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## Encapsulation of an anticancer drug Isatin inside a host nano-vehicle SWCNT: a molecular dynamics simulation

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The use of carbon nanotubes as anticancer drug delivery cargo systems is a promising modality as they are able to perforate cellular membranes and transport the carried therapeutic molecules into the cellular components. Our work describes the encapsulation process of a common anticancer drug, Isatin (1H-indole-2,3-dione) as a guest molecule, in a capped single-walled carbon nanotube (SWCNT) host with chirality of (10,10). The encapsulation process was modelled, considering an aqueous solution, by a molecular dynamics (MD) simulation under a canonical NVT ensemble. The interactions between the atoms of Isatin were obtained from the DREIDING force field. The storage capacity of the capped SWCNT host was evaluated to quantify its capacity to host multiple Isatin molecules. Our results show that the Isatin can be readily trapped inside the volume cavity of the capped SWCNT and it remained stable, as featured by a reduction in the van der Waals forces between Isatin guest and the SWCNT host (at approximately  $-30 \text{ kcal mol}^{-1}$ ) at the end of the MD simulation (15 ns). Moreover, the free energy of encapsulation was found to be  $-34 \text{ kcal mol}^{-1}$  suggesting that the Isatin insertion procedure into the SWCNT occurred spontaneously. As calculated, a capped SWCNT (10,10) with a length of  $30 \text{ \AA}$ , was able to host eleven (11) molecules of Isatin, that all remained steadily encapsulated inside the SWCNT volume cavity, showing a potential for the use of carbon nanotubes as drug delivery cargo systems.

Cancer is the second-leading cause of death in the world<sup>1</sup> and several therapeutic approaches have been developed to obviate the cancer cells from a body, including the use of surgical, radiation therapy and chemotherapy methods<sup>2,3</sup>. Despite the widespread application of chemotherapy in cancer treatment, the drawbacks of using anticancer drugs including their toxic side effects on fast-growing healthy cells, and multi-drug resistance are believed to limit the application of chemotherapy in cancer treatment<sup>4,5</sup>. Nanoscale delivery systems are promising modalities that have a potential to overcome the barriers associated with chemotherapy treatment by delaying drug resistance<sup>6,7</sup>, improving drug release profile into tumor micro-environment<sup>8-11</sup>, facilitating penetration of drugs into cancer cells and tissues<sup>12</sup>, while shielding drugs from premature chemical reactions and/or an

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induced physical tension<sup>13,14</sup> and, most importantly, are able to apportion the required dosage of the drugs via their unique carrier profile<sup>15,16</sup>.

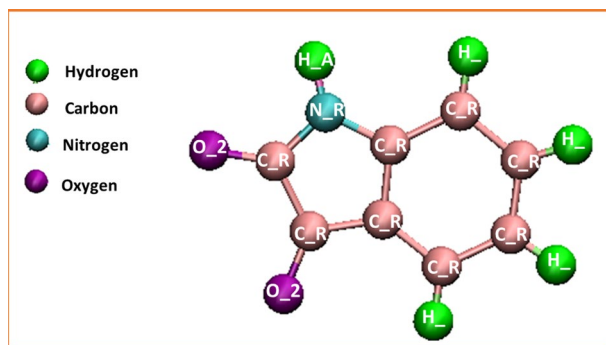
An open-ended or end-capped, carbon nanotubes (CNTs) have attracted a significant attention as possible drug nano-carriers owing to their remarkable features including mechanical robustness and exceptionally large surface area per unit mass ratios, large enough to provide an ample volume for possible encapsulation and adsorption of nano-sized media such as drug molecules, compact storage and delivery characteristics<sup>17</sup>, and a relatively uncomplicated functionalization profile that allows CNTs to be doped or grafted with desired chemical structures and/or molecules<sup>18–20</sup>. Single-walled CNTs (SWCNTs) are  $sp^2$  hybridized allotropes of carbon, similar to fullerenes<sup>21</sup>. The structure of a SWCNT is comprised of 6-membered carbon rings, as in graphite<sup>22</sup>. The cylindrical SWCNTs may have one or both ends capped with a hemisphere of the buckyball or fullerene structure<sup>23</sup>.

