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# Fabrication of biocidal materials based on the molecular interactions of tetracycline and quercetin with hydroxyapatite via In Silico- and In vitro approaches

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# ABSTRACT

Synthetic hydroxyapatite (HA) materials with antibacterial and biocompatible properties have potential for biomedical applications. The application of various computational methods in silica is highly relevant for the optimal development of modern materials. In this work, we used molecular docking to determine the binding constants of tetracycline (TET) and quercetin (QUE) with hydroxyapatite and compared them to experimental data of the adsorption of tetracycline (TET) and quercetin (QUE) on the HA surface. The experimental adsorption study was performed via the UV-VIS method. The fabricated biocidal powders were characterized via X-ray powder diffraction (XRD), Fourier transform infrared (FTIR) spectroscopy, and scanning electron microscopy (SEM). The electrical charge of the HA particle surface was determined via zeta potential measurements. The molecular docking method was used to predict the binding affinities of TET and QUE for HA. We also performed molecular docking studies to predict the binding affinity of TET and QUE for HA. These affinities correlate with the experimental binding constants, suggesting that molecular docking is a good tool for material property prediction. In addition, the antimicrobial activity of the HA/TET and HA/QUE powders was determined against 2 g-positive bacterial strains: S. aureus and E. faecalis. The obtained HA powders were evaluated for biocompatibility in vitro with the myoblast cell line C2C12.

# 1. Introduction

The demand for multifunctional materials, which regulate their physicochemical and biomechanical properties, have therapeutic effects, etc., is high in all areas of science and technology [1-4]. To effectively use such materials for their intended purpose, various aspects of the interaction of their constituent parts need to be studied in more detail: changing the crystal structure and surface charge, determining binding constants, and finding the most favorable spatial conformation [5–8].

Bone tissue engineering is an advanced approach to the development of functional scaffold materials for repairing and regenerating

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damaged bones [9]. A major research direction in this field is to produce scaffolds with desirable shapes, structures, and physical, chemical, and biological features [10]. Hydroxyapatite (HA), a main inorganic component of bones and teeth, is one of the most promising materials in bone tissue engineering [11]. Owing to its biocompatibility, biodegradability, and good adsorption properties, HA has been widely used in the repair of hard tissues and drug delivery [12,13]. However, HA has lower properties than natural bones do, so it is necessary to improve HA-based materials by adding other components [14]. For example, poor antibacterial activity can easily cause bacterial infection and chronic inflammation, eventually resulting in implant failure [15]. Unfortunately, research findings have shown that pure HA has a low ability to deactivate bacteria [16]. As a result, additional components are needed to improve antibacterial activity and maintain biocompatibility. The addition of bioactive phytochemicals such as flavonoids can also inhibit osteoclastogenesis, prevent bone resorption and bone loss, or promote bone formation via the promotion of osteoblast genesis [17].

Tetracycline (TET) is one of the most effective antibiotics and the second most common antibiotic worldwide. It has a broad spectrum of activity and has been extensively used in medical applications to prevent bacterial infections [18]. Quercetin (QUE), a plant flavonoid, notably enhances osteoblast proliferation, differentiation, and mineralization, whereas osteoclast cell proliferation and differentiation are dramatically suppressed by it [19]. Thus, HA-based biomaterials modified with these substances have good prospects for use in tissue engineering. Despite many articles in which various properties of the HA/TET and HA/QUE systems are considered [20–25], further investigations are still needed. In particular, few scientists have resorted to the use of theoretical methods to study the interaction of HA with substances and compared the obtained results with experimental data. However, modern theoretical chemistry is a powerful tool that can provide insights into many phenomena and processes in chemical systems, increasing the efficiency of their use in practice.

In this regard, this study aims to apply molecular docking studies for the prediction of the binding affinity of TET and QUE to HA and the fabrication of biocidal materials based on HA particles adsorbed on the surface of TET and QUE molecules. The novelty of our approach lies in combining theoretical calculations and confirming them with experimental data as the gold standard. Notably, the calculated binding affinity constants of tetracycline and quercetin to the hydroxyapatite surface via molecular docking are correlated with experimental data obtained in the laboratory. UV–VIS spectroscopy was used as a standard experiment to confirm the calculations. In addition, the antibacterial activity of the prepared biocidal HA powders was tested against the bacterial strains *S. aureus* and *E. faecalis*, which are the main causative agents of osteomyelitis and other bone diseases [26,27]. Biocompatibility was evaluated through in vitro experiments in the presence of the C2C12 mouse myoblast cell line. We also performed molecular docking studies to predict the binding affinity of TET and QUE for HA. The molecular docking results were also compared with the experimental binding constants determined via the UV–VIS spectrophotometric method.

