experiments. Differences between groups were determined using one-way analysis of variance (ANOVA) followed by a Newman-Keuls post hoc test. The *p* value of ≤ 0.05 was considered statistically significant.

(A)

Aromatic ring (A-ring)



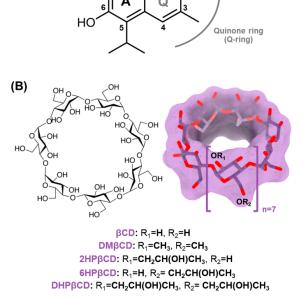


Figure 1. (**A**) 2D structure of MG. (**B**) 2D and 3D structures of β CD, where the functional substitutions used in this study are shown below.

3. Results and Discussion

From docking study, MG could form two possible inclusion complexes with β CDs through its aromatic ring (A-ring, form I) or quinone ring (Q-ring, form II) insertion into the hydrophobic cavity. However, after performing all-atom MD simulations for 90 ns in triplicate on both forms (altogether 30 independent simulations), only form I of MG/ β CD, MG/DM β CD, and MG/2HP β CD is likely stable, whilst the dissociation of inclusion complex is detected in the form II of all systems as well as the form I of MG/ β CD and MG/DHP β CD. The latter finding is consistent well with previous study showing that the HP modification on β CD at C6-position (DS = 7) led to the dissociation of the encapsulated mansonones E and H from hydrophobic inner cavity [42]. Therefore, the MG/ β CD, MG/DM β CD (DS = 14), and MG/2HP β CD (DS = 7) complexes in form I were further focused in this study.

3.1. Ligand Mobility Inside *βCD's* Hydrophobic Cavity

The dynamics behavior of the encapsulated MG inside hydrophobic interior of β CD(s) along the simulation time was studied using the calculations of the distance between the center of mass (C_m) of each ring of MG (A-ring (black) and Q-ring (grey)) and the C_m of β CD without taking into account the functional substituents. The obtained results are shown in Figure 2A, whereas the representative binding mode of MG/ β CD(s) taken from the final MD snapshot is depicted in Figure 2B. The horizontal grey box ranging from –3.95Å to 3.95Å (~7.9Å) represents the positions of the primary (narrow) and secondary (wider) rims of native β CD, respectively [43]. In the case of MG/ β CD and MG/2HP β CD systems, MG stably positions nearby the wider rim of β CD(s) and preferentially inserts the A-ring inside the hydrophobic inner cavity. By considering the MG/DM β CD models, MG rapidly moves toward the narrow rim of DM β CD after the heating step and subsequently positions in this orientation until the end of the simulations.

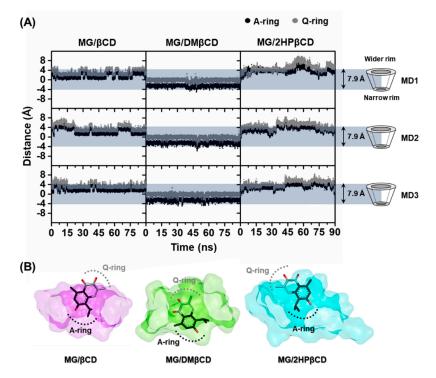


Figure 2. (**A**) The distance between C_m of MG ring (A/Q) and C_m of β CD of all studied inclusion complexes for three independent MD simulations (MD1-3). (**B**) The binding orientation of MG inside β CD (violet), DM β CD (green), and 2HP β CD (cyan) cavities drawn from the last MD snapshot.

