Synergistic Antibacterial Potential and Cell Surface Topology Study of Carbon Nanodots and Tetracycline Against E. coli

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Increasing drugs and antibiotic resistance against pathogenic bacteria create the necessity to explore novel biocompatible antibacterial materials. This study investigated the antibacterial effect of carbon dot (C-dot) against E. coli and suggested an effective synergistic dose of tetracycline with C-dot, using mathematical modeling of antibacterial data. Colony count and growth curve studies clearly show an enhanced antibacterial activity against E. coli synergistically treated with C-dot and tetracycline, even at a concentration ten times lower than the minimum inhibitory concentration (MIC). The Richards model-fit of growth curve clearly showed an increase in doubling time, reduction in growth rate, and early stationary phase in the synergistic treatment with 42% reduction in the growth rate (μm) compared to the control. Morphological studies of E. coli synergistically treated with C-dot + tetracycline showed cell damage and deposition of C-dots on the bacterial cell membrane in scanning electron microscopy imaging. We further validated the topological changes, cell surface roughness, and significant changes in the height profile (ΔZ) with the control and treated E. coli cells viewed under an atomic force microscope. We confirmed that the effective antibacterial doses of C-dot and tetracycline were much lower than the MIC in a synergistic treatment.

Keywords: carbon dot, antibacterial, carbon nanomaterial, synergistic, atomic force microscopy, antibiotic resistance

INTRODUCTION

Several studies suggest that nanoparticles (NPs) can be an excellent replacement to antibiotics and effective drug carriers and are suitable for multifunctional designs with their broad-spectrum antibacterial properties, considering the availability of various functional groups (Mühling et al., 2009; Pelgrift and Friedman, 2013; Beyth et al., 2015). Nanoparticles not only combat antibiotic resistance but also act as a carrier of drugs/antibiotics such as liposomal NPs (Daeihamed et al., 2016) and solid lipid (SL) NPs (Thuokral et al., 2014; Naseri et al., 2015) used in drug delivery and a wide...
range of therapeutics (Pissuwan et al., 2011; Gholipourmalekabadi et al., 2017). Carbon nanodots (C-dots) or carbon quantum dots are one of the most recently discovered and effective antimicrobial agents (Thakur et al., 2014; Lim et al., 2015; Maas, 2016; Meziani et al., 2016; Yang et al., 2016; Dong et al., 2017; Xin et al., 2019; Muktha et al., 2020). They are carbon-based nanoparticles with a size less than 10 nm and have surface passivation properties (Sun et al., 2006; LeCroy et al., 2016). C-dots possess photo excited state properties and redox processes similar to conventional nonscale semiconductors (Rocha et al., 2002; Hu et al., 2015; Jelinek, 2017; Righetto et al., 2017), which initiate the efficient photoinduced charge separation. This process allows the formation of radical anions and cations (Cao et al., 2013; Fernando et al., 2015; LeCroy et al., 2016). These photo-generated electrons and holes in carbon dots are the basis of catalytic processes (Cao et al., 2013; Fernando et al., 2015), providing a strong photodynamic effect to C-dots (Luo et al., 2021). Due to their photoinduced redox potential, C-dots are excellent candidates functioning as antibacterial agents and are effective against bacterial cells, particularly under normal visible/natural light illumination without giving an excess dose of the photon (Lim et al., 2015; Meziani et al., 2016). An excess amount of photons also triggers C-dot-dependent photodamage of tetracycline and other similar antibiotics (López-Peñalver et al., 2010; Nagamine et al., 2020; Semeraro et al., 2020; Xu et al., 2020; Liu et al., 2021).

When traditional antibacterial reagents are compared with C-dot, the latter proves to be advantageous as they are intrinsically non-toxic both in vitro and in vivo (Najjar et al., 2007; Li et al., 2014a) and environment-friendly, whereas antibacterial chemicals mostly exhibit potential toxicity to the environment and ecological elements. As discussed earlier, the uncheckered use of antibiotics and other antibacterial substances has led to the emergence of multidrug-resistant bacteria (MDR). Thus, the C-dot being non-toxic and environment-friendly can be effectively used against these MDR bacteria and helps control infections in public health sectors. Studies have suggested a direct interaction between carbon-based nanoparticles with various bacterial and fungal cells (Zou et al., 2016). Studies on the antimicrobial activities of carbon dots are increasing with time, and several studies suggested mechanisms that include oxidative stress (Akhanv et al., 2011; Li et al., 2014b). Currently, few reports are available on the potential of carbon dots having an antimicrobial effect on E. coli and other microorganisms (Thakur et al., 2014; Meziani et al., 2016; Dong et al., 2018; Dong et al., Elcodous et al., 2020; 2020; Muktha et al., 2020).