## 2. Materials and methods

# 2.1. Materials

Tetracycline hydrochloride ( $C_{22}H_{24}H_{20}$ 8·HCl,  $\geq$ 99.9 %) was purchased from Fisher BioReagents (Thermo Fisher Scientific Inc., Pittsburgh, Pennsylvania, USA). Quercetin dihydrate ( $C_{15}H_{10}O_7 \cdot 2H_2O$ ,  $\geq$ 98.9 %) and acridine orange were obtained from Vecton (Vecton Corp., Saint Petersburg, Russia). Propidium iodide (PI) was obtained from Invitrogen<sup>TM</sup> (Burlington, Canada). Fetal bovine serum (FBS), Dulbecco's modified Eagle's medium (DMEM), and penicillin/streptomycin were purchased from BioloT (BioloT Ltd., Saint Petersburg, Russia).

# 2.2. Synthesis of HA powder

Hydroxyapatite (HA) powder was synthesized according to the literature via a sol-gel method [28]. Briefly, an aqueous solution of 0.3 mol/mL NaH<sub>2</sub>PO<sub>4</sub> was added dropwise to a 1.4 mol/mL CaCl<sub>2</sub> solution at room temperature. To control the pH of the solution, NH<sub>4</sub>OH was also added in a dropwise manner. After the addition was completed, the solution was maturated for 60 min and centrifuged. The as-precipitated HA was oven-dried overnight at 60 °C. The powders were sifted, and a fraction of particles with a size of 81  $\mu$ m was used.

#### 2.3. Adsorption experiments

The adsorption study was conducted by adding 10 mg of HA powder to different tubes containing 1.5 mL of tetracycline and quercetin ethanol solutions at different concentrations (1, 10, 100, or 1000  $\mu$ g/mL) with 10-fold dilutions and shaking in IKA mini G (IKA-Werke GmbH & Co. KG, Staufen im Breisgau, Germany) at 1610×g for 5–10 min at room temperature. After mixing, the supernatants were collected, and the powders were dried in an oven at 60 °C for 3 h. For the kinetics studies, solutions at initial concentrations of 10  $\mu$ g/mL were used. The substance concentration was then analyzed at different time points. The time required for the adsorption of TET and QUE was estimated by decreasing the absorbance peaks in the UV–VIS spectra. The amount of adsorbed biologically active compounds ( $q_e$ ,  $\mu$ g/mg) and the adsorption rate (%) at the equilibrium state were obtained as follows:

$$q_e = \frac{(C_0 - C_e)V}{m} \tag{1}$$

$$\% A dsorbed = \frac{(C_0 - C_e)}{C_0} \bullet 100$$
<sup>(2)</sup>

where  $q_e$  (µg/mg) is the adsorption amount;  $C_0$  and  $C_e$  (µg/mL) are the initial and equilibrium concentrations of the substance in solution, respectively; V (mL) is the volume of the substance solution; and m (µg) is the weight of the adsorbent.

#### 2.4. Characterization method

The HA particle morphology was characterized via a scanning electron microscope (Tescan Vega 3) equipped with an energy dispersive X-ray (EDX) attachment (TESCAN GROUP, a.s., Brno, Czech Republic). An Au film was sputtered onto the samples to enhance conductivity via SPI sputter installation (Structure Probe, Inc., West Chester, PA, USA). The local qualitative elemental compositions of the samples were identified via energy dispersive X-ray analysis. The measurements were carried out with secondary electrons at a voltage of 20 kV. In addition to the conventional SEM image and the local qualitative elemental composition spectra, detailed maps of the elemental distribution were also generated to estimate the allocation of the elements.

Adsorption studies were carried out via a microvolume UV–visible spectrophotometer (Bioevopeak SP-MUV1000; Bioevopeak Co., Ltd., Jinan, Shandong, China) in the range of 190–850 nm. The droplet size was 2  $\mu$ l. The residual amount of the drug in the solution was quantified at 240 nm for TET and 241 nm for QUE. The drug solution in ethanol with an initial concentration of 10  $\mu$ g/mL was used as a baseline.

X-ray analysis of the obtained powders was carried out via CuK $\alpha$  radiation ( $\lambda = 1.5406$  Å) on a D2 PHASER (Bruker, Billerica, Massachusetts, USA) with a 2 $\theta$  range from 10 to 50° and a step size of 0.02. Tetracycline and quercetin powders were used as controls.

The presence of functional groups in pure HA and after the adsorption of tetracycline and quercetin on the HA surface were analyzed via Fourier transform infrared (FTIR) spectroscopy on an FSM 2201/2202 spectrometer (Infraspek Ltd., Saint-Petersburg, Russia). FTIR spectra of the materials were recorded in transmittance mode via KBr pellets over the wavenumber range of 500–4000 cm<sup>-1</sup> at a resolution of 4 cm<sup>-1</sup>. For each spectrum, 30 scans were performed. The mass ratio of the test substance to KBr was 1:100.

The zeta potential of each sample was determined via a ZetaSizer Compact-Z (Photocor Ltd., Moscow, Russia). Measurements were performed in ethanol solutions.

