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EDITED BY

Xuetao Xu,
Wuyi University, China

REVIEWED BY

Guanghao Zhu,
Shanghai University of Traditional Chinese
Medicine, China
Danying Huang,
Guangdong University of Petrochemical
Technology, China
Haoxing Zhang,
Shenzhen University, China

*CORRESPONDENCE

Faliang Liang,
✉ faliang2024@163.com

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Inhibition mechanism investigation of quercetagenin as a potential tyrosinase inhibitor

Faliang Liang*

Pharmacy Department, Jiang Men Maternity and Child Healthcare Hospital, Jiangmen, China

Tyrosinase is one important rate limiting enzyme in melanin synthesis, directly affecting the melanin synthesis. Quercetagenin is one active ingredient from marigold. Thence, the inhibition effects of quercetagenin against tyrosinase were investigated. The results showed quercetagenin could inhibit tyrosinase activity with IC_{50} value of 0.19 ± 0.01 mM and the inhibition type was a reversible mixed-type. Results of fluorescence quenching showed quercetagenin could quench tyrosinase fluorescence in static process. CD and 3D fluorescence results showed the interaction of quercetagenin to tyrosinase could change tyrosinase conformation to inhibit activity. Moreover, docking revealed details of quercetagenin's interactions with tyrosinase.

KEYWORDS

nature product, quercetagenin, tyrosinase, inhibition effects, inhibitor

1 Introduction

As we all know, various diseases affect people's health (Zhang et al., 2020; Zhang X. et al., 2021; Shi et al., 2023; Zhou S. et al., 2022). As a therapeutic target, tyrosinase is one important metal enzyme containing two copper, which widely presents in the organism (Min et al., 2023; Fan et al., 2017). Tyrosinase has been confirmed to be involved in the synthesis of melanin (Li et al., 2023; Hassan et al., 2023; Li J. et al., 2021). The first reaction process is monophenolase activity, in which L-tyrosine is hydroxylated into L-dopa and the second reaction process is diphenolase activity, in which L-dopa is subsequently oxidized into dopaquinone (Djafarou et al., 2023; Lee et al., 2023; Zargaham et al., 2023). In the organism, melanin acts crucial roles to protect the skin from UV radiation (Lu et al., 2023a; Znajafi et al., 2023). But, excessive elevated melanin content leads to lots of pigmentation disorders, including age spots and melanoma (Broulier et al., 2023; De Barros et al., 2023; Xue et al., 2023). Inhibition of tyrosinase activity would reduce melanin generation, thence, the finding on novel tyrosinase inhibitors is attracting more attention due to their potential application in medicine and cosmetics fields (Wang and Mu, 2021). Although kojic acid and arbutin (Figure 1) are applied as tyrosinase inhibitors in the medical and industrial fields, they still have been found to lots of adverse side effects (Wang et al., 2021; Esma et al., 2023). Now, to find novel tyrosinase inhibitors is essential for treatment of melanin synthesis (Romagnoli et al., 2022; Nasab et al., 2023).

Natural products have been the vital sources for clinical drugs (Li Y. et al., 2021; Wu et al., 2021; Chen et al., 2022; Zang et al., 2022). Many natural products display widely biological activities, such as antioxidant (Zhang Y. et al., 2021; Tao et al., 2022; Tang et al., 2023), anti-tumor (Liu et al., 2022; Chen et al., 2023; Song et al., 2023), anti-inflammatory (Wang et al., 2020; Wang X. et al., 2022; Zhou Y. et al., 2022), and so on (Wang Y. et al., 2022; Qi et al., 2022; Ma et al., 2023). In particular, natural products show low toxicity (Shao et al., 2020; Jiang et al., 2021;

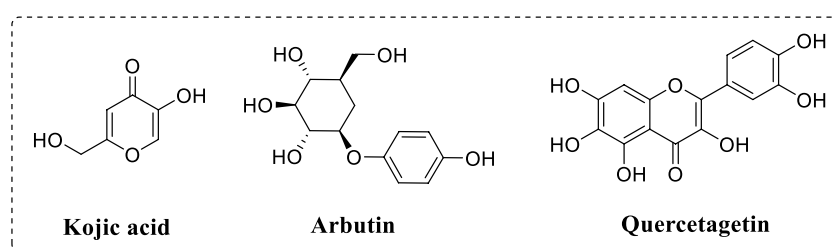


FIGURE 1
Chemical structure of kojic acid, arbutin and quercetagenin.