C-dots obtained from gum arabic showed potential antimicrobial activity against Gram-positive and Gram-negative bacteria (Thukral et al., 2014). Nanocarbons synthesized by green synthesis from turmeric smudge, pomegranate peel, and watermelon peel have also been reported with potent antibacterial activity against E. coli, Bacillus subtilis, and Pseudomonas aeruginosa (Chun et al., 2016; Muktha et al., 2020). In recent years, even though many studies have emerged on the use of C-dots for various applications, more systematic studies on the antimicrobial property are required to understand the exact mechanisms of their actions. Detailed analysis of the antibacterial effect and the mathematical modeling of the obtained data are necessary to understand the minute changes to identify the effective doses of such broad-spectrum antibacterial agents (Zwietering et al., 1990; Qu et al., 2004; Liu et al., 2005; Mouton and Vinks, 2005; Rao et al., 2018). Several nanomaterials combined with other antibacterial reagents, chemicals, and antibiotics, such as H2O2, Na2CO3, acetic acid, methylene blue, toluidine blue, tetracycline, and neomycin, exhibited enhanced synergistic effects against S. aureus, Bacillus subtilis, Acinetobacter, P. aeruginosa, and E. coli even at concentrations below the MIC (Thakur et al., 2014; Meziani et al., 2016; Dong et al., 2017; Ehsan and Sajjad, 2017; Dong et al., 2018; Abd Elkodous et al., 2020). In the current study, we confirmed the synergistic biocidal effect of C-dots along with tetracycline well below the MIC, showing an enhanced antibacterial effect, reduction in doubling time, and morphological and topological changes on E. coli cells. Atomic force microscopy (AFM) studies can provide details about the outer membrane of cells. The earliest surface studies of E. coli cells revealed that the force–distance curve was able to detect the surface structure using functionalized probes in AFM imaging and show a detection method of bacteria at the single-cell level (Tanaka et al., 1999; Amro et al., 2000; Miller et al., 2006). After treating with antimicrobial agents such as chitosan, fosfomycin, and other molecules, the images obtained in AFM measurements revealed antibacterial effects, cell wall collapse, and significant morphological changes (Neethirajan and DiCicco, 2014; Eaton et al., 2008; Alves et al., 2010; Ipte and Satpati, 2020; Zhang et al., 2021).

**MATERIALS AND METHODS**

**Materials and Instrumentation**

Carbon nanodots obtained from Sigma-Aldrich Inc. were used for this study (Product no. 900414). Antibiotic tetracycline, Luria–Bertani broth, Luria–Bertani agar, Muller–Hinton agar, and Muller–Hinton broth were all research grade (Hi Media laboratories). The instruments used for this study are a spectrophotometer (Chemito UV2300, Shimatduz Corp.), sonicator (Sonics and materials), shaker, incubator (Classic Scientific), laminar airflow, laboratory centrifuge, scanning electron microscope (ZEISS EVO18), and NTMDT atomic force microscope (NTEGRA). All the glassware used for the experiments were of Borosil, Riviera, Vensil, or Schott Duran.

**Characterization**

Carbon dots used for this study were research grade and purchased from Sigma-Aldrich (Product no. 900414). We further characterized the size, absorption spectra, and emission spectra of C-dots using a scanning electron microscope, spectrophotometer, and fluorescence spectrophotometer, respectively (Supplementary Figure S1).