#### 2.5. Antibacterial activity

The antimicrobial activity of the biocidal powders was investigated against the growth of the following pathogens: *Staphylococcus aureus* (ATCC 25912) and *Enterococcus faecalis* (ATCC 29812) microbial strains. The antimicrobial assays were performed according to 0.5 McFarland standard microbial cultures. The experiments to evaluate the antibacterial properties were performed according to the recommendations of the Clinical and Laboratory Standards Institute (CLSI) [29] and the European Committee on Antimicrobial Susceptibility Testing (EUCAST) [30]. Mueller–Hinton (MH) agar and Columbia agar containing 5 % sheep blood were sterilized by autoclaving at  $121 \pm 2$  °C and  $103 \pm 5$  kPa for 30 min. Warm medium (20 mL) was poured into Petri dishes (Nunc Inc., Roskilde, Denmark) and allowed to cool at ambient temperature. Inocula for MICs were prepared in sterile Mueller–Hinton broth (Oxoid). The microorganisms were grown at 37 °C for 24 h. The analysis was carried out in five experiments for each of the analyzed samples. Afterward, the samples were inoculated with 1.5 mL of microbial suspension at a density of  $1 \times 10^8$  CFU/mL, prepared in phosphate-buffered saline (PBS), and incubated for 24 h. Then, 5 µL of each of the tested HA powders was applied in 3 repetitions and incubated on MH agar media for 24 h at 37 °C.

# 2.6. Cell culture

The C2C12 mouse skeletal myoblast cell line was purchased from Sigma Aldrich (Merck KGaA, Darmstadt, Germany and Sigma – Aldrich Co., LLC, St. Louis, Missouri, USA) and cultured in high-glucose DMEM containing 10 % fetal bovine serum (FBS) and 1 % penicillin/streptomycin at 37 °C and 5 %  $CO_2$  in humidified (95 %) air. The culture medium was changed every 3 days. The culture medium was subsequently removed, and the cells were detached from the surface of the 75 cm<sup>2</sup> cell culture flask with 1 mL of 0.25 wt% trypsin solution. To neutralize the trypsin, 5 mL of the culture medium was added.

# 2.7. Acridine orange/propidium iodide staining

The C2C12 myoblast cell line was treated with biocidal powders (1 g/mL) at  $1 \times 10^4$  cells/well in a 12-well culture plate. An aliquot of 20 µL of acridine orange (AO) stock solution (1 mg/mL) was added to each well with PBS and cooled at +4 °C. Then, an aliquot of 20 µL of propidium iodide (PI) stock solution (1 mg/mL) was added to each well and incubated in the dark for 10 min at +4 °C. The fluorescence was measured via a confocal microscope (Leica DMi 8, Germany) with an excitation wavelength of 460 nm, an emission wavelength of 550 nm for AO, an excitation wavelength of 525 nm and an emission wavelength of 595 nm for PI.

#### 2.8. Experimental binding constant

For calculations of the experimental binding constants, we used the radioligand binding assay method. The residual concentration

of the biologically active compound (TET or QUE) in the solution at time *t* was calculated via the following equation:

$$C_t = \frac{A_t}{A_0} \bullet C_0 \tag{3}$$

where  $C_0$  and  $C_t$  are the initial and residual concentrations and where  $A_0$  and  $A_t$  are the absorbance values at time zero and time t, respectively. The absorbance values were retrieved from the highest absorbance peaks (240 nm for TET and 241 nm for QUE). The time t changed from 0 to 530 min when the adsorption process reached equilibrium. Since the experiment was conducted 3 times, the mean and standard deviation were calculated for each  $C_t$  value. The resulting mean values were input into GraphPad Prism 9.5.1 software, and the dissociation rate constants were obtained through the one-phase exponential decay equation:

$$Y = (Y_0 - NS) \bullet exp (-K_d \bullet X) + NS$$
(4)

where  $Y_0$  is the binding at time zero, in units of the Y axis; NS is the binding (nonspecific) at infinite times, in units of the Y axis; and  $K_d$  is the dissociation rate constant in inverse units of the X-axis. The binding constants, or association constants, of TET and QUE to HA were obtained as the inverse of the respective dissociation constants via the following equation:

$$K_a = \frac{1}{K_i} \tag{5}$$

where  $K_a$  and  $K_i$  are the association and inhibition rate constants, respectively [31].