A number of theoretical studies were performed recently to evaluate the performance of SWCNTs as possible host nano-vehicles for anticancer drug delivery including the study by Moradnia et al.<sup>24</sup> that evaluated the adsorption function of Gemcitabine (AKA. Gemzar) anticancer drug on the surface of the functionalized SWCNT. The study evaluated the van der Waals (vdW), electrostatic (Elec), and binding free energy energies for the interaction of Gemcitabine employing a molecular dynamics (MD) simulation. The negative value for the free energy and vdW energy confirmed the conformational stability of Gemcitabine on the surface of SWCNTs. Roosta et al.<sup>25</sup> evaluated the changes in the Elec and vdW energies during the encapsulation process of the Cisplatin inside the CNT by using MD simulation. They reported that the vdW contribution to solute–solvent interaction energy and Elec contribution to solute–solvent interaction energy were equal to the values of  $-0.322 \text{ kcal mol}^{-1}$  and  $-0.260 \text{ kcal mol}^{-1}$ , respectively. Maleki et al.<sup>26</sup> studied the conformational attachment of another anticancer drug Doxorubicin (AKA. Adriamycin, Rubex) on a SWCNT. Their calculations for vdW and Elec energies between the host SWCNT and Doxorubicin guest have shown that the value of the Elec energy was zero while the vdW interaction was solely responsible for the adsorption process. Mirsalari et al.<sup>27</sup> studied the adsorption of the anticancer drug Dacarbazine into the CNT by means of MD simulation. They reported that Decarbazine was inserted spontaneously inside the cavity of the CNT so that the vdW energy decreased and reached the value of  $-160 \text{ kJ mol}^{-1}$  at the end of the simulation. Also, the adsorption of other anticancer drugs including Sunitinib (AKA. Sutent, SU11248), Streptozotocin or Streptozocin (INN, USP) (STZ), and Sorafenib (AKA. Nexavar) on functionalized SWCNTs were studied using the MD simulation by Dehnesin et al.<sup>28</sup>. The adsorption of Sorafenib on SWCNTs occurred through the  $\pi$ - $\pi^*$  stacking and hydrogen (H-) bonding, whereas STZ was absorbed exclusively through H-bonding. The interactions between anticancer drug Penicillamine (AKA. Cuprimine) and SWCNT was investigated using the MD simulation by Shaki et al.<sup>11</sup>; their study have shown that the interaction energies had negative values confirmed that the adsorption process of Penicillamine guest on SWCNT was exothermic and spontaneous. The co-adsorption process of Doxorubicin with various dye molecules on/into the SWCNT (30,0) is studied by using MD simulation by Panczyk et al.<sup>29</sup>. They reported that the drug can be released in acidic environment due to the presence of dye molecules facilitating the free energy barrier elimination against release. In another works done by the same team<sup>30</sup>, it was predicted that the Cisplatin can be adsorbed on SWCNT with free energy of  $-25 \text{ kJ mol}^{-1}$ . Wolski et al.<sup>31</sup> investigated the encapsulation of Carmustine anticancer drug inside the SWCNT, then capped it using functionalized magnetic nanoparticles by means of MD simulation. It was observed that the magnetic nanoparticle having diameters larger than  $35 \text{ \AA}$  were able to absorb huge amounts of energy from the magnetic field and cause the release of the drug.

Considering the main concepts understood from the recent theoretical studies<sup>11,26,28</sup> on the use of SWCNT-anticancer drug delivery systems, it is evident that non-bonded interaction energies such as vdW and electrostatic interactions between the enclosed or engrafted drugs and SWCNT determine the adsorption function of the drug on the surface or inside the SWCNT. It is however, important to gain a deeper understanding about the performance of SWCNTs as possible host nanocarriers of anticancer drugs through the calculation of interaction energies between the drug and the SWCNTs to ascertain their suitability as controlled drug delivery systems and gain an insight about their drug encapsulation and release functions.

The MD simulation in this regard, is a promising tool that has been widely applied to investigate the performance of complex nano-sized systems bridging diverse areas of cell biology and materials science<sup>32–34</sup>. Besides the great ability that the MD offers to evaluate of the thermal and mechanical features of nanostructures<sup>35–40</sup>, the method provides the ability to obtain a deeper insight into the interatomic interactions of nano-sized complex systems containing bio-molecules<sup>41</sup>. The anticancer drug Isatin (1H-indole-2,3-dione), a natural alkaloid in a form of red–orange powder (aka. Tribulin), derived from indole with a formula  $C_8H_5NO_2$  that was first obtained by Otto Linné Erdman and Auguste Laurent<sup>42</sup> in 1840 as a by-product of indigo dye oxidation by nitric and chromic acids. Isatin is naturally found in a human body as an endogenous compound for metabolite of tryptophan or epinephrine<sup>43</sup>. Isatin is soluble in hot water, alcohol, acetic acid, and benzene but is sparingly soluble in ether. Remarkably, Isatin remains stable and soluble in concentrated hydrochloric acid and in concentrated sulfuric acid<sup>44</sup> these properties allow encapsulation of Isatin as a guest molecule inside the host CNTs and theoretically permit an *in-situ* functionalization of Isatin-containing CNTs.