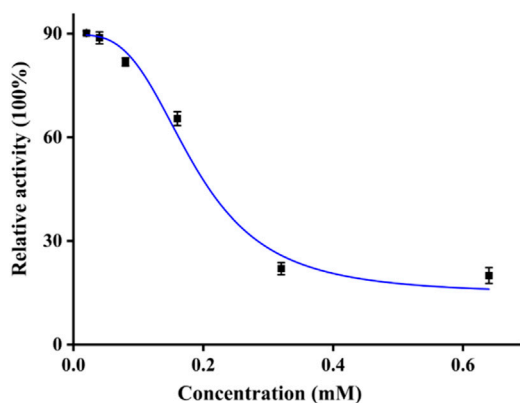


FIGURE 2
Tyrosinase inhibitory activity of quercetagenin.

Pei et al., 2021). Thence the development of natural products as tyrosinase inhibitors attracts much attention.

Quercetagenin (Figure 1) is one active ingredient from marigold and has a chemical structure of 3,3',4',5,6,7-hexahydroxyflavone (Bulut and Yilmaz, 2021; Wang X. et al., 2022). As one polyhydroxyphenol molecule with multiple hydrogen donor substituents, quercetagenin represents rich biological activities (Wu F. et al., 2023). For example, quercetagenin shows strong antioxidant activity and can effectively scavenge DPPH and ABTS (Fuentes et al., 2021). Quercetagenin also enhances the antioxidant enzymatic activities in broilers tissue by Nrf2/ARE signal pathway (Wu et al., 2022). In addition, quercetagenin displays immunomodulatory and anti-inflammatory to inhibit the release of macrophage-derived chemokine (Rufino et al., 2021). Besides, quercetagenin display other beneficial functions, including anti-virus and anti-diabetes (Seyedi et al., 2016). However, to our knowledge, the inhibition effects of quercetagenin against tyrosinase have not been reported yet.

Hence, in this study, we investigated the inhibition effects of quercetagenin against tyrosinase by the multispectral method, followed by the molecular docking.

2 Results and discussion

2.1 Inhibitory activity

The tyrosinase inhibitory activity of quercetagenin was examined on mushroom tyrosinase. The tyrosinase activity was measured

under different concentrations of quercetagenin (Figure 2). It could be observed that tyrosinase relative activity gradually reduced with quercetagenin concentration (0–0.64 mM), meaning that quercetagenin could inhibit the tyrosinase activity with quercetagenin concentration. Its IC_{50} value was calculated to be 0.19 ± 0.01 mM, which was lower than that of kojic acid. This result showed that quercetagenin could be used as a natural tyrosinase inhibitor.

2.2 Kinetic study

There are two kinds of inhibitors, reversible and non-reversible inhibitors. For a reversible inhibitor, it can reduce the enzyme activity by the binding to enzyme, which can restore the enzyme activity through the remove of inhibitor. There are three reversible inhibitors, including competitive, non-competitive and mixed-type inhibitors. The inhibition type of quercetagenin on tyrosinase was subsequently investigated. With different concentration of quercetagenin and tyrosinase, the absorbance changes were measured (Figure 3A) and found that the lines of quercetagenin with different concentration passed origin and slopes reduced with quercetagenin concentration. The results suggested quercetagenin as a reversible inhibitor. With different concentration of quercetagenin and substrate, the absorbance changes were analyzed using Lineweaver-Burk plots (Figure 3B). The lines of quercetagenin intersected in the third quadrant and their slopes increased with quercetagenin concentrations. The results indicated that quercetagenin inhibited tyrosinase in a mixed-type, meaning that quercetagenin bound to tyrosinase and tyrosinase-substrate complex to inhibit its activity. Similar phenomena were obtained in inhibition type of indole-carbohydrazides (Iraji et al., 2022).

Moreover, as shown in Figures 4A, B, the inhibition constants K_i and K_{is} were obtained from the secondary curves of inhibitory kinetics to be 0.24 and 0.12 mM. Smaller K_{is} value than K_i meant that binding force of quercetagenin with tyrosinase-substrate complex was stronger than with tyrosinase. That was to say that quercetagenin preferred to bind with tyrosinase-substrate complex.