# 2.9. Molecular docking

The 3D coordinates of the TET (CID 54675776) and QUE (CID 5280343) molecular structures were retrieved from the PubChem database as 3D SDF files. The HA structure was generated via the HA unit cell CAR file, which is available from the INTERFACE-MD software developed by the Heinz group at the University of Colorado at Boulder. The HA unit cell consists of six  $PO_4^{3-}$  groups, two  $OH^{-}$  groups, and ten  $Ca^{2+}$  ions. The HA slab was designed via the TopoTools v.1.8 VMD plugin. Molecular docking was performed via the AutoDock and AutoDock Vina (ADVina) programs with default settings. Genetic and gradient optimization algorithms were employed. The ligand-HA binding site at the HA center for each ligand-HA complex was identified via Cartesian coordinates: x = 9.2 Å, y = 15.94 Å, and z = 3.52 Å. Grid maps were created with a grid spacing of 0.375 Å and a dimension size of 60 Å. To prepare the structures for molecular docking, Gasteiger partial charges were assigned, and rotatable bonds were defined according to standard protocols published elsewhere. The docking output results were represented by the approximation function as the estimated Gibbs free energy of binding ( $\Delta G_{bind}$ ).

# 2.10. Statistical analysis

All experiments were reproduced at least three times using three samples from each group. The obtained data were averaged with the standard error of the mean. Statistical analysis to compare the results of the cell antibacterial activity and cell viability assays was performed via ANOVA (\*p < 0.05).



**Fig. 1.** Zeta potentials of pure HA, HA with tetracycline (HA/TET-1, HA/TET-2), and HA with quercetin (HA/QUE-1, HA/QUE-2). HA/TET-1, HA/TET-2, HA/QUE-1, and HA/QUE-2 indicate that tetracycline and quercetin ethanol solutions were used at 1 mg/mL and 100 µg/mL, respectively.

# 3. Results and Discussion

### 3.1. Characterization of synthesized materials

To estimate the possibility of antibiotic adsorption on HA particles, the surface charge of HA particles was initially measured. Fig. 1 shows the results of the surface charge and zeta potential measurements of the HA particles after antibiotic adsorption at 25 °C. As shown in Fig. 1, the initial surface charge of HA is negative and equal to -14.4 mV. Notably, the negative value of the zeta potential correlates with literature data for hydroxyapatite powders synthesized in aqueous solutions [32–35]. In addition, the zeta potential of natural bone-derived HA particles was found to be -9.25 ± 0.9 mV at a typical physiologic pH of 7.4 [36]. The negative values of the zeta potential are suggested to have a significant favorable effect on the attachment and proliferation of bone cells [36]. The adsorption of tetracycline from an ethanol solution led to a significant change in the surface charge to +6.1 mV (Fig. 1, plot for HA/TET-1). However, the changes in the HA surface charge and zeta potential value in the less concentrated solutions (the tetracycline concentration was 100  $\mu$ g/mL) were smaller. The surface charge of the HA/TET-2 particles remains negative and equal to -4.5 mV (Fig. 1, plot for HA/TET-2). The decrease in or change in the zeta potential in the presence of TET may be caused by the lower adsorption of TET on the surface of HA, the greater adsorption of the antibiotic and the thick adsorption layer. The interaction between the negatively charged tetracycline and quercetin with the negatively charged HA is probably due to the interaction of the functional groups of the adsorbed molecules with the hydroxyl groups of the HA surface [31]. This explains the noticeable reduction in the zeta potential of HA particles with adsorbed TET. A higher adsorption level of TET considerably changes the surface charge and zeta potential of HA. Notably, tetracycline has a protonated form via NH<sub>2</sub> groups at the fourth carbon atom in concentrated solutions. It determines the positive charge of a molecule. An increase in the amount of adsorbed TET also increases the number of NH<sup>+</sup><sub>3</sub> groups at the interface that react with the surface of the HA. The positive charge originating from these groups has a direct effect on lowering the negative charge and zeta potential of HA in the presence of TET, as demonstrated in the present study.

The XRD profiles of pure HA and powders after adsorption of tetracycline (HA/TET-1) and quercetin (HA/QUE-1) are shown in Fig. 2A. In all XRD patterns (Fig. 2 A, plots 1–3), there are peaks at 2O: 25.9°, 31.879°, 32.232°, 32.917°, 34.14°, 39.91°, 46.75°, and 49.53°. These peaks correspond to the main Miller planes and correspond to HA crystal angles of 25.879° (002), 31.774° (211), 32.197° (112), 32.902° (300), 34.049° (202), 39.819° (310), 46.713° (222), and 49.469° (213) according to ICDD-PDF# 9–432 and literature data [18,36–38]. The characteristic peaks of tetracycline and quercetin were not detected at 10–50° in the HA/TET-1 and HA/QUE-1 composites, respectively. This could be related to the amorphous structure of tetracycline and quercetin [38,39]. Owing to this phenomenon, samples (HA/TET-2 and HA/QUE-2) obtained from less concentrated solutions are not included in Fig. 2B. In addition to zeta potential measurements, FTIR analysis was used to estimate the possibility of adsorption of tetracycline and quercetin on HA or binding between HA particles and tetracycline or quercetin.