**Biocidal Effect of Tetracycline**

The susceptibility of tetracycline on E. coli culture was tested on agar plates by a well diffusion assay. In brief, 5.0 ml of 50 μg/ml...
Working stock of tetracycline was prepared and filter-sterilized. From the working stock of tetracycline, different concentrations (0.25 μg/ml, 0.5 μg/ml, 1.0 μg/ml, 1.5 μg/ml, 2.0 μg/ml, 2.5 μg/ml, 3.0 μg/ml, 3.5 μg/ml, 4.0 μg/ml, 4.5 μg/ml, and 5.0 μg/ml) were used for the antibacterial susceptibility assay. An overnight grown culture of *E. coli* (100 μL) spread-plate on Mueller–Hinton agar plates at room temperature (RT), wells were cut in the agar plate with a 5-mm well borer, and 50 μL of each of the tetracycline concentrations, as mentioned before, were added to the individual wells. The aforementioned experiments were performed under sterile conditions in the laminar flow. The plates were incubated at 37°C for 16 h, and the diameter of the zone of inhibition was recorded.

A tetracycline susceptibility test was also carried out in liquid culture to validate the inhibitory effect of the zone of inhibition assay. From the working stock of 50 μg/ml tetracycline, concentrations of tetracycline (0.25 μg/ml, 0.5 μg/ml, 1.0 μg/ml, 1.5 μg/ml, 2.0 μg/ml, 2.5 μg/ml, 3.0 μg/ml, 3.5 μg/ml, 4.0 μg/ml, 4.5 μg/ml, 5.0 μg/ml, 5.5 μg/ml, and 6.0 μg/ml) were used by making up the final volume to 5 ml in Muller–Hinton broth. Each test tube was inoculated with 10 μL of the overnight grown culture. The tubes were kept at 37°C on a shaker for 24 h. The O.D. was recorded at 595 nm.

### Biocidal Effect of C-Dots

From the sonicated stock solution of 2.0 mg/ml of C-dot, 1.0 μg/ml, 2.0 μg/ml, 4.0 μg/ml, 8.0 μg/ml, 16 μg/ml, 32 μg/ml, and 64 μg/ml concentrations were used to assess the antibacterial effect. The 1.0 ml of overnight grown *E. coli* culture was centrifuged, pelleted, washed twice with PBS, and mixed in 1.0 ml PBS. The bacterial culture was treated with the same volume of the C-dot mixture solution to maintain the C-dot concentrations and incubated for 1 h with intermittent mixing at room temperature under visible light conditions. Various studies reported nanomaterial-mediated photodamage of tetracycline. Therefore, we used only visible light in C-dot treatment experiments to avoid nanomaterial-dependent photoinduced tetracycline damage. After incubation, 10 μL of the treated samples were inoculated in individual test tubes containing 5.0 ml of sterile LB. All tubes were incubated in a 37°C incubator shaker at 115 RPM for 8 h. The O.D. was recorded at 595 nm for all the samples.

Four C-dot samples of 8.0 μg/ml, 16 μg/ml, 32 μg/ml, and 64 μg/ml from the treated samples were used to study effect on the bacterial colony count. We serially diluted the treated culture and spread plated the 100 μL of each sample onto the LB agar plates. The plates were incubated at 37°C overnight, and the CFU was calculated from the obtained colony count.

### Synergistic Biocidal Effect of C-Dot and Tetracycline

For the synergistic treatment, 0.1 μg/ml, 0.25 μg/ml, 0.5 μg/ml, 0.75 μg/ml, 1.0 μg/ml, and 1.5 μg/ml concentrations of tetracycline and 2.0 μg/ml, 4.0 μg/ml, 8.0 μg/ml, and 16 μg/ml of C-dots were used. Synergistic doses were maintained by mixing a 1:1 ratio of 500 μL tetracycline +500 μL C-dot. The 1.0 ml of overnight grown *E. coli* culture was centrifuged, pelleted, washed twice with PBS, and mixed in 1.0 ml PBS. The bacterial culture was treated with an equal volume of the tetracycline + C-dot mixture solutions prepared before. And 50 μL of *E. coli* cells were added to each tube to maintain a final volume of 1.0 ml. The culture tubes were incubated for 1 h with intermittent mixing at room temperature. After incubation, 10 μL of the treated samples were inoculated in individual test tubes containing 5.0 ml of sterile LB broth along with the untreated control. All tubes were incubated in a 37°C incubator shaker at 115 RPM for 8 h. The O.D. was recorded at 595 nm for all the samples.