3.2. BCDs Conformations Upon MG Binding

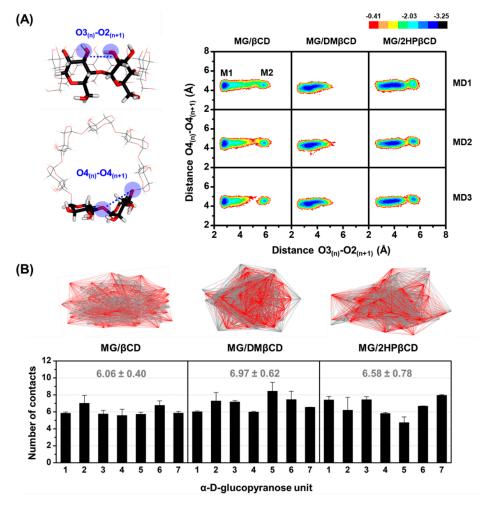
The β CD(s) conformational changes upon MG encapsulation were investigated by calculating (*i*) the distance of the secondary hydroxyl groups on the wider rim of β CDs (O3_(n)–O2_(n+1), dO₃₋₂), corresponding to a possibility of intramolecular hydrogen bond (H-bond) generation (dO₃₋₂ of \leq 3.5 Å), and (*ii*) the distance of glycosidic oxygen atoms (O4_(n)–O4_(n+1), dO₄₋₄) (Figure 3A, left). Afterward, the Equation (4) was used to transform the distributions of these two parameters into the free energy value, F(*x*,*y*):

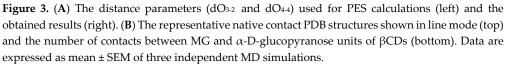
$$\mathbf{F}(x,y) = -k_{\rm B}T \log[P(x,y)] \tag{4}$$

where k_{B} is the Boltzmann constant, *T* is the temperature (303 K), and P(x,y) is the probability of dO₃₋₂ (x) and dO₄₋₄ (y). The potential energy surface (PES) is shown in Figure 3A, in which the M1 area is the most stable form of the β CD(s), whilst the M2 region represents the rotation of the α -D-glucopyranose units.

As compared to the unbound form of β CDs exhibiting three local minima areas: M1 (dO₃₋₂ of ~3.0–4.5 Å and dO₄₋₄ of ~4.3–4.7 Å), M2 (dO₃₋₂ of ~5 Å and dO₄₋₄ of ~4.5–5.2 Å), and M3 (dO₃₋₂ of ~5 Å and dO₄₋₄ of ~6 Å) [30], the molecular encapsulation of MG toward DM β CD and 2HP β CD dramatically induces the stable conformation of β CD(s) by enhancing the formation of intramolecular H-bonds on the wider rim, as evidenced by the obviously increased population in M1 region (dO₃₋₂ of ~2.5–4.5 Å and dO₄₋₄ of ~4.1–4.9 Å, Figure 3A). In addition, the M3 region with both lengthened distances, which was only detected in the free form of β CDs, is completely disappeared in all MG-bound systems. These H-bond-operated conformational changes of β CD(s) upon the ligand binding correlate well with previously published research [44]. However, the distortion of glucopyranose units represented by M2 area is found in MG/ β CD and MG/2HP β CD systems, but not in MG/DM β CD models, indicating that MG/DM β CD is the most stable complex, which is consistent with Δ Gbind, MM/GBSA calculations (Table 1) as well as the experimental *K*_c and Δ Gbind, exp values (Tables 2 and 3) as discussed later.

We further characterized the native contact points within 3 Å between MG and β CDs during the last 20-ns MD simulations. The obtained results (Figure 3B) reveal that MG/DM β CD inclusion complex displays the highest number of contacts per α -D-glucopyranose units (6.97 ± 0.62) followed by MG/2HP β CD (6.58 ± 0.78) and MG/ β CD (6.06 ± 0.40) complexes, respectively, as clearly shown by the representative 3D contact structures (Figure 3B, top) demonstrating that MG/DM β CD exhibits the most compact feature. Altogether, the native-contact-driven structures consistently support the PES calculations.





3.3. Solvent Accessibility Toward Inclusion Complexes

The effect of water accessibility on MG/ β CD(s) inclusion complex formation was characterized by solvent accessible surface area (SASA) calculations using MG as the atomic radii for solventexposed area. The entire SASA results are depicted in Figure 4A, whereas the average SASA values taken from the last 20-ns MD simulations are illustrated in Figure 4B.