Fig. 2B shows the FTIR spectra of the pure HA, HA/TET-1, and HA/QUE-1 composites. In the spectra of all the samples, there are two intense peaks at 2355 cm<sup>-1</sup>, and those at 3740 cm<sup>-1</sup> are associated with OH<sup>-</sup> groups [40]. The peak at 1049 cm<sup>-1</sup> is attributed to  $PO_4^3$  vibrations [41]. The two peaks at 1427 cm<sup>-1</sup> and 1462 cm<sup>-1</sup> are associated with  $CO_3^2$  impurities [42]. The  $PO_4^3$  peak decreases with increasing content of tetracycline in HA, whereas a strong peak attributed to the OH<sup>-</sup> group at 2355 cm<sup>-1</sup> [43,44] can be observed in all the samples (Table 1). To study the interaction between the hydroxyapatite surface and the adsorbed molecules, we deconvoluted the FTIR spectra (Fig. 2C–H). Deconvolution of the spectra separates overlapping bands in the  $v_4PO_4^3$ ,  $v_2CO_3^2$ ,  $v_3CO_3^2$ - and amide regions, allowing a more detailed analysis of changes at the atomic level [45]. The  $PO_4^3$  peak is deconvoluted into two peaks, which could be due to its overlap with the HPO\_4^2- apatite peak. Usually, separating these peaks and distinguishing them from each other by colorimetry are very difficult [45]. Fig. 2C–H shows that the intensity of the OH<sup>-</sup> band vibration is lower in the HA/QUE-1 and HA/TET-1 composites than in pure HA. This may be due to the interaction and formation of hydrogen bonds between TET or QUE and the hydroxyapatite surface. The interaction of positively (protonated form) charged tetracycline and quercetin with

Table 1
Comparison of the bands observed in the FT-IR spectra of the samples with the literature data

Sample	Domain, Assignment	IR (cm <sup>-1</sup> )	Reference
Pure HA	$v_3 PO_4^{3-}$ and $HPO_4^{2-}$ symmetric and asymmetric stretching	1052, 1101	41
HA/TET-1		1052, 1096	
HA/QUE-1		1054, 1097	
Pure HA	$v_2 CO_3^{2-}$ type A1 doublet	1531	40
HA/TET-1		1531	
HA/QUE-1		1530	
Pure HA	H <sub>2</sub> O bending	1694	3,4, 40
HA/TET-1		1693	
HA/QUE-1		1693	
Pure HA	H <sub>2</sub> O	3290	40
HA/TET-1	Symmetric stretching	3269	
HA/QUE-1		3282	
Pure HA	OH <sup>-</sup> stretching	3741	40, 43, 44
HA/TET-1		3738	
HA/QUE-1		3739	



**Fig. 2.** A) XRD profiles of pure HA particles (plot 1), HA/TET-1 (plot 2) and HA/QUE-1 (plot 3). HA/TET-1 and HA/QUE-1 indicate that tetracycline and quercetin ethanol solutions, respectively, were used at a concentration of 1 mg/mL. The HA peaks with intensities greater than 15 % are highlighted as (\*). B) FTIR spectra of pure HA particles (plot 1), HA/TET-1 (plot 2) and HA/QUE-1 (plot 3). C–H) Deconvolution of pure HA and HA/QUE-1 (C), pure HA and HA/TET-1 (F) showing the  $v_4PO_4^{3-}$  and  $HPO_4^{2-}$  bands (800–1300 cm<sup>-1</sup>). Deconvolution of the  $v_3CO_3^{2-}$  band (1300–1900 cm<sup>-1</sup>) for 80 °C dried pure HA and HA/QUE-1 (D), pure HA and HA/TET-1 (G). Deconvolution of the  $a_1H_2O$  and  $a_1HO^{-}$  bands (2800–3800 cm<sup>-1</sup>).

negatively charged HA is probably due to the interaction of the functional groups of the adsorbed molecules with the hydroxyl groups of the HA surface [44].

In addition, in a plot of the HA/QUE-1 composite, we also observed an increase in peak intensity at approximately  $570 \text{ cm}^{-1}$ , which is responsible for the C–H bending of aromatic hydrocarbons (out-of-plane) [43]. Thus, we can assume that tetracycline and quercetin, which have aromatic groups, are adsorbed on the HA surface.

The surface morphology of pure HA was analyzed via SEM and is shown in Fig. 3A–G. The pure HA particles are roundish conglomerates of *ca.* 20–40  $\mu$ m. The conglomerates consist of hexagon-shaped structures. The surface of the particles is smooth, and the diameter of the pores is approximately 100 nm. Elemental mapping analysis revealed calcium and phosphorous in the pure HA, HA/ TET-1 and HA/QUE-1 powders (Fig. 3H–J). A uniform distribution and good dispersion of Ca, P, O and C throughout the surface of the HA were observed. The quantitative chemical compositions of HA/TET-1 and HA/QUE-1 were estimated from EDX (Table 2). The EDX spectra presented in Fig. 3 K–M reveal that the pure HA, HA/TET-1 and HA/QUE-1 composites are composed of four main elements, Ca, P, O and N. The presence of the N constituent element from the tetracycline in the HA/TET-1 powder was statistically significant. Therefore, we did not include nitrogen in the results. EDX analysis revealed that the Ca/P ratio in the samples varied from 1.7 for pure HA to 1.61  $\div$  1.67 for the HA/TET-1 and HA/QUE-1 composites. Thus, we can assume a high stoichiometry of hydroxyapatite [46].