For synergistic treatment, we used 8.0 μg/ml and 16 μg/ml C-dot concentrations with 1.5 μg/ml tetracycline. Two samples were mixed in a 1:1 ratio (500 μL tetracycline +500 μL C-dot) to maintain the aforementioned concentrations and further mixed in 1.0 ml volume of the overnight grown *E. coli* culture. Synergistic treatment was carried out for 1 h at room temperature with intermittent mixing. The treated samples were centrifuged, pelleted, washed twice with PBS, and mixed in 1.0 ml PBS. After treatment, the culture was serially diluted and 100 μL of the treated sample spread onto LB agar plates. The plates were incubated at 37°C overnight; the number of colonies was recorded to calculate CFU.

### Growth Curve Study of Synergistic C-Dot and Tetracycline Treatment

The growth curve of *E. coli* was studied by culturing the bacteria in 25 ml of LB broth overnight at 37°C on a shaker at 115 RPM. Tetracycline and C-dot mixture were prepared, as discussed earlier. A control of LB media together with 1.5 μg/ml tetracycline, 16 μg/mL C-dot, and 1.5 μg/ml tetracycline +16 μg/mL C-dot was used for the growth curve measurement. *E. coli* cells were treated with all the aforementioned dilutions for 1.5 h at 37°C with intermittent mixing. After incubation, 200 μL of each sample was inoculated in conical flasks containing 25 ml of sterile LB broth and kept in an incubator shaker at 37°C at 115 RPM. The O.D. was recorded at 595 nm after every 1-h interval.

### Analysis of Growth Kinetics

Growth rate kinetic analysis of bacteria is oriented toward understanding the cellular process that links the two or more cellular functions, for example, the relationships between growth and cell doubling time, fitted using empirical functions by mathematical modeling based on multiple steps involved in the progression of cell growth. The experimental data sets require the analysis of depth parameter statistics to estimate the precision of the results by applying descriptive methods. Virtually, all of the models developed for kinetic studies are non-linear (Qu et al., 2004). These functional models describe the number of organisms or logarithm of the number of organisms as a function of time. Growth curves obtained in spectroscopic experiments can be modeled using various growth functions such as logistic [Eq. (1)], Gompertz [Eq. (2)], Richards, and Stannard [Eq. (3)]; those are mathematically equivalent (Zwietering et al., 1990;
We process the experimental growth curve data using the following mathematical model described in Eq. 1, Eq. 2, and Eq. 3.

\[ y(t) = \frac{A}{1 + \exp\left(\frac{\mu_{\text{max}}(t - t)}{A}\right)} \]  

Gompertz function:

\[ y(t) = A\exp\left( - \exp\left(\frac{\mu_{\text{max}}\exp\left(\frac{1}{A}\right)(\lambda - t) + 1}{\lambda} \right) \right), \]  

Richard function:

\[ y(t) = A\left[1 + \exp\left(1 + v\right)\exp\left(\frac{\mu_{\text{max}}}{A}\right)\left(1 + \frac{1}{v}\right)(\lambda - t)\right]^{\frac{1}{v}}. \]

The generation time \( t_{\text{gen}} \) or the population doubling time for logistic growth is related to the growth rate as \( t_{\text{gen}} = \ln(2)/\mu_{\text{max}} \).

The experimental optical density (O.D.) data are a function of time for control without any treatment. Bacterial growth often shows a phase in which the specific growth rate starts at a value of zero and then reaches a maximal value (μ) in a certain period, resulting in a lag time (λ). The growth progress also contains a final phase in which the rate decreases and finally reaches zero, so that an asymptote (A) reached or maximum O.D. We fitted the growth curve of \( E. coli \), treated with C-dot, tetracycline, and C-dot + tetracycline, employing the logistic, Gompertz, Richards, and Stannard function using the Levenberg–Marquardt algorithm as implemented in Gnuplot (Williams and Kelley, 2010). These functions are useful to obtain easily interpretable metrics such as population kinetics, the growth rate, the initial population size, and doubling time to summarize microbial growth curve data. In the previous modeled-fit functions, the growth curve is considered a function of O.D. plotted against time. The changes in the growth rate shows sigmoidal behavior (Figure 3), with a lag phase immediately after \( t = 0 \), followed by an exponential phase and then a stationary phase.