In the present work, first, the MD simulation was employed to evaluate the encapsulation of Isatin inside the end-capped SWCNT (10,10) host through the calculation of vdW between the host SWCNT and the drug, followed by computing the potential of mean force (PMF) of the encapsulated Isatin inside the SWCNT in a aqueous environment. Secondly, the storage capacity of the capped SWCNT (10,10) with the length of  $30 \text{ \AA}$  to encapsulate the Isatin was evaluated.



**Figure 1.** The molecular structure of anticancer drug Isatin and its atoms as represented using the DREIDING force field.

### Simulation method

In the current work, the MD simulation of the encapsulation processes and storage of Isatin guest inside the capped SWCNT was performed by employing the Large-Scale Atomic/Molecular Simulator (LAMMPS) software (Ver. March 2018, <https://www.lammps.org/download.html>)<sup>45</sup>. Inspired by previous investigations, CHARMM27 force field was applied to obtain the interaction parameters of hydrogen and oxygen atoms in water molecules (TIP3P)<sup>46</sup>, a well-known force field widely applied in drug delivery simulation<sup>47</sup>. The Tersoff potential was used to consider the interaction between the carbon atoms in the SWCNT<sup>48</sup>. Evidently, several works have addressed successful application of the aforementioned force fields<sup>49,50</sup>. Atoms of modeled nanotube were fixed during the simulation. The molecular structure of Isatin is shown in Fig. 1. The interaction parameters between the atoms of Isatin were obtained from the DREIDING force field, which is a smart approach successfully used in prediction of the structure and dynamics of organic, biological, and main-group inorganic molecules<sup>51</sup>. The optimization of geometry of the drug was carried out by means of CVff force field to find charges of atoms, as proposed by developers of DREIDING methodology<sup>51</sup>. The Isatin force field data can be found in supplementary data.

The temperature and the pressure of the systems were locked to 300 K and 101.3 kPa, respectively, by applying NPT Ensemble<sup>52</sup>. The cutoff distance for the Lennard–Jones potential and Coulombic potential was considered to be 12 Å. The parameters of Lennard–Jones potential for cross vdW interactions between non-bonded atoms was estimated using Lorentz–Berthelot combination rule<sup>53</sup>. The visualization was obtained by means of visual MD (VMD) simulation (Ver. 1.9.3, <https://www.ks.uiuc.edu/Development/Download/download.cgi?PackageName=VMD>)<sup>54</sup>.

The simulation steps of this research are described as following:

1. At first step, the insertion process of the peptide Isatin into the capped CNT, and subsequently the stability of encapsulated drug inside the capped SWCNT host were studied. Considering the size of the Isatin guest molecule, the capped SWCNT in order to serve as a host drug nano-carrier was selected the chirality of an armchair (10,10) and displaying the length of 30 Å. At the beginning of the MD simulation, Isatin was situated at the initial distance of 2 Å from the nanotube. The axial direction of the SWCNT was set parallel to the z-axis of the simulation box. The complex comprised of the capped SWCNT- Isatin was immersed in the simulation box consisting of TIP3P 3-point water molecules as well as counter-ions to neutralize the simulated solution with periodic boundary condition. To assess the encapsulation process of the peptide, in the first stage, the minimization of the system was performed in the canonical NVT ensemble at 300 K, where moles (N), volume (V) and temperature (T) gradients were conserved. Next, the MD runs were performed in the NPT ensemble for 15 ns with the time step of 1 fs.