2.3 Fluorescence quenching

The fluorescence quenching process of tyrosinase by quercetagenin was investigated. For fluorescence spectra of tyrosinase, they all showed characteristic peaks at 344 nm at 295,

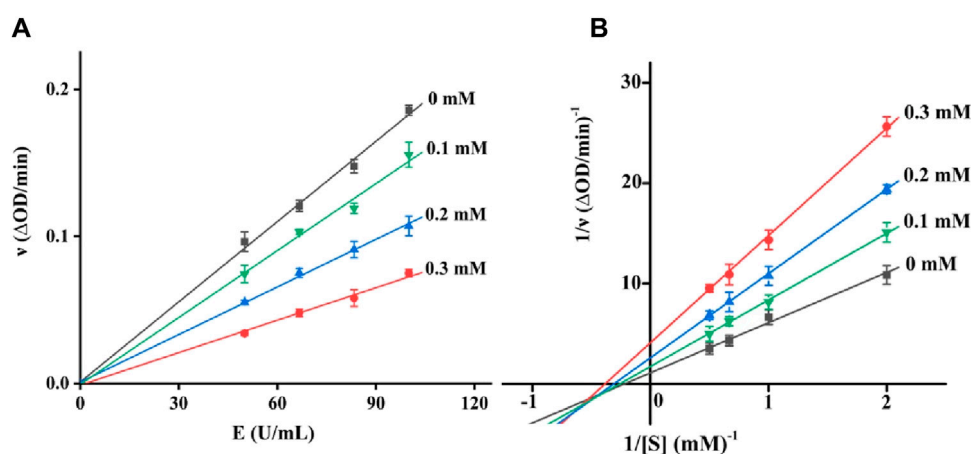


FIGURE 3 (A) Reversible assay of quercetagenin; (B) Lineweaver-Burk plots of quercetagenin.

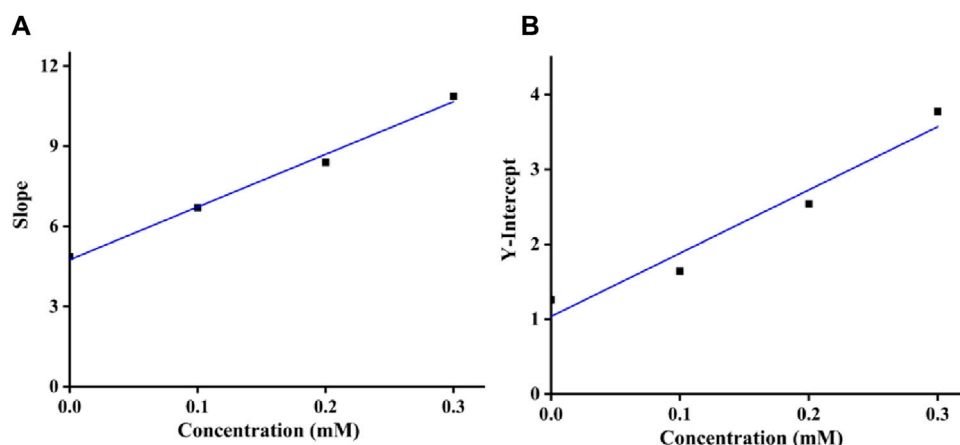


FIGURE 4 (A) K_i assay of quercetagenin; (B) K_{is} assay of quercetagenin.

298, and 305 K, respectively (Figures 5A–C). But, quercetagenin did not show effective fluorescence spectra. Moreover, when treated by quercetagenin, tyrosinase presented the gradually decreasing peak intensity (Figures 5A–C), which indicated that quercetagenin bound to tyrosinase.

The fluorescence quenching data at 295, 298, and 305 K were further analyzed by Stern-Volmer plots. As is shown in Figure 5D the Stern-Volmer plots of quercetagenin, lines with different temperature presented good linearity, meaning that there was only one quenching type of static or dynamic in quenching process. Then the quenching constant (K_{sv}) the bimolecular quench rate constant (K_q) were obtained (Table 1). The K_{sv} results found that K_{sv} values decreased with the temperature. And K_q values at 295, 298, and 305 K were higher than $2 \times 10^{10} \text{ L mol}^{-1} \text{ S}^{-1}$. Above results suggested the quenching process of quercetagenin against tyrosinase was a static process. This quenching type presented in the quenching of indole derivatives on α -glucosidase (Hu et al., 2024). The quenching process also was

the process of thermodynamic changes. Thence, the thermodynamic parameters were calculated (Table 1). The negative ΔG value unlocked one spontaneous process of quercetagenin binding to tyrosinase. The positive ΔH and ΔS values suggested hydrophobic interactions as the important forces between quercetagenin and tyrosinase.