**Nanoparticles Interaction and Cellular Topology Study**

Direct visualization of the cell under high magnification is the best way to assess nanomaterial-dependent toxicity effects in bacterial cells. We used scanning electron microscopy (SEM) and atomic force microscopy (AFM) imaging to study the interaction between the bacteria and nanoparticles, morphological changes, and the topological changes on the cell surface in the independent and synergistically treated samples with C-dots and tetracycline.

**Morphological Study With Scanning Electron Microscopy**

The overnight grown \( E. coli \) cells (2.0 ml) for control, treated culture with C-dot and tetracycline, and their combinations were prepared. \( E. coli \) cells were grown in LB media for 6 h and centrifuged at 5,000 rpm for 5 min. The supernatant was discarded; the pellet was washed twice with PBS and centrifuged for 5 min at 5,000 rpm. The supernatant was again discarded, and the pellet was resuspended in 50 µL of PBS buffer. Under sterile conditions, 5.0 µL of the \( E. coli \) sample was spread on a glass slide and allowed to air dry at 20°C. The cells on slides were fixed with 2.5% paraformaldehyde for 20 min. The fixed cells were gently washed twice with PBS and then finally with double distilled water. The fixed cells were post-fixed with 1% osmium tetroxide for 1 h. The samples were dehydrated by flooding the slide using a series of ethanol concentrations of 35, 50, 75%, and 95% ethanol for 10 min each and finally with absolute alcohol for 10 min. Then the samples were air-dried and dehumidified for 20 min under vacuum and sputter-coated with gold for SEM analyses using a ZEI SS EVO 18 SEM imaging system.

**Surface Topology Study by Atomic Force Microscopy**

For AFM imaging, 2.0 ml of the overnight grown \( E. coli \) culture was treated with C-dot and C-dot + tetracycline along with untreated control. Treated cells were centrifuged at 5,000 rpm for 5 min. The supernatant was discarded; the pellet was washed twice with PBS and centrifuged at 5,000 rpm. The collected cell pellets were resuspended in 50 µL PBS, and 20 µL cells spread on the 1.0-cm × 1.0-cm mica sheet. Cells were air-dried and fixed on a mica sheet using 2.5% glutaraldehyde for 30 min. The fixed cell samples were air-dried at 20°C room temperature for AFM analysis. The experiment was carried out with an NTegra Prima AFM mounted on an Olympus inverted optical microscope and operated with the NTMDT controller, placed on a k-technology vibration isolator. This technique detects the van der Waals forces existing between a probe tip and molecular material (or the surface sample or biological material), scanned along the cantilever tip with a piezo scanner (maximum XYZ scan range of 100 µm × 100 µm × 20 µm). The cantilever tip is attached at one end to a cantilever which is built in at the other end. This technique can generate information detected on surface topography and mechanical properties at the nanometer scale. All data presented in this article were generated with NSG01 series cantilevers (single crystal silicon, N-type, resistivity 0.01–0.025 Ω-cm, antimony-doped), whose spring constant was about 40 N/m, with a maximum resonant frequency of 150 kHz and at a scan rate of 0.2–1 Hz. The sample was mounted on the freshly cleaved mica substrate surface 5 mm × 5 mm using the drop-cast method. We performed experiments at 18°C in a dehumidified and acoustic room. We utilized two non-contact atomic force microscopy (NC-AFM) modes; amplitude modulation, which gave height profile, and frequency modulation that provide phase images of the sample. We processed the data obtained in AFM measurements using open source software Gwyddion (Něcás and Klapeček, 2012; Klapeček et al., 2017). The height profile data are processed by subtracting plain surface data to eliminate tilt in the images and perform facet leveling.
RESULTS