However, we indirectly determined the presence of adsorbed tetracycline and quercetin molecules in the sample. The HA/TET-1 and HA/QUE-1 composites were soaked in distilled water for 24 h, and then, the resulting solutions were analyzed via the UV–VIS method. The resulting composites released tetracycline and quercetin when kept in solution.

# 3.2. Adsorption kinetics studies

The adsorption of tetracycline and quercetin on the HA powders was also confirmed via the UV–VIS method. UV–VIS spectroscopy revealed a peak at approximately 240 nm in the spectra of HA/TET-1 and HA/QUE-1. A weaker peak at 377 nm was also observed in the case of free quercetin (Fig. 4A and D). The diluted solutions of tetracycline and quercetin were used to determine the binding



**Fig. 3.** A–G) SEM images of pure HA (A, D), HA/TET-1 (B, E), and HA/QUE-1 (C, G). HA/TET-1 and HA/QUE-1 indicate that tetracycline and quercetin ethanol solutions, respectively, were used at 1 mg/mL. H–J) Elemental mapping of the pure HA particle surface (H), HA/TET-1 particle surface (I), and HA/QUE-1 particle surface (J) presented for Ca P, O and C. K–M) EDX spectra of the pure HA (K), HA/TET-1 (L) and HA/QUE-1 (M) powders.

## Table 2

The elemental composition of the pure HA, HA/TET-1 and HA/QUE-1 powders was estimated from EDS analysis. HA/TET-1 and HA/QUE-1 indicate that tetracycline and quercetin ethanol solutions, respectively, were used at 1 mg/mL.

Sample	Atomic Composition (%)				
	Са	Р	0	С	
HA	20.41	12.01	48.04	19.54	
HA/TET-1	18.80	11.65	47.78	21.78	
HA/QUE-1	21.14	12.64	46.45	19.78	

constant.

Binding constants provide a fundamental measure of the affinity of a solute to a ligand, and their determination is an important step in describing and understanding molecular interactions. The binding constant is a basic experimental parameter used in a variety of studies, such as the prediction of drug efficiency or drug interactions. In general, a large binding constant between the carrier and the drug is required to maintain a stable complex and improve bioavailability after implantation. The concentrations of tetracycline and quercetin gradually decreased in the solutions with immersed HA particles. The experiment was carried out until the residual value reached equilibrium. The adsorption amounts and removal rates of TET and QUE are shown in Table 3. Optical spectrophotometry revealed that HA particles adsorbed approximately 1.34 µg per milligram and 1.23 µg of tetracycline and quercetin, respectively.

In the case of TET, the residual concentration of the substance in the solution will reach the equilibrium state later than in the case of QUE, which is fully consistent with the obtained values of the binding constants.



**Fig. 4.** UV–VIS spectra of TET (A) and QUE (D) ethanol solutions at an initial concentration of 1000 μg/mL after the HA powder was adsorbed; molecular docking affinity values of TET (B) and QUE (E) on the HA surface; experimental binding constants of TET (C) and QUE (F) on the HA surface. The adsorption experiment lasted 9 h until the system reached equilibrium.

#### Table 3

Adsorption characteristics and predicted equilibrium constants of TET and QUE on the HA surface. The initial substance concentration in the ethanol solutions was 10 µg/mL in the experimental settings.

Substance	Average adsorption amount ( $\mu g/mg$ )	Adsorption rate (%)	Κί (μΜ)	Ka(pred) ( $\mu M^{-1}$ )
TET	1.34	89	28.68	0.035
QUE	1.23	82	217.18	0.005

All these observable effects could be explained by the chemical structure of tetracycline, which contains multiple functional groups that are able to interact with the HA surface through a combination of electrostatic, hydrogen bonding, and van der Waals interactions. Quercetin, on the other hand, has a simpler chemical structure and fewer functional groups that can interact with the HA surface.

# 3.3. Molecular docking

In our study, the experimental binding constants were compared with the binding affinities of tetracycline hydrochloride and quercetin dihydrate to the HA surface, which were calculated via molecular docking (Fig. 4B and E). Molecular docking is a widely known computational method that can be used to predict the binding affinity of small molecules to a target protein or biomolecule.

The binding constants have been calculated via molecular docking because this widely known computational method can be used to predict the binding affinity of small molecules for a target protein or biomolecule. TET and QUE are two small molecules that have been shown to have potential therapeutic effects, and their binding affinity to HA is of interest for understanding their potential as drug candidates.