2.4 CD spectra

The effect of quercetagenin on conformation change of tyrosinase was investigated using CD spectra. Figure 6 showed CD spectra of tyrosinase, the characteristic peaks around 210–220 nm stood for the peptide chains of tyrosinase. When treated with quercetagenin, the CD spectra of tyrosinase appeared some changes, meaning its conformation change. Then its secondary structure contents were calculated (Table 2) and found that quercetagenin treatment (molar ratio: 2:1) caused reduction of

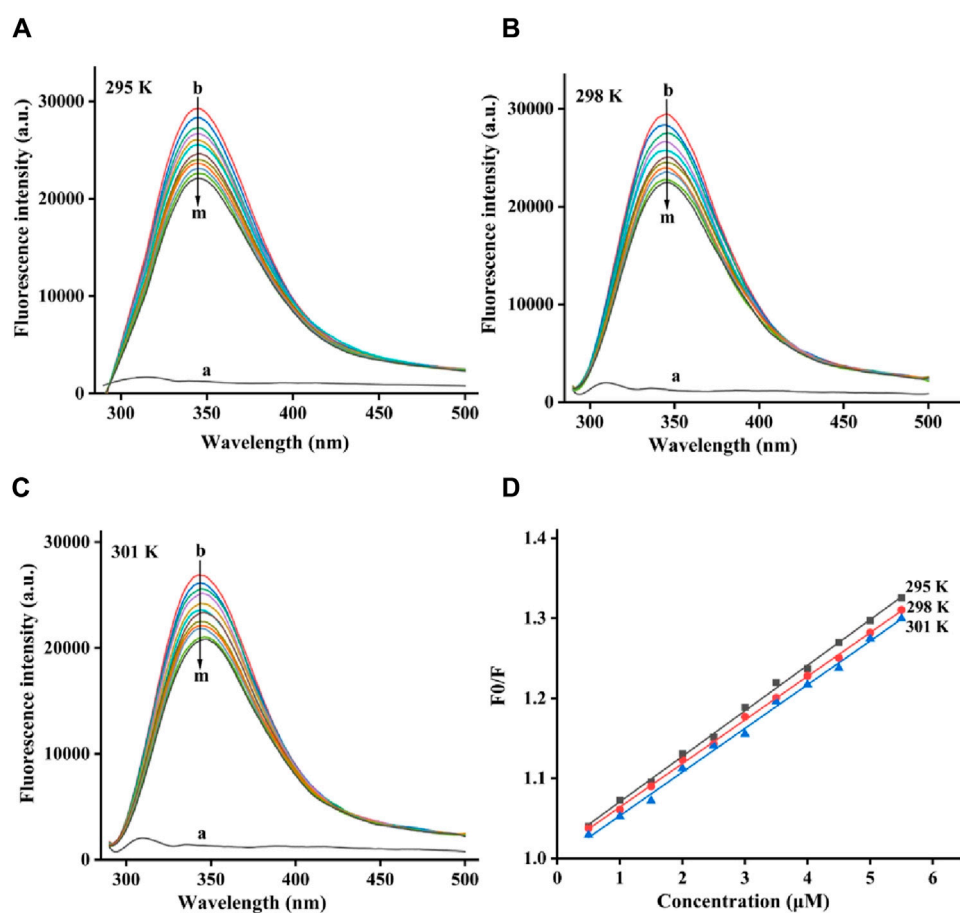


FIGURE 5 (A–C) Fluorescence quenching spectra of quercetagenin on tyrosinase at 295, 298, and 305 K, respectively; (D) Stern-Volmer plots of quercetagenin.

TABLE 1 Parameters of quercetagenin with tyrosinase.