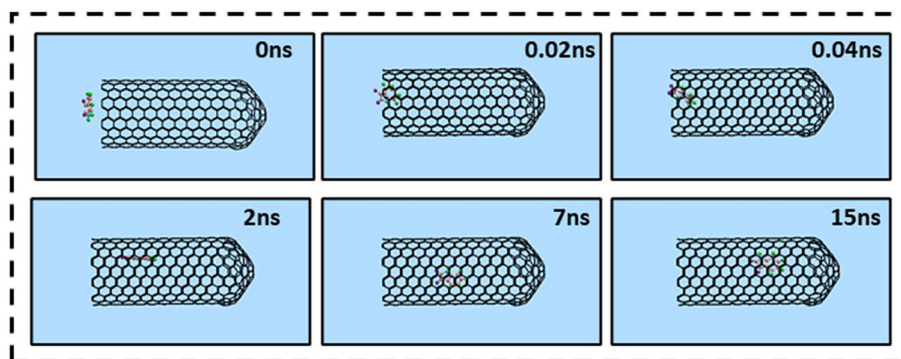
To evaluate the stability of the encapsulated Isatin inside the capped SWCNT, an external force was loaded on the encapsulated Isatin along the z-axis of the nanotube to pull it out from the capped CNT host in the direction opposite to the penetrating process. The spring constant  $k$  and pulling velocity were chosen equal to 15 kcal mol<sup>-1</sup> Å<sup>-2</sup> and 0.0005 Å ps<sup>-1</sup>, respectively<sup>47</sup>. The pulling process was simulated ten times to compute the potential of PMF using Jarzynski's equality as noted below<sup>46</sup>:

$$e^{-\beta\Delta G} = e^{-\beta W} \quad (1)$$

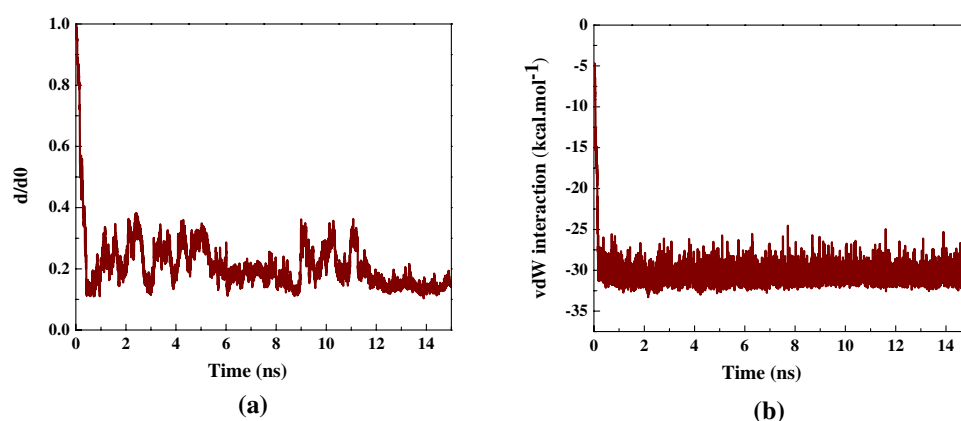
where  $\Delta G$  and  $W$  correspond to the free energy discrepancy between two states and the performed work on the system respectively. The  $\beta$  is equal to  $(K_B T)^{-1}$ , where  $K_B$  stands for the Boltzmann constant.

2. At the second step, the storage capacity of the capped SWCNT (10,10) host was investigated. For this, 15 molecules of Isatin were placed inside the capped SWCNT. The axial directions of nanotubes were set to be parallel to the z-axis of the simulation box. The minimization of the system was done in the canonical NVT ensemble at 300 K while the SWCNT were fixed. Then, the MD run was performed in the NPT ensemble for 15 ns with the time step of 1 fs.





**Figure 2.** Representative snapshots of insertion of a single molecule of Isatin into an armchair capped SWCNT (10,10) host at various times. For clarity, the water molecules have been omitted from the image.



**Figure 3.** (a) The  $d/d_0$  values (normalized to the CoM distance, where  $d_0$  is the initial CoM distance) between the drug Isatin and capped SWCNT (10,10) as a function of simulation time, (b) the vdW interaction between the drug Isatin and the capped SWCNT as a function of simulation time.