Data in Figure 4A indicate that the SASAs of MG/ β CD and MG/ $DM\beta$ CD systems are more stable than those of MG/2HP β CD complexes. In the case of MG/ β CD, the SASAs fluctuate in the range of ~150–250 Å², whereas the SASAs of MG/ $DM\beta$ CD are relatively steady at ~150 Å² along the simulation times. By considering the MG/2HP β CD models, the SASA values considerably fluctuate (~200–300 Å²) at 40–70 ns and subsequently decrease to ~150 Å² after 70 ns (except MD1). The average SASA values over the last 20 ns from three independent simulations in Figure 4B reveal that the complexation of MG with DM β CD significantly decreases the water accessibility toward the MG molecule inside hydrophobic inner cavity as compared to the MG/ β CD and MG/2HP β CD (* $p \le 0.05$). These observations correlate well with previous study [45] demonstrating that the lowest solvent exposed inclusion complex displayed the highest binding affinity. However, the SASAs of MG/ β CD and MG/2HP β CD are not significantly different, suggesting that DM β CD is the most preferred encapsulating agent for MG, which is in good correlation with MM/GBSA-based free energy calculations as discussed in the next section.

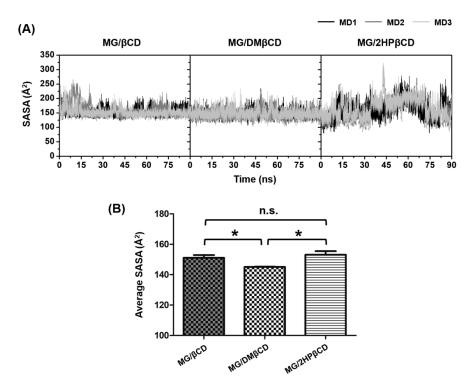


Figure 4. (**A**) SASA of MG/ β CD, MG/DM β CD, and MG/2HP β CD for three different MD runs. (**B**) The average SASA over the last 20-ns MD simulations for each inclusion complex. Data are expressed as mean ± SEM (n = 3). n.s., not significantly different. * $p \le 0.05$.

3.4. Binding Free Energy of Inclusion Complexes

To estimate the binding affinity of the inclusion complexes, we applied MM/GBSA [46] calculation on each MG/ β CD(s) using 300 snapshots extracted from the last 20-ns MD simulations. The average $\Delta G_{\text{bind, MM/GBSA}}$ and its energy components, vdW (ΔE_{vdW}) and electrostatic (ΔE_{ele}) energies, for MG/ β CDs are summarized and compared in Table 1. As expected, due to the poor solubility of MG, the host-guest complexation in gas phase is driven predominantly by vdW interactions (~ -27 to -29 kcal/mol). Similarly, the summation of $\Delta G_{\text{solv,non-polar}} + \Delta E_{\text{vdW}}$ energies (~ -30 to -32 kcal/mol) is much lower than the $\Delta G_{\text{solv,polar}} + \Delta E_{\text{ele}}$ terms (~8 to 10 kcal/mol), indicating the vdW forces play a pivotal role in the formation of MG/ β CDs in an aqueous environment. These findings strongly correlate with those reported for other hydrophobic ligands in complex with β CDs [47–49].

Several studies have shown that the methyl and hydroxypropyl modifications on β CD can significantly enhance the stability of many lipophilic guest molecules [50–52]. In good agreement with these reports, our results demonstrate that the $\Delta G_{\text{bind, MM/GBSA}}$ of MG in complex with modified β CDs is significantly lower than that of MG in complex with unsubstituted β CD, which can be ranked in the order of MG/ β CD (–2.34 ± 0.35 kcal/mol) > MG/2HP β CD (–3.35 ± 0.14 kcal/mol, *) > MG/DM β CD (–5.73 ± 0.04 kcal/mol, ***). Taken altogether, all structural and energetic analyses convince that DM β CD analog is the most suitable host for MG encapsulation.

Table 1. The average Δ*G*_{bind}, MM/GBSA</sub> and its energy components (kcal/mol) of MG/βCDs inclusion complexes. Data are expressed as mean ± SEM from three independent MD simulations. Δ*E*_{MM}, molecular mechanics energy; Δ*G*_{solv}, solvation free energy comprising polar (Δ*G*_{solv,polar}) and non-polar (Δ*G*_{solv,polar}) terms; Δ*S*, entropy. * $p \le 0.05$ and *** $p \le 0.001$ vs. MG/βCD.