Biocidal Effect of Tetracycline and C-Dot

Tetracycline is a very well-known broad-spectrum antibiotic for the management of bacterial infections. However, the overuse of antibiotics is creating antibiotic resistance against pathogenic strains. Reducing the effective dose of antibiotics is a significant challenge. We have explored the antibacterial efficacy of C-dot. The inhibitory effect was confirmed on E. coli cells treated with varying tetracycline concentrations. The results showed inhibition of the E. coli growth with tetracycline concentrations of 1.5 μg/ml to 6.0 μg/ml (Supplementary Figure S2 and Supplementary Table S1). Complete inhibition was observed below 1.5 μg/ml concentrations without an inhibition zone. When tested in liquid culture, a decrease in growth was observed up to 1.5 μg/ml concentration. The minimum inhibitory concentration (MIC) for tetracycline against E. coli was 1.5 μg/ml. The antibacterial effect on E. coli was further studied with a liquid culture using C-dot. The increase in C-dot concentration from 1.0 μg/ml to 32 μg/ml showed growth inhibition and 50% reduction in growth observed at 16 μg/mL C-dot concentration (Figure 1A). The C-dot–treated cells showed a decrease in the number of colonies compared to the control plate, where more than 50% of the colonies decreased at 8.0 μg/mL. C-dot concentration (Figure 1B, Supplementary Figure S3 and Supplementary Table S2). CFU decreased as the C-dot concentration increased from 8 μg/ml to 64 μg/mL. A ten time CFU decrease was observed at 64 μg/mL.
C-dot concentration. A correlation between the O.D. in broth and the CFU of *E. coli* was observed with a correlation coefficient of 0.8955, which indicates good correlation between two independent treatments between O.D. in broth and CFU (Figures 1C,D).

**Synergistic Biocidal Effect of C-Dot and Tetracycline**

Our data for synergistic treatment showed a similar trend in the growth of *E. coli* cells treated with tetracycline and C-dot for the aforementioned concentrations. Significant growth inhibition was observed below and near the MIC of tetracycline together with the MIC of C-dot. The results showed considerable inhibition of the growth of *E. coli* cells with six time lower tetracycline (0.25 μg/ml) along with 0.75 μg/mL C-dot concentration than the MIC (Figure 2A, Supplementary Table S3). Growth of *E. coli* decreased twofold in the 1.5 μg/ml tetracycline (MIC) + 8.0 μg/mL C-dot treated samples compared to independent treatments. The number of colonies reduced eightfold when the concentration of C-dot increased to 16 μg/ml compared to the MIC concentration of tetracycline (Figure 2B, Supplementary Figure S4 and Supplementary Table S3). *E. coli* cell counts were reduced by 30% with 0.75 μg/ml tetracycline along with 1.0 μg/mL C-dot; 50% with 1.5 μg/ml tetracycline with 2.0 μg/mL C-dot; and 75% with 1.5 μg/ml tetracycline with 16 μg/mL C-dot concentrations. In all the aforementioned cases, the concentration of tetracycline was lower or near to the MIC (Figure 2A). These synergistic treatment results indicate a dramatic improvement of the antibacterial efficiency compared to the independent treatment of either tetracycline or C-dot.

**Growth Curve Study for Synergistic Treatment**

Time-dependent growth retardation and early stationary phase were observed from the growth curve data of synergistic treatment compared to independent treatments either with 1.5 μg/ml tetracycline or 16 μg/mL C-dot (Figure 3 and Supplementary Figure S5A). At 7 h, comparing the control, a 20% decrease in growth of the cells treated with 1.5 μg/ml tetracycline and a 40% reduction in the cells treated with 16 μg/mL C-dot were observed. However, nearly 80% growth reduction in the logarithmic phase was observed on treating the cells with a combination of 16 μg/mL C-dot and 1.5 μg/ml tetracycline. The stationary phase was not attained in both independent treatments at 7 h (Figure 3). These data support the enhanced antibacterial effect in the synergistic treatment. The control culture exhibited a sigmoidal growth curve, and the treated culture sample showed a reduction in growth with an increase in the concentration of the C-dot and tetracycline. The model’s parameters predict that the synergistic treatment with C-dot + tetracycline treatment has a higher inhibitory effect on *E. coli* than the independent treatments. The experimental data support the model’s prediction (Figure 3 and Supplementary Figure S4). The model-fit data revealed that the growth rate
parameter $\mu_m$ decreased by 32, 44, and, 42% for logistic function, Gompertz, and Richards function compared to the control (Supplementary Figure S5 and Supplementary Table S1). The Richards function gives a better fit to the experimental growth data (Figure 3).