T (K)	K_q ($\times 10^{12}$ Lmol $^{-1}$ S $^{-1}$)	K_{sv} ($\times 10^4$ Lmol $^{-1}$)	ΔH (KJ/mol)	ΔG (KJ/mol)	ΔS (J/(mol·K))
295	5.69	5.69	25.34	-23.27	164.78
298	5.45	5.45		-23.76	
301	5.40	5.40		-24.26	

α -helix and random coils and increase of β -sheet and β -turn. These results indicated that quercetagenin treatment could lead to conformation changes of tyrosinase.

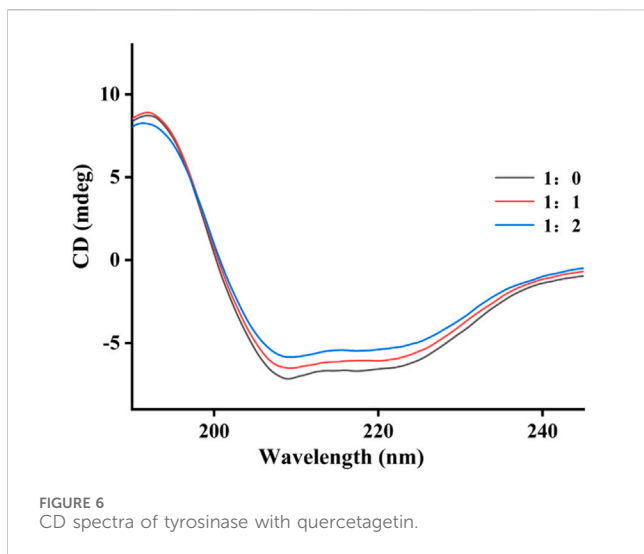
2.5 3D fluorescence spectra

3D fluorescence spectra were monitored to further investigate the effect on conformation of tyrosinase by quercetagenin. As shown in Figure 7A the 3D fluorescence spectra tyrosinase, two characteristic peaks appeared, including Peak A for Tyr and Trp residues and Peak B for peptide backbone. When treated with quercetagenin, Peak A and Peak B were reduced the intensity by 35.4% and 13.3%, respectively (Figure 7B), which suggested that quercetagenin treatment would cause the changes of Tyr and Trp

residues, and peptide backbone. That was to say that quercetagenin treatment would cause the conformation change of tyrosinase.

2.6 Molecular docking

The interaction of quercetagenin to tyrosinase was simulated by molecular docking method. As shown in Figure 8A the docking results, quercetagenin bound to the active pocket of tyrosinase with trihydroxychromone section in the active catalytic zone of active pocket. From Figure 8B the docking results in detail, it could be observed that quercetagenin made one hydrogen bond with His259 and hydrophobic interaction with Val248, His263, Val283, and Ala286. The results indicated that the interactions



between quercetagenin and tyrosinase resulted in the reduction of tyrosinase activity.

2.7 Copper-chelating activity

The copper-chelating activity of quercetagenin was finally assayed using copper sulfate. Figure 9A showed the quercetagenin-copper sulfate mixture. Quercetagenin showed its UV spectra with characteristic peak at 370 nm. While, after added copper sulfate, the UV characteristic peak of quercetagenin was gradually decreased. These results indicated that copper might complex with

quercetagenin to change its UV characteristic peak. The peak intensity at different molar ratios of quercetagenin to copper sulfate was analyzed (Figure 9B) and found that the decreasing trend of quercetagenin peak intensity become equilibrium, meaning that the binding molar ratio of quercetagenin with copper sulfate was 1. The copper-chelating activity of quercetagenin also might be one reason of the reduction of tyrosinase activity.

3 Conclusion

Tyrosinase is one important rate limiting enzyme in melanin synthesis, directly affecting the melanin synthesis. Quercetagenin is one active ingredient from marigold. In this study, the inhibition effects of quercetagenin against tyrosinase were investigated. The results showed that quercetagenin could inhibit the tyrosinase activity with IC_{50} value of 0.19 ± 0.01 mM and the inhibition type was a reversible mixed-type. Results of fluorescence quenching showed that quercetagenin could quench tyrosinase fluorescence in static process. CD and 3D fluorescence results showed interaction of quercetagenin to tyrosinase could change tyrosinase conformation to inhibit activity. Moreover, docking revealed details of quercetagenin's interactions with tyrosinase.