Energy Component (kcal/mol)	MG/βCD	MG/DMβCD	MG/2HPβCD
$\Delta E_{ m vdW}$	-27.61 ± 1.19	-29.19 ± 0.29	-28.59 ± 0.64
$\Delta E_{ m ele}$	-11.79 ± 0.40	-4.43 ± 0.19	-6.40 ± 0.90
$\Delta E_{ m MM}$	-39.40 ± 1.58	-33.62 ± 0.26	-34.99 ± 0.26
$\Delta G_{ m solv,polar}$	22.14 ± 0.91	12.72 ± 0.13	17.17 ± 0.54
$\Delta G_{ m solv,non-polar}$	-2.83 ± 0.04	-2.98 ± 0.01	-3.10 ± 0.02
$\Delta G_{ m solv}$	19.31 ± 0.86	9.73 ± 0.12	14.06 ± 0.57
$\Delta G_{ m solv,polar}$ + $\Delta E_{ m ele}$	10.35 ± 0.53	8.29 ± 0.19	10.77 ± 0.47
$\Delta G_{ m solv,non-polar}$ + $\Delta E_{ m vdW}$	-30.44 ± 1.24	-32.18 ± 0.31	-31.69 ± 0.66
$T\Delta S$	-17.74 ± 0.36	-18.16 ± 0.13	-17.57 ± 0.42
ΔG bind, MM/GBSA	-2.34 ± 0.35	-5.73 ± 0.04 (***)	-3.35 ± 0.14 (*)

3.5. Phase Solubility Study and Thermodynamic Parameters

As the molecular modeling results suggest that DM β CD derivative is the most preferential host for MG, we further confirmed our findings by conducting the experimental phase solubility study and Van't Hoff-based thermodynamic parameter evaluation.

Phase solubility diagrams of all studied inclusion complexes at temperatures 30, 37, and 45 °C are summarized in Figure 5, where the corresponding K_c values are given in Table 2. The obtained results show that the increased β CD(s) concentrations can enhance the solubility of MG in a liner manner (ranked in the order of DM β CD >> 2HP β CD > β CD, see Table S1), indicating that the stoichiometric ratio between MG and β CD(s) is 1:1 (A_L type) [41]. By considering the stability of all studied inclusion complexes at 30 °C, the highest K_c value is detected in MG/DM β CD (2245 M⁻¹) followed by MG/HP β CD (684 M⁻¹) and MG/ β CD (562 M⁻¹), respectively. Moreover, the solubility of MG is dramatically increased up to ~7 times, ~28 times, and ~10 times by complexation with β CD, DM β CD, and HP β CD, respectively (Table 4). These observations are consonant well with several reports demonstrating that β CD derivatives, especially DM β CD, could significantly enhance the stability and solubility of hydrophobic guest molecules better than natural β CD [25,53,54]. However, the increased temperature remarkably reduces the stability of all investigated complexes, in a manner similar to previous works [55,56].

Using Van't Hoff plot (see Supporting Information, Figure S1), the obtained thermodynamic values, i.e., ΔH , ΔS , and $\Delta G_{\text{bind, exp}}$, are summarized in Table 3. The negative ΔH (–20.77, –23.27, and –17.39 kcal/mol for MG/ β CD, MG/DM β CD, and MG/HP β CD, respectively) suggests an exothermic process. Moreover, the formation of all inclusion complexes is spontaneous, as evidenced by the negative sign of $\Delta G_{\text{bind, exp}}$ values at 30 °C (–3.71, –4.56, –3.85 kcal/mol for MG/ β CD, MG/DM β CD, and MG/HP β CD, respectively). Notably, the trend of MM/GBSA-based free energies strongly agrees with $\Delta G_{\text{bind, exp}}$ values, indicating that MM/GBSA serves as a suitable method for predicting the binding affinity of MG/ β CDs inclusion complexes, which is in good agreement with several studies on other host-guest systems [56–60].