Nanoparticle Interaction and Bacterial Morphology Study

Scanning Electron Microscopy Analysis

Morphological changes were observed in C-dot, and C-dot-tetracycline-treated E. coli cells. The bacterial cells showed the deposition or binding on the cell surface, deformity in the cellular membranes, and surface roughness (Figure 4). In the samples treated with the combination of C-dot with tetracycline, the E. coli cells showed extensive deposition and destruction of cell morphology compared to the cells treated with only C-dot. The red arrow in the image indicates the morphological changes, cell damage, and deposition of C-dot. Compared to the control (Figure 4A), cells treated with C-dot and C-dot–tetracycline clearly show damage and binding on the outer membrane of cells (Figure 4B). However, the synergistic treatment indicates the enhanced biocidal effect, where the cells show severe damage on outer membranes and distorted morphology (Figure 4C).

Atomic Force Microscopy Analysis

The AFM studies of the three-dimensional imaging of the E. coli bacterial samples treated with C-dot and tetracycline revealed morphological/topological changes. The plotted graph for cell height and surface roughness suggests the high degree of roughness on C-dot–treated cells compared to control cells (Figure 5 and Supplementary Figure S6). We performed AFM measurements under ambient conditions (air) at a non-/contact intermittent mode to get the advantage of the harmonic oscillator response behavior of the AFM cantilever. This operating mode measures minimal force (quantum chemical van der Waals force) that prevents the indentation of bacteria by the AFM tip. This will protect the outer membrane of the E. coli cells, which is reflected in the observed phase-contrast and height images. Imaging of E. coli under physiological conditions would make more sense, but this will lower the accuracy of the nanoscale resolution on the outer membrane. Phase images (or

FIGURE 4 | E. coli cells observed under scanning electron microscopy (40.00 K X). (A) Control E. coli cells. (B) E. coli cells treated with C-dot. (C) E. coli cells treated with C-dot together with tetracycline. Scale bars in the panels left to right are 2 µm, 1 µm, and 200 nm, respectively.
Frequency modulation mode have been used to achieve the detail information of outer membrane which is not possible in height imaging (or amplitude mode). The Phase imaging mode is the phase difference between the exciting signal and the response of the cantilever. This mode is susceptible to the increase in interaction between the AFM tip and the surface. The dissipative processes happening in the AFM tip allow the study of the electromechanical properties of bacterial membranes at nanoscale. The results obtained in the present experiment are shown in Figure 5. The height information has been extracted from a linear map across a sample cell in the control cell. These data predict that the control cell has a length of ~1.1 μm with a height of about 38 nm and a width of 80 nm (Figures 5A–D). The (linear path) cross section across the control cell shows the smooth topological distribution, with vertical height distribution (ΔZ) of 8.0 nm. However, the C-dot– and C-dot + tetracycline–treated samples show ΔZ distribution of ~31 and ~25 nm, respectively. These data indicate the higher morphological change with C-dot + tetracycline synergistic treatment (Figures 5E–L). The average line profile over multiple cells is shown in Supplementary Figure S6B. A minor deviation is observed for the peak of line profile cross section ~ 8 nm for control cells, while ~22 and ~30 nm for cells treated with C-dot and C-dot + tetracycline.

**DISCUSSION**

Due to the emerging antibiotic resistance, there is a need to reduce the exposure and the heavy dose of antibiotics or replace other biocompatible antibacterial materials. The problem of antibiotic resistance has become a global issue, especially in the public health sector. To combat this, nanoparticles and their combination with other broad-spectrum bactericidal agents can be a potential solution. In the present study, we have used a tetracycline and C-dot combination to minimize biocidal treatment. Tetracycline inhibits bacterial protein synthesis and prevents the association of aminoacyl-tRNA with the bacterial ribosome (Chopra et al., 1992). C-dot is a biocompatible nanomaterial with a potent antibacterial effect. This study revealed a synergistic approach using antibiotics and C-dots.
below their MIC, resulting in enhanced bactericidal properties against E. coli. The proposed mechanism for the toxicity of C-dot is due to the effect of photoinduced electrons and holes responsible for their antimicrobial properties under visible light exposure (Lim et al., 2015; Meziani et al., 2016). In case of synergistic antibacterial treatment, bactericidal activity was observed with a ten time lower concentration of the MIC of tetracycline (0.1 μg/ml, 0.25 μg/ml, and 0.5 μg/ml) combined with low doses of C-dots (1.0 μg/ml, 2.0 μg/ml). An enhanced inhibitory effect was observed when the concentration of C-dot increased to 16 μg/ml and 32 μg/ml. The combination of C-dot and tetracycline indicates an effective bactericidal effect.