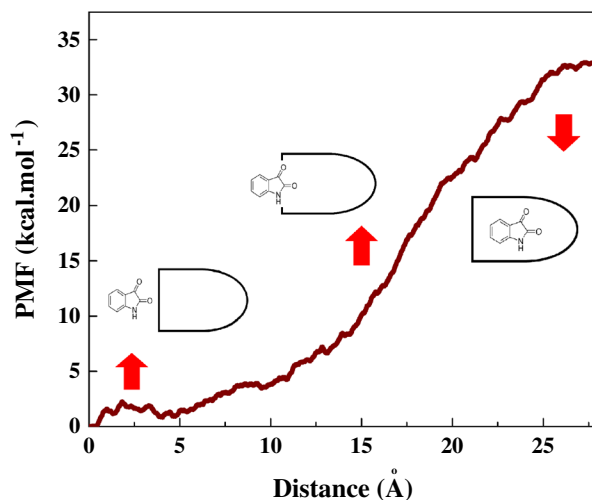
## Results and discussion

**Localization of Isatin within the capped SWCNT-drug complex.** The encapsulation process of a single Isatin guest molecule inside the capped SWCNT (10,10) host was investigated by means of the MD simulation. Figure 2 illustrates the snapshots of positions of the drug at various times in the simulation box which obtained using VMD software. As shown in Fig. 2, Isatin molecule was successfully encapsulated inside the capped SWCNT host. Moreover, the drug remained stable in the cavity of the nanotube owing to the vdW interaction interactions between the Isatin's conjugated aromatic rings and the interior wall (side wall and capped end) of the nanotube up to the end of the simulation time (15 ns). Another point is that DREIDNG, in the current work, appeared as a powerful potential for keeping the Isatin's structure stable without any deformation. Therefore, it can be concluded that choosing this potential for the modeled anticancer drug was a smart choice<sup>55</sup>.

Figure 3a shows the changes of the normalized center of mass (CoM) distance between Isatin guest and the capped SWCNT (10,10) host,  $d/d_0$ , during the simulation. The  $d/d_0$  value was found to reduce significantly during the first 0.5 ns of the simulation in good agreement with rapid adsorption of the drug into the capped SWCNT cavity owing to the marked vdW interaction forces between Isatin and the nanotube. The same decreasing trend in  $d/d_0$  value of at the beginning of the encapsulation process was observed by Kang et al.<sup>47</sup> reporting on SmtA protein experiencing self-adjustment through the conformational variations before entering the SWCNT. After the complete insertion of the drug occurred towards the end of the simulation, the value of the  $d/d_0$  were found to fluctuate continuously in the specified range due to the surrounding vdW interaction forces between the SWCNT host interior wall and the Isatin molecule.

According to the neutralized nature of the capped SWCNT (10,10), the value of the electrostatic interactions such as dipole–dipole, and H-bonding interactions between the nanotube and the Isatin molecule was found to be zero. Therefore, the vdW interaction energy was considered as an absorbance interaction between Isatin and the capped nanotube. Figure 3b demonstrates the changes in the value of the vdW interaction energy between the drug and the capped SWCNT (10,10) during the simulation.

Evidently, once the simulation was initialized, the value of the vdW energy decreased significantly from  $-5 \text{ kcal mol}^{-1}$  to approximately  $-30 \text{ kcal mol}^{-1}$  and fluctuated around the latter value until the end of the



**Figure 4.** The PMF for the system comprised of a single Isatin guest molecule inside the capped SWCNT host computed from ten (10) pulling through the MD simulation. The images represent the positions of Isatin molecule corresponding to the z-coordinate along the capped SWCNT at a close proximity-, entering/exiting- and in a fully-enclosed position inside the SWCNT host.

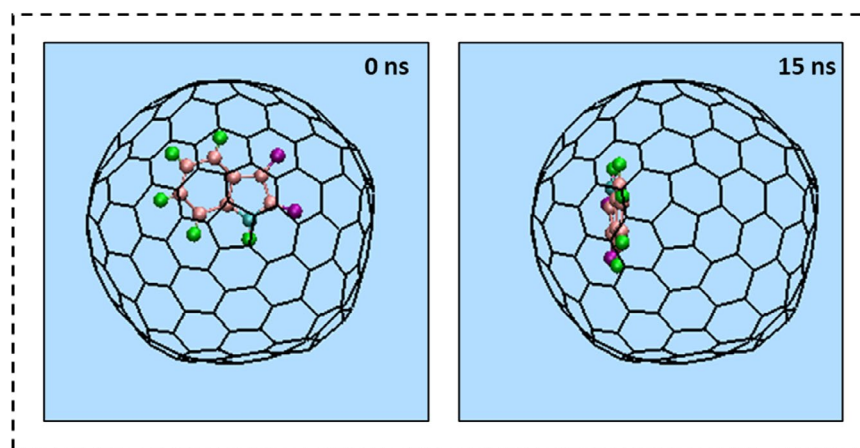
simulation. These changes revealed that the Isatin-SWCNT vdW interaction energy that is attributed to the relatively small size of the drug molecule, caused a rapid adsorption of Isatin molecule into the capped SWCNT host volume cavity. Moreover, the obtained low value for the vdW energy shows that a desirable interaction between the guest molecule and SWCNT host occurred in favor of both the adsorption and desorption functions of the drug in the Isatin-SWCNT nano-delivery system<sup>56</sup>.