Taken together, based on theoretical and experimental investigations, we selected the MG/DM β CD, showing the highest stability, for further structural characterizations in comparison to the MG/ β CD and the free form of MG.

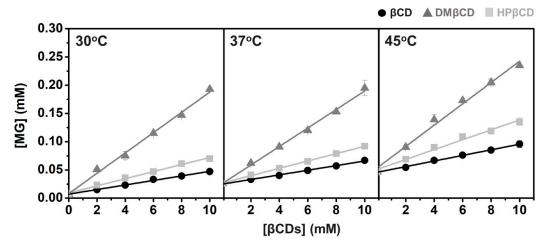


Figure 5. Phase solubility diagram of MG with all studied β CDs in water at 30 °C, 37 °C, and 45 °C. An excess amount of MG was added to β CD(s) solutions (0–10 mM). The mixtures were shaken (250 rpm) at 30 °C, 37 °C, and 45 °C for 72 h. Subsequently, the suspensions were centrifuged at 12,000 rpm for 15 min. The amount of MG presenting in solution was measured by UV-Vis spectroscopy at 415 nm. Data are expressed as mean ± SEM of three independent experiments.

Table 2. *Kc* of MG/ β CDs inclusion complexes at different temperatures. Note that the 95% confidence interval and the other statistic values for the slope and the y-intercept of linear regression are shown in Table S2.

	Stability Constant (Kc, M ⁻¹)			
Temperature (°C)	MG/βCD	MG/DMβCD	MG/HPβCD	
30	562	2245	684	
37	165	643	245	
45	109	358	173	

Table 3. Thermodynamic values for the inclusion complex formation of MG with β CDs derived from Van't Hoff plots (using R of 1.985 × 10⁻³ kcal·mol⁻¹·K⁻¹ and T of 303 K) in comparison to the Δ G_{bind}, MM/GBSA. Note that the 95% confidence interval and the other statistic values for the slope and the y-intercept of linear regression are shown in Table S3.

Thermodynamic Parameter (kcal/mol)	MG/βCD	MG/DMβCD	MG/HPβCD
ΔΗ	-20.77	-23.27	-17.39
$T\Delta S$	-17.06	-18.71	-13.54
$\Delta G_{\text{bind, exp}}$ (30 °C)	-3.71	-4.56	-3.85
$\Delta G_{ ext{bind, MM/GBSA}}$ (Table 1)	-2.34 ± 0.35	-5.73 ± 0.04	-3.35 ± 0.14

Table 4. Solubility of MG and its inclusion complexes at 30 °C.

	Solubility of MG (mg/L)
MG	1.7
MG/βCD	11.5
MG/DMβCD	47.2
MG/HPβCD	17.1

3.6. Inclusion Complex Characterization

3.6.1. Thermal Behavior of MG and Its Inclusion Complexes

The thermal behaviors of the starting materials (MG, β CD, and DM β CD) and the inclusion complexes (MG/ β CD and MG/DM β CD) were characterized in the solid state by DSC analysis (Figure 6). The characteristic endothermic/exothermic peaks of the free compounds are as follows: MG at 208.7 °C and 213.0 °C, β CD at 118.9 °C, and DM β CD at 77.1 °C, 184.9 °C, and 293.9 °C. The broad endothermic peak of β CD detected at 118.9 °C corresponds to the release of water molecules from the hydrophobic cavity [61]. As shown in supplemental Figure S2, the distinct thermal peaks of the free MG were expressed in all the physical mixture products, indicating that the physical mixing method does not provide a real inclusion complex.

As a host-guest complexation process is occurred, the thermal features of the acquired product are dramatically changed [62]. In accordance with these facts, our data in Figure 6 reveal that the thermal peaks of the free MG were totally disappeared in all the freeze-dried inclusion complexes. In addition, in the case of MG/ β CD inclusion complexation, the characteristic broad peak of β CD at 118.9 °C is shifted to 57.2 °C by the influence of the MG-induced water replacement. The similar change of dehydration peak of β CD after ligand complexation was also found by Rajendiran and coworkers [63]. Similarly, the broad endothermic peak (77.1 °C) and the exothermic peaks (184.9 °C and 293.9 °C) of DM β CD are dramatically shifted with a decreased intensity after complexing with MG, in good agreement with previous study [64]. Therefore, the freeze-drying method can successfully generate the new solid phase between MG and β CDs.