Our research revealed that TET exhibited greater affinity for HA than did QUE. This was due to the lower Gibbs free energy of binding ( $\Delta G = -6.2 \text{ kcal/mol}$ ) for TET than for QUE ( $\Delta G = -5.0 \text{ kcal/mol}$ ). Gibbs free energy is a thermodynamic parameter that describes the energy change that occurs when a system undergoes a chemical reaction. In the case of molecular docking,  $\Delta G$  can be used to estimate the strength of the interaction between the receptor and the ligand. The lower  $\Delta G$  value for TET indicates that it has a stronger binding affinity to HA. This finding correlates with the experimental results, where TET has been found to have a higher

binding constant to HA than does QUE. The higher binding affinity value of TET to HA suggests that TET may be a more effective drug candidate for applications such as drug delivery and bone regeneration.

Our studies revealed that TET has a greater binding constant to HA ( $K_a = 78.74 (\mu g/mL)^{-1}$ ) than to QUE ( $K_a = 50.4 (\mu g/mL)^{-1}$ ) and a lower dissociation constant (Fig. 4C and F). This means that TET is removed from the solution more slowly and that equilibrium is reached later (but with tighter binding). In practice, the absorbance values practically do not change after 230 min for QUE and after 350 min for TET (Fig. 4A and D).

The results obtained from the adsorption experiments and molecular docking could be explained by the chemical structure of tetracycline, which contains multiple functional groups able to interact with the HA surface through a combination of electrostatic, hydrogen bonding, and van der Waals interactions. Quercetin, on the other hand, has a simpler chemical structure and fewer functional groups that can interact with the HA surface.

# 3.4. Antibacterial activity

Fig. 5A shows the antibacterial activity of the biocidal powders. The samples were examined via the disk diffusion method using *E. faecalis* and *S. aureus*. As shown in Fig. 5B, the inhibition zone revealed that HA particles loaded with TET from concentrated solutions (HA/TET-1) had strong antibacterial activity against 2 g-positive bacterial strains, *S. aureus* and *E. faecalis*. Notably, tetracycline and quercetin bind to HA at ratios of 1.34 µg per 1 mg of HA and 1.23 µg per 1 mg of HA, respectively. According to the literature [47–49], most gram-positive bacteria are sensitive to a TET concentration of 1 µg/mL, whereas most gram-negative bacteria are sensitive to 1–25 µg/mL TET. Quercetin also inhibits the growth of various bacterial isolates, including *Escherichia coli* and *Staphylococcus aureus*, with minimum inhibitory concentrations (MICs) ranging from 20 to 400 mg/mL [50]. Since we used 100 µg of the composites, the resulting concentration that inhibited bacterial growth was below the specified limit. This finding could prove the effectiveness of tetracycline as an additional component that gives hydroxyapatite the desired antibacterial properties. For HA powders with low concentrations of tetracycline and quercetin, no zone of inhibition was observed. The diameters of the inhibition zones of HA/TET-1 against *S. aureus* and *E. faecalis* (region 3) are 15 and 11 mm, respectively. For sample HA/TET-2 (region 4), the inhibition zone diameters are approximately 7 and 2 mm, respectively. In comparison, the HA sample taken as a control with quercetin (region 6) did not show antibacterial activity against either strain. This could be related to a decrease in the amount of quercetin adsorbed on HA particles [51].

#### 3.5. Cytotoxicity

Since biomaterials for bone tissue restoration should have not only bactericidal properties but also osteoinductive potential, the biocompatibility of the obtained powders was tested using C2C12. The choice of TET is because this antibiotic is well studied and has been on the world market for a long time [51,52]. The choice of QUE is because it is not hemolytic [53,54]. Moreover, it rather exhibits hemolysis inhibition properties. At a concentration of 10  $\mu$ g/mL, quercetin inhibited hemolysis by 35.5 %, ranking third among the tested flavonoids for this effect1. Additionally, quercetin has been shown to reduce hemolytic activity in the context of bacterial infections, indicating its protective role against erythrocyte lysis23. Overall, quercetin has beneficial effects on red blood cells rather than causing hemolysis [53].

The mouse myoblast C2C12 cell line is known to differentiate into myoblasts with further potential to differentiate into osteoblasts [54]. The viability of C2C12 cells treated with biocidal powders was confirmed by acridine orange/propidium iodide staining (Fig. 6 A). The fluorescent dye acridine orange (AO) can bind to nucleic acids within cells, causing altered spectral emission [55]. Double-stranded DNA, which is a sign of a living system, is colored green by AO, whereas damaged DNA is colored red [56]. Another



**Fig. 5.** (A) Antibacterial activity of HA powders with tetracycline ethanol solutions: 1 - tetracycline ( $500 \ \mu g/mL$ ), 2 - tetracycline ( $10 \ \mu g/mL$ ), 3 - tetracycline ( $1 \ m g/mL$ ), 4 - tetracycline ( $100 \ \mu g/mL$ ), and 5 - tetracycline ( $1 \ \mu g/mL$ ). (B) Inhibition zones of *S. aureus* and *E. faecalis* were treated with two samples of HA with tetracycline (HA/TET-1 and HA/TET-2, respectively; tetracycline ethanol solutions were used at 1000  $\mu g/mL$  and 100  $\mu g/mL$ ).