After the sudden drop, the smooth fluctuation of the value of the vdW energy around the specified value showed that Isatin was affected by the vdW interaction between its conjugated aromatic rings of the and the interior wall (side wall and capped end) of the nanotube and remained stably encapsulated until the end of the simulation. A similar variation in vdW values was reported by Maleki et al.<sup>26</sup> determining the electrostatic and vdW interactions between the Doxorubicin molecule and SWCNT. According to their results, a zero value of electrostatic interactions provides the absorbance of Doxorubicin on the SWCNT rendered by the vdW interactions with a favorable negative value. An MD simulation performed by Hashemzadeh et al.<sup>57</sup> revealed that a reduction of the vdW energy between the guest Paclitaxel (AKA. Taxol) and SWCNT host to the negative value, which supports the drug natural encapsulation inside the SWCNT.

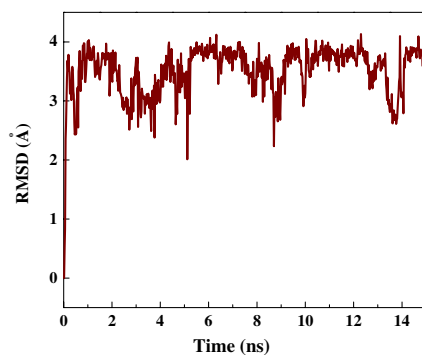
**Calculation of free energy from the MD simulation.** Following the termination of the MD simulation, once Isatin was completely inserted inside the capped SWCNT (10,10) at 15 ns, the PMF profile of the encapsulated drug was determined through pulling the drug molecule out by means of the MD simulation at the speed of  $0.0005 \text{ \AA ps}^{-1}$ , which has been selected according to the encapsulation process speed. Repeating the simulation ten times the average value of work ( $W$ ) at each pulling distance, was obtained as shown in the PMF functional profile (see Fig. 4) that illustrates the Isatin's positions along the z-axis of the capped SWCNT host relative to the distance travelled in Fig. 4. The free energy of this simulated system was found to increase relative to the pulling distance reaching its maximum value of  $34 \text{ kcal mol}^{-1}$  at the pulling distance of  $28 \text{ \AA}$ . The free energy of encapsulated Isatin molecule is  $-34 \text{ kcal mol}^{-1}$ , a signature of spontaneous encapsulation process. A similar value is reported by Veclani et al.<sup>58</sup> for the Ciprofloxacin (AKA. Cipro, Cifran) antibiotic encapsulated in SWCNT as  $-21 \text{ kcal mol}^{-1}$ , and  $-9.5 \text{ kcal mol}^{-1}$  for the drug adsorbed on the surface of a SWCNT.

**The variations in the conformation of Isatin.** The alterations in the conformation of a single Isatin molecule at 0 and 15 ns of the MD simulation are shown in Fig. 5a. The left image in Fig. 5a depicts the drug molecule immersed in aqueous solution outside the capped SWCNT host at 0 ns. The right image in Fig. 5a corresponds to the state in which the conformation of the drug was adjusted to the interior wall of the SWCNT at 15 ns. Notably, at the end of the simulation, Isatin's cyclic rings were adjusted to be situated parallel to the side wall of the SWCNT so that the highest the  $\pi$ - $\pi'$  stacking interaction was established.

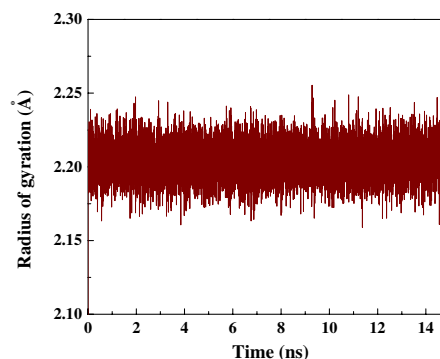
The root-mean-square-deviation (RMSD) corresponding to the variations in the conformation of Isatin inside the capped SWCNT (10,10) host as a function of the simulation time is shown in Fig. 5b. The RMSD varied significantly by 0.1 ns to the value of  $4 \text{ \AA}$  which revealed the variations in the both the conformation and the position of the drug molecule under the influence of the capped SWCNT-Isatin vdW interaction. Following the insertion of Isatin molecule inside the capped SWCNT host, the value of RMSD was found to fluctuate in the specified range confirming that the conformation of the drug could not vary much due to the physical confinement inside the cavity of the capped SWCNT host. The similar changes in the RMSD value of the platinum-based anticancer during the encapsulation inside the silicon-carbon nanotube was recently observed by Hasanzade et al.<sup>59</sup>. The alteration of the gyration radius of Isatin is shown in Fig. 5c. Isatin gyration radius