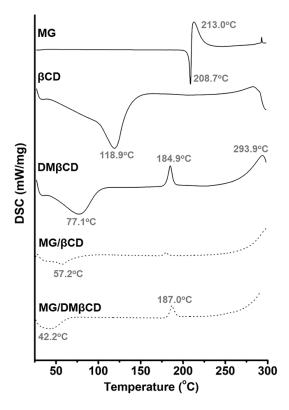


Figure 6. DSC thermogram of MG, β CD, DM β CD, and the freeze-dried inclusion complexes (MG/ β CD and MG/DM β CD). Note that prior to perform DSC, all compounds were dissolved in deionized water and were subsequently lyophilized for 3 days.

3.6.2. Surface Morphological Changes upon Inclusion Complexation

It was evident that the inclusion complex formation dramatically induces the alterations of the particle shape and surface morphology of the resulting products [65–67]; thus, we performed SEM analysis in order to characterize the morphological changes upon host-guest encapsulation. The SEM micrographs at 300 times magnification of all investigated samples are summarized in Figure 7. In the case of the unbound forms, β CD and DM β CD are defined as rod-shaped structure, in accord with previous reports [68,69], whereas MG displays a flake-like feature. Upon molecular complexation, the surface morphology and the particle shape/size of the obtained freeze-dried inclusion complexes MG/ β CD and MG/DM β CD (Figure 7D,E) are totally different from those of the free forms, confirming the successful formation between MG and β CDs.

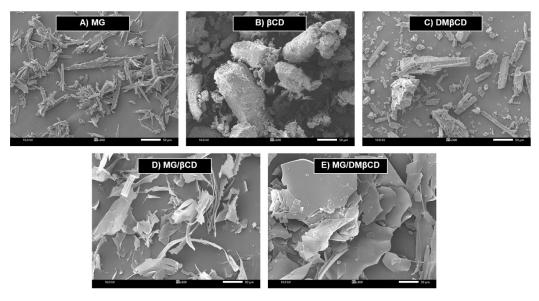


Figure 7. SEM images of (**A**) MG, (**B**) βCD, (**C**) DMβCD, (**D**) MG/βCD, and (**E**) MG/DMβCD at 300 times magnification.

3.7. Cytotoxicity of MG/BCDs Inclusion Complexes toward Lung Cancer

Lung cancer is the first leading cause of cancer-related death globally with its five-year survival rate of only 17.8% [70]. In this study, we evaluated the cytotoxic activity of MG and its inclusion complexes (MG/ β CD, MG/DM β CD, and MG/HP β CD) against A549 human lung cancer cell line using MTT assay. The cell viability (% of control) results are summarized in Figure 8A, whereas the IC₅₀ values (μ M) are shown in Figure 8B. The obtained results reveal that MG, MG/ β CD, MG/DM β CD, and MG/HP β CD decrease cell viability in a concentration-dependent manner, in which MG/HP β CD exhibits the lowest cell viability followed by MG/DM β CD, MG/ β CD, and MG, respectively (Figure 8A).