4 Materials and methods

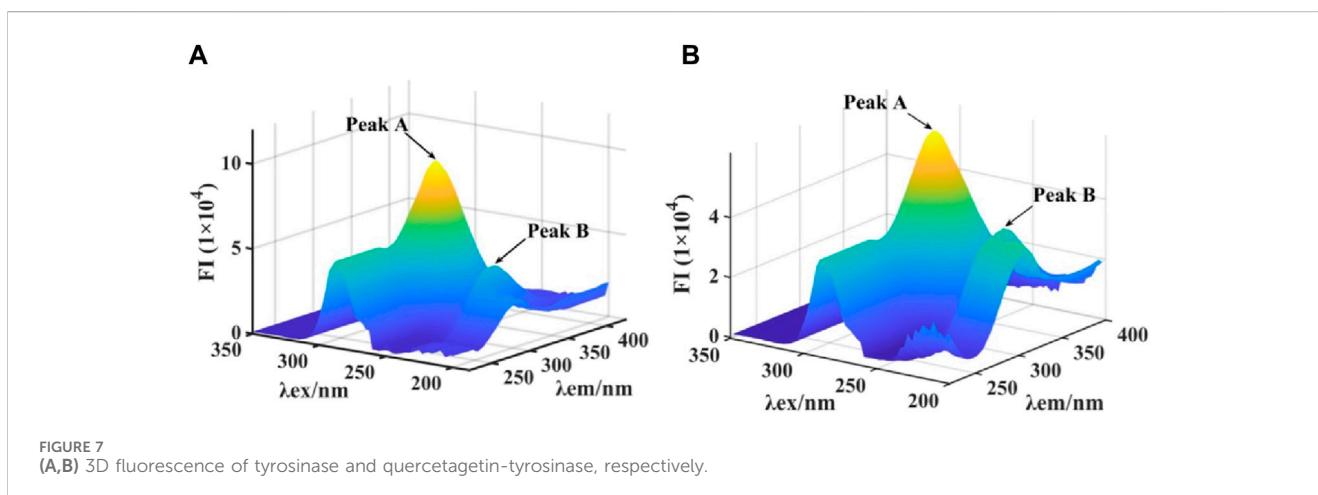
4.1 Tyrosinase activity assay

Tyrosinase inhibitory activity of quercetagenin was measured (Lu et al., 2023b). 10 μ L of quercetagenin in DMSO solutions was added into 140 μ L of tyrosinase in PBS solution and incubated for

TABLE 2 The secondary structure contents of tyrosinase with quercetagenin.

Molar ratio	α -Helix (%)	β -Sheet (%)	β -Turn (%)	Random coils (%)
1:0	27.0	23.1	15.6	44.7
1:1	22.2	27.6	16.9	41.1
1:2	19.2	31.4	17.9	38.8

Note: Molar ratio means [tyrosinase]: [quercetagenin].



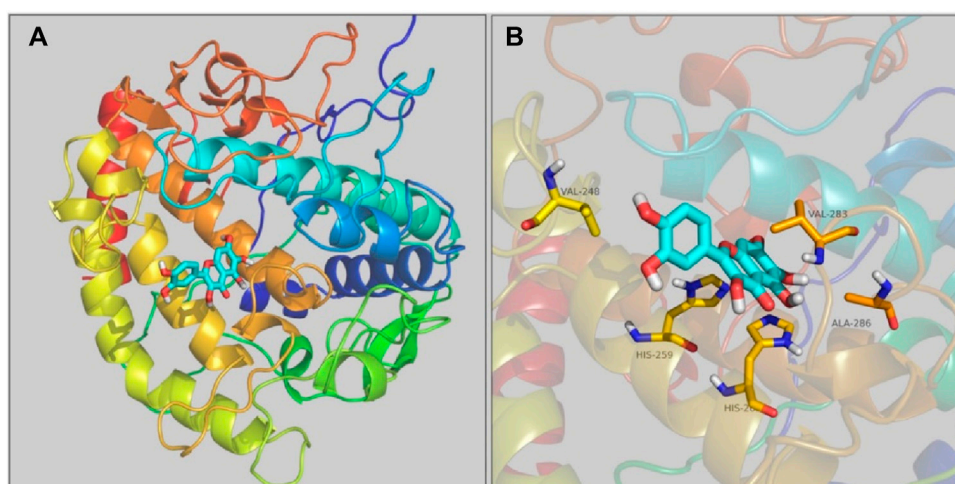


FIGURE 8
(A,B) The molecular docking of quercetagenin to tyrosinase.