(a)

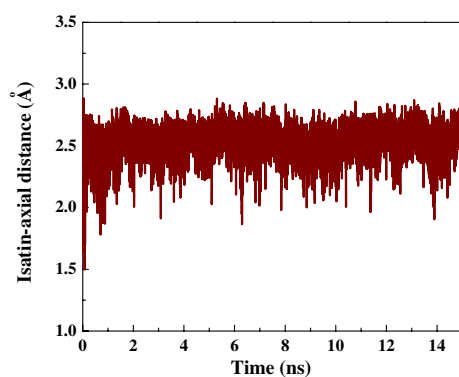


(b)

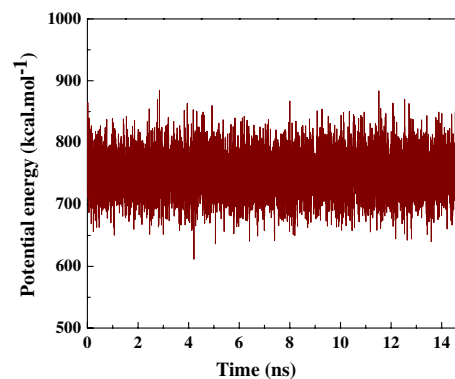


(c)

**Figure 5.** (a) Axial views of a single Isatin guest molecule at 0 ns and 15 ns in the MD simulation. For clarity, the water molecules have been omitted from the image. (b) The RMSD values of Isatin as shown as a function of simulation time. (c) Isatin radius gyration is shown as a function of simulation time in the capped SWCNT-Isatin complex.



(a)

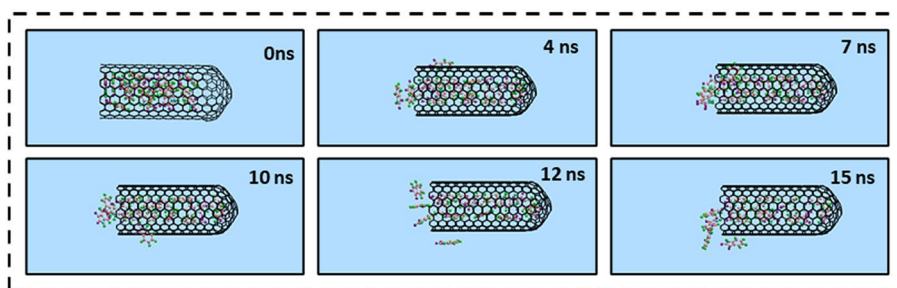


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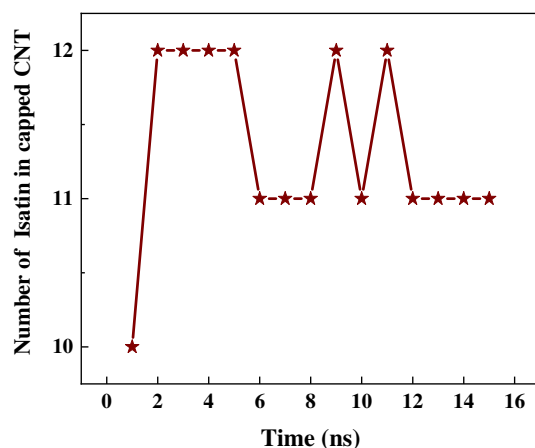
**Figure 6.** (a) The distance between Isatin's CoM and the central axis of capped SWCNT as a function of simulation time. (b) The potential energy variation of Isatin inside the capped SWCNT-Isatin complex as a function of simulation time.

was found to be essentially unaltered but varied continuously in a narrow range of 2.175–2.225 Å owing to a relatively small size and rigid molecular structure of the drug.