Multiple lines of evidence have shown that β CD inclusion complexations can potentially enhance the antitumor activity of lipophilic guest compounds [25,67,71]. In correlation with these reports, the present study shows that the inclusion complexes of MG significantly increase the anticancer effect on A549 cells, which can be ranked in the order of MG/HP β CD (IC₅₀ of 5.62 ± 0.40 μ M, ***) > MG/DM β CD (IC₅₀ of 13.45 ± 0.24 μ M, ***) > MG/ β CD (17.63 ± 0.42 μ M, ***) >> MG (42.86 ± 2.09 μ M) (Figure 8B). The enhancement of anticancer activity of MG by complexation with β CDs is likely due to the reason that β CDs can infiltrate into the drug permeation barrier, called unstirred water layer (UWL) consisting of a large number of strong H-bond networks [72], better than the free form of lipophilic MG. It is likely that the introduced hydrophilic HP (HP β CD) and the hydrophobic methyl groups on β CD (DM β CD) could respectively (*i*) enhance the flux of drug though UWL [73] and (*ii*) interact with biological membrane greater than unmodified CD [74]. Therefore, the cytotoxic effect on A549 of MG/HP β CD and MG/DM β CD is higher than that of MG/ β CD. Importantly, the free form of β CD, DM β CD, and HP β CD does not affect the antitumor property of MG/ β CDs inclusion complexes, as evidenced by the cell viability values of >90% (Figure 8C).

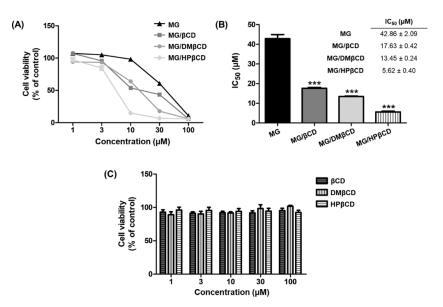


Figure 8. (**A**) Cell viability of A549 human lung cell line after treatment with logarithmic concentrations (1, 3, 10, 30, and 100 μ M) of MG, MG/ β CD, MG/DM β CD, and MG/HP β CD for 48 h. (**B**) The IC₅₀ (μ M) of all investigated compounds. (**C**) Cell viability of the free form of β CD, DM β CD, and HP β CD. Data are expressed as mean ± SEM of three independent experiments. *** $p \le 0.001$ vs. MG.

4. Conclusions

In the present study, we combined theoretical and experimental studies to identify the most suitable β CD host molecule for MG, a promising anticancer agent extracted from *Mansonia gagei*, for improving the aqueous solubility and the antitumor potential. The 90-ns MD simulations in triplicate revealed that MG preferentially positioned inside β CD, DM β CD, and 2HP β CD cavities through its aromatic ring. The MG binding led to the rigidity of β CDs by increasing the intramolecular H-bond formations on the wider rim. Among the three different MG/ β CDs, the ΔG_{bind} values from MM/GBSA calculations and experimental phase solubility study revealed that MG/DM β CD was the most stable complex, as supported by the lowest water accessibility toward MG atomic radii and the highest number of contacts between host and guest molecules. The successful formation of MG/ β CDs was confirmed by DSC and SEM techniques. The anticancer activity of MG toward A549 lung cancer cells was significantly enhanced (~2–7 fold) by complexation with β CDs, especially HP β CD and DM β CD analogs. Altogether, the obtained results confirmed the working hypothesis and demonstrated the good potentiality of β CD derivatives as suitable formulations of MG for further pharmaceutical and medicinal applications.

Supplementary Materials: The following are available online at www.mdpi.com/xxx/s1, Figure S1: Van't Hoff plots of MG/βCDs inclusion complexes., Figure S2: DSC thermogram of MG, βCD, DMβCD, and the physical mixtures MG/βCD and MG/DMβCD. Note that prior to perform DSC, all compounds were not dissolved in deionized water and were not subsequently lyophilized., Table S1: Linear equation of MG/βCDs inclusion complexes at different temperatures derived from Figure 5., Table S2: The 95% confidence interval and the other statistic values for the slope and the y-intercept of linear regression of phase solubility study derived from Figure 5., Table S3: The 95% confidence interval and the other statistic values for the slope and the y-intercept of linear regression of Van't Hoff plot derived from Figure S1.

Author Contributions: P.M. and T.R.; conceptualization. P.M. conducted theoretical and experimental studies. T.C. and P.W.; methodology. M.P. and P.W.; resources. W.C. purified, synthesized, and identified MG using

NMR. P.M., T.R., P.W., and T.C. analyzed the data. P.M. wrote the original manuscript. All authors reviewed and edited the manuscript. T.R.; funding acquisition.

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