**Fig. 6.** (A) Fluorescence microscopy images of C2C12 cells treated with HA/TET-1 and HA/QUE-1 samples after 24 h of cultivation. The green fluorescence indicates viable cells, and the red fluorescence indicates the viable nucleus of cells. (B) Cell density (number of cells/mm<sup>2</sup>) of C2C12 cells cultured for 24 h, 48 h and 72 h in the presence of the control, HA/TET-1 and HA/QUE-1 samples. Mean  $\pm$  SE, \*P < 0.05 (one-way ANOVA). The samples were tested per area type, and three independent experiments were conducted. Glass was used as a control, and pure HA particles were used as a control2. \*\*p < 0.05, n = 3.

dye that binds to nucleic acids, propidium iodide (PI), always fluoresces with an orange-red color [57]. In the case of HA adsorbed on its surface of tetracycline (HA/TET-1), prominent red fluorescence could be visualized, possibly due to the penetration of PI stain through the damaged cell membrane. This may be due to the diffusion of tetracycline through the cell membrane [58].

Fig. 6B shows that there was no significant difference in the cellular density of the control sample. In contrast, in HA/TET-1 and HA/QUE-1, there was a significant decrease (P < 0.05) in the number of C2C12 cells compared with that in the control sample. However, the cell density of the HA/QUE-1 sample was greater than that of the HA/TET-1 sample. In addition, the cell viability for all the samples was high even after contact with the HA/TET-1 and HA/QUE-1 samples. These results are also consistent with the literature data. Researchers have repeatedly shown that HA itself is not cytotoxic [59,61]. However, HA particles are capable of adsorbing nutrients from the medium and are hygroscopic, which may slightly reduce cell growth in control 2. It has also been demonstrated that the addition of components such as tetracycline and quercetin to biomaterials does not significantly increase cytotoxicity in relation to mouse and human fibroblasts and osteoblast-like cells [60–64]. In our study, we confirmed the absence of cytotoxicity for the HA/TET and HA/QUE systems.

# 4. Conclusions

In this work, we analyzed the adsorption of tetracycline and quercetin on the surface of hydroxyapatite. The charge of HA particles changes from -14.4 mV to +6.1 mV upon adsorption of a concentrated solution of tetracycline and to 5.4 mV for adsorption of quercetin. The results of the antibacterial tests revealed that the HA samples with tetracycline had high antibacterial activity against 2 gram-positive bacteria: *S. aureus* and *E. faecalis*. Composites with tetracycline and quercetin molecules adsorbed on the surface exhibit antibacterial activity at much lower concentrations than the same substances in unbound form in solution. In addition, the results of the cell experiments revealed that HA composites with tetracycline and quercetin have good biocompatibility.

Our study also provides important insights into the use of TET and QUE in medical applications. Our findings revealed that TET has greater binding affinity to HA than does QUE, likely due to its lower Gibbs free energy of binding. These results are consistent with

experimental data, where we have shown that TET has a larger binding constant to HA than does QUE. These findings suggest that TET may be a more effective drug candidate for applications such as drug delivery and bone regeneration and highlight the potential of molecular docking as a tool for drug discovery and biomaterial development.

## CRediT authorship contribution statement

Anastasiia M. Isakova: Writing – original draft, Visualization, Validation, Investigation, Formal analysis. Maxim A. Kutyrev: Writing – original draft, Visualization, Validation, Investigation, Formal analysis. Aleksandra S. Kudasheva: Validation, Investigation. Elizaveta V. Rogacheva: Methodology, Investigation, Formal analysis. Lyudmila A. Kraeva: Writing – review & editing, Supervision, Project administration, Methodology, Investigation, Conceptualization. Sergey Shityakov: Writing – original draft, Supervision, Investigation, Formal analysis. Mikhail V. Zhukov: Writing – original draft, Investigation. Sviatlana A. Ulasevich: Writing – review & editing, Supervision, Project administration, Methodology, Investigation, Funding acquisition, Conceptualization. Ekaterina V. Skorb: Writing – review & editing, Supervision, Resources, Project administration, Conceptualization.

# Data availability statement

Data will be made available on request.

# Ethical statement

This study did not involve human or animal subjects, and thus, no ethical approval was required.

# Declaration of competing interest

The authors declare that they have no known competing financial interests or personal relationships that could have appeared to influence the work reported in this paper.

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