Figure 6a shows the alterations in distance between Isatin's CoM and the central axis of the capped SWCNT (10,10) host as a function of simulation time. At the beginning of the simulation, the drug in the simulation box was positioned in a way that the CoM of the guest drug was aligned on the central axis of the SWCNT host. Once



**Figure 7.** Representative snapshots of storage capacity of Isatin guest molecules inside a single capped SWCNT (10,10) host at various simulation times. For clarity, the water molecules have been omitted from the image.



**Figure 8.** A potential number of Isatin molecules enclosed inside a single capped SWCNT (10,10) host carrier as a function of simulation time.

the simulated commenced, Isatin molecule moved suddenly towards one side of the capped SWCNT owing to vdW interaction between the SWCNT wall and the drug resulting in an immediate increase of the drug-axis distance to the value of 2.7 Å. After the complete insertion of Isatin molecule inside the SWCNT host, the distance between the CoM of the drug and central axis of the SWCNT host fluctuated slightly confirming the stability of Isatin molecule inside the SWCNT due to the desirable SWCNT-to-Isatin vdW interaction.

The changes in the potential energy of Isatin during the self-encapsulation process is shown Fig. 6b. The potential energy curve of Isatin is essentially a linear function that shows minor fluctuations around the value of 600 kcal mol<sup>-1</sup> during the insertion process confirming that the simulated system has reached its equilibrium state. A similar trend in potential energy changes of the Flutamide (AKA. Eulexin) anticancer drug in the SWCNT- Flutamide complex was recently reported by Kamel et al.<sup>60</sup>

**The storage of Isatin inside the capped SWCNT.** Due to the possible cytotoxicity associated with CNTs at high concentrations, the encapsulation of more than one guest Isatin molecule inside a single host SWCNT is likely lower the amount the required CNTs in nanomedicine therapy. Figure 7 illustrates the snapshots of the MD simulation of the storage of the several Isatin guest molecules inside a single capped SWCNT host at various times.

Figure 8 shows the number of Isatin guest molecules encapsulated in a single capped SWCNT (10,10) host as a function of simulation time. At 0 ns of the simulation, fifteen (15) molecules of Isatin were positioned inside the cavity of a single capped SWCNT host. As the simulation commenced, a few Isatin molecules were expelled from the SWCNT cavity and shown attached to the surrounding exterior wall of the SWCNT due to the vdW interactions. Our simulation have shown that an average eleven (11) Isatin guest molecules could be stably encapsulated inside a single capped SWCNT host carrier.

## Conclusion

Our work described the encapsulation process of the anticancer drug Isatin inside the capped SWCNT (10,10) and, evaluated a number of Isatin guest molecules that can be readily encapsulated inside the SWCNT cavity employing the MD simulation. Our results revealed that a single Isatin guest molecule can be readily and rapidly adsorbed inside the capped SWCNT host and remained stably encapsulated until the end of the MD simulation. The value of the capped SWCNT-Isatin vdW interaction energy was found to reach the value of - 30 kcal mol<sup>-1</sup>

at the end of the simulation (15 ns) that showed a high probability of complete adsorption and desorption of Isatin guest inside the SWCNT host SWCNT-Isatin drug nano-delivery system. In first 0.1 ns of the simulation, the RMSD was found to vary significantly reaching the value of 4 Å. The latter revealing the variations in both the conformation and the position of Isatin guest under the influence of the capped SWCNT-Isatin vdW interaction. The obtained value of free energy of encapsulation process, namely,  $-34 \text{ kcal mol}^{-1}$  has confirmed that the insertion procedure of Isatin guest into the SWCNT host occurred spontaneously and naturally. The MD simulation performed to calculate the probable storage capacity of the capped SWCNT (10,10) with length of 30 Å have shown that approximately eleven (11) Isatin guest molecules could be stably encapsulated inside a single capped SWCNT host cavity, confirming a potential for SWCNT-Isatin drug delivery systems in future nano-medicine applications.

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### Author contributions

M.Z.D. performed the simulation and wrote the first draft. A.H.M. designed and conceptualized the study. F.Y. and F.S. developed research methodology and checked chemical interactions/possibility of encapsulation, respectively. B.B. and G.N. checked data collection. F.Y. and A.H.M. developed the theoretical formalism and analyzed the data. A.E. and O.A. drafted the manuscript. A.H.M. and S.H. supervised the study. M.R.S. coordinated the study. M.R. analyzed the methodology and critically revised the manuscript. All authors read, commented on the manuscript, gave the final approval for publication and agreed to be held accountable for the work performed herein.

### Competing interests

The authors declare no competing interests.

### Additional information

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