The stationary phase was achieved in 5 h compared to the independent treatment, where the bacteria were still in their log phase even after 10 h. The growth rate analysis data predict that mathematical models such as logistic, Gompertz, and Richards function models predict the experimental data very well (Table 1). The growth rate parameter ($\mu_m$), which controls the steepness of the function, is the primary variable that describes the cellular growth. The growth rate parameter ($\mu_m$) shows a significant decrease (~40%) in C-dot and tetracycline samples compared to the control. Attachment of C-dots and cell damage observed with C-dot treatment in the SEM study indicated that the particle and cell surface interaction was prominent with C-dot treatment. There were drastic morphological changes and cell damage observed in the E. coli cells in the synergistic treatment of C-dot together with tetracycline. The AFM study with C-dot and tetracycline also indicates outer membrane roughness in the height profile (amplitude imaging) and cellular morphological changes in phase images compared to the control. The exact mechanism of how these particles interact with the cell surface either directly or indirectly still remains unexplored. This study needs to be extended further with antibiotic-resistant pathogens to identify the intricate dose and other parameters for synergistic use. Thus, this study establishes a framework to combat antibiotic-resistant bacteria, presently demonstrated using E. coli as a model organism and inhibition of growth with a much lower MIC dose of tetracycline concentration when used in combination with C-dot.

### Table 1

|          | $A$  | $\mu_m$ | $\lambda$ | $\nu$ |
|----------|------|---------|-----------|-------|
| Control  |
| Logistic | 2.06 | 0.49    | 2.84      | 1.0   |
| Gompertz | 2.2  | 0.46    | 2.65      | 1.0   |
| Richards | 1.97 | 0.67    | 3.23      | 3.16  |
| Stannard | 1.97 | 0.67    | 3.23      | 3.16  |
| Tet      |
| Logistic | 1.65 | 0.33    | 2.46      | 1.0   |
| Gompertz | 1.81 | 0.31    | 2.22      | 1.0   |
| Richards | 1.62 | 0.38    | 2.56      | 1.48  |
| Stannard | 1.62 | 0.38    | 2.56      | 1.48  |
| C-dot    |
| Logistic | 1.24 | 0.27    | 2.06      | 1.0   |
| Gompertz | 1.31 | 0.25    | 1.87      | 1.0   |
| Richards | 1.22 | 0.29    | 2.12      | 1.33  |
| Stannard | 1.22 | 0.29    | 2.12      | 1.33  |
| C-dot + tet |
| Logistic | 0.85 | 0.28    | 2.23      | 1.0   |
| Gompertz | 0.86 | 0.33    | 2.43      | 1.0   |
| Richards | 0.84 | 0.34    | 2.36      | 2.1   |
| Stannard | 0.84 | 0.34    | 2.36      | 2.1   |

### CONCLUSION

This study demonstrates that the C-dot with tetracycline is effective for synergistic antibacterial treatment with a ten time lower tetracycline concentration than the MIC. Our SEM and AFM imaging data prove the cellular damage and change in the cell topology. We also ascertain our work with logistic, Gompertz, and Richards mathematical model, which suggest that all the models reveal the antibacterial effect as seen in the bacterial growth curve. The Richards model gives the best fit. These models will help understand and extrapolate the minimal effects on bacterial growth. We summarize that C-dot and tetracycline are ideal candidates for antibacterial cocktails and possibly effective treatment for pathogenic MDR bacterial strains with minimal antibiotic dose.

### DATA AVAILABILITY STATEMENT

The original contributions presented in the study are included in the article/Supplementary Material, further inquiries can be directed to the corresponding authors.

### AUTHOR CONTRIBUTIONS

DKT conceptualized the plan and obtained the funding, GJ, DKT, and MT performed antibacterial assay SEM imaging, sample preparation, and data analysis. VVG performed AFM experiments and mathematical analysis of growth curve data. VVG and DKT performed AFM imaging analysis. MT and DKT wrote the manuscript and prepare figures. SK contributed to microbiological structuring. SD and DKT performed characterization of C-dot. All authors read and finalized the manuscript.

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### SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fgene.2021.626276/full#supplementary-material
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