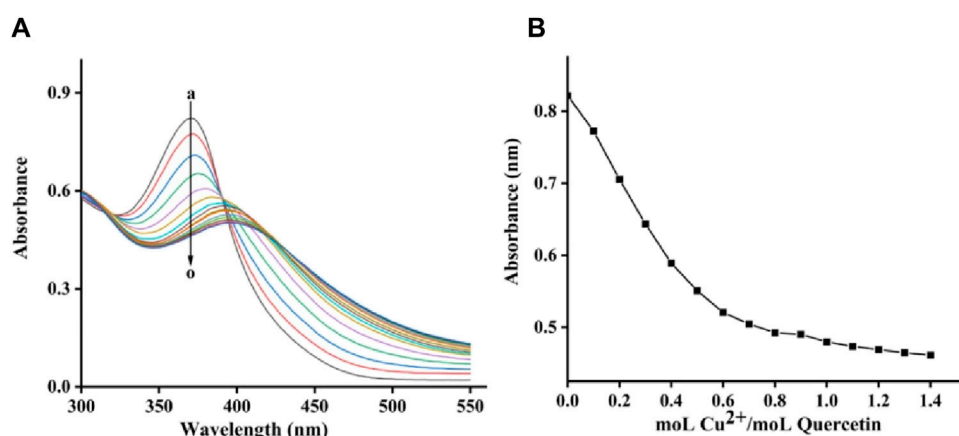


FIGURE 9
(A) UV spectra of quercetagenin-copper sulfate mixture; (B) UV peak of quercetagenin versus molar ratios of quercetagenin to copper sulfate.

10 min. 50 μL of L-dopa (in PBS) was added as the substrates. Then, the absorbance was measured at 475 nm on a microplate reader. Kojic acid was selected as the positive control. All experiments were tested in triplicate. Inhibition rate (%) = $[(\text{OD}_1 - \text{OD}_0) / \text{OD}_0] \times 100\%$. OD_0 and OD_1 were the absorbance of tyrosinase and tyrosinase-quercetagenin mixture.

4.2 Inhibitory kinetics

The reversibility of quercetagenin against tyrosinase was analyzed at different concentration of quercetagenin with tyrosinase or L-dopa. And the inhibitory kinetics of quercetagenin against tyrosinase was analyzed at different concentration of quercetagenin with tyrosinase or L-dopa. The inhibition constants K_i and K_{is} were obtained from the secondary curves of inhibitory kinetics (Xu et al., 2020; Deng et al., 2022; Lin et al., 2023).

4.3 Fluorescence quenching

To the 3.0 mL of tyrosinase solution, 1 μL of quercetagenin was added by titration method (Li et al., 2024a; Min et al., 2024). The mixture was measured fluorescence spectra at excitation of 280 nm. The experiments were conducted at temperatures of 295, 298, and 305 K, respectively. The Stern-Volmer equation and the van't Hoff equation were employed to obtained parameters.

4.4 CD spectra

To the 100 μL of tyrosinase solution, 1 μL of quercetagenin was added. CD spectra were measured at room temperature (Xiao et al., 2023; Li et al., 2024b). Tyrosinase solution without quercetagenin was also measured its CD spectra.

4.5 3D fluorescence spectra

To the 3.0 mL of tyrosinase solution, 1 μ L of quercetagenin was added. 3D fluorescence spectra were measured (Wu X. et al., 2023).

4.6 Copper-chelating activity

To the 2 mL of quercetagenin solution, 5 μ L of copper sulfate solution was gradually added. UV absorption spectra were detected. The molar ratios of quercetagenin to copper sulfate ranged from 10: 0 to 10: 14.

4.7 Molecular docking

Docking of quercetagenin with tyrosinase was simulated using the SYBYL software (Zhang et al., 2022; Feng et al., 2024). The tyrosinase crystal structure (PDB: 2Y9X) was optimized by removing water, adding hydrogen, and generation of active pocket. Quercetagenin structure was performed energy minimization. Thence, docking procedure was run in the default format.

Data availability statement

The raw data supporting the conclusion of this article will be made available by the authors, without undue reservation.

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Author contributions

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Conflict of interest

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