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Honokiol as an α -glucosidase inhibitor

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Honokiol, a naturally occurring compound from *Magnolia obovata Thunb.*, has many biological activities, but its anti- α -glucosidase activity is still unclear. Therefore, we determined its inhibitory effects against α -glucosidase. Activity assays showed that honokiol was a reversible mixed-type inhibitor of α -glucosidase, and its IC_{50} value was $317.11 \pm 12.86 \mu\text{M}$. Fluorescence results indicated that the binding of honokiol to α -glucosidase caused a reduction in α -glucosidase activity. 3D fluorescence and CD spectra results indicated that the binding of honokiol to α -glucosidase caused conformational change in α -glucosidase. Docking simulated the detailed interactions between honokiol and α -glucosidase, including hydrogen and hydrophobic bonds. All findings showed that honokiol could be used as a natural inhibitor to develop α -glucosidase agents.

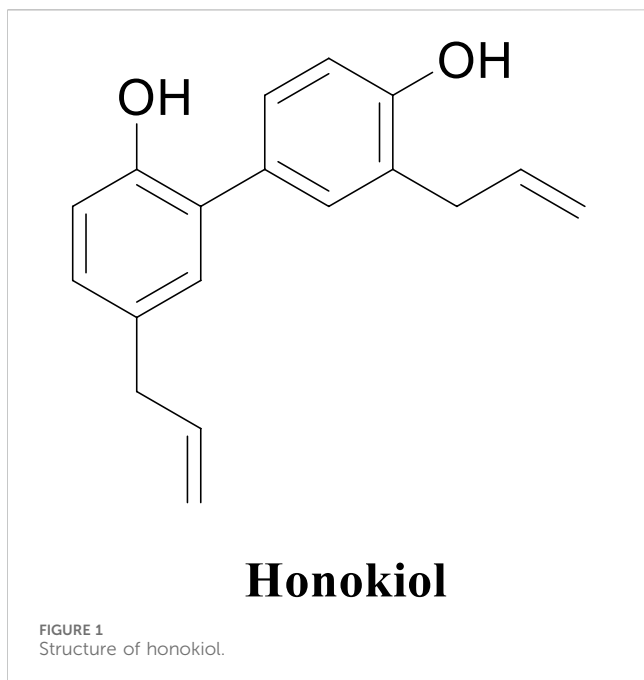
KEYWORDS

nature product, honokiol, α -glucosidase, inhibition mechanism, inhibitor

1 Introduction

Type 2 diabetes mellitus (T2DM) is a growing health concern with increased prevalence (Xu et al., 2020; ElSayed et al., 2023; Zhou et al., 2023). Now, T2DM has become a significant global health issue (Hu et al., 2024; Li et al., 2024). Epidemiological trends indicate that the prevalence of diabetes could reach an alarming 643 million individuals worldwide by 2030 (Song et al., 2022; Lin et al., 2023). The hallmark clinical feature of T2DM is elevated blood glucose levels, or hyperglycemia, which can result in a spectrum of debilitating complications (Jiang et al., 2020; Ding et al., 2021; Xing et al., 2021). Therefore, management of hyperglycemia is a critical aspect for T2DM patients (Hu et al., 2020).

One key characteristic of T2DM is postprandial hyperglycemia, which is intricately linked to the breakdown of carbohydrates (Hameed et al., 2019; Davies et al., 2022). α -Glucosidase, an enzyme present in enterocytes of the small intestine, facilitates the hydrolysis of glycosidic bonds to liberate glucose (Basri et al., 2023; Xiao et al., 2023). The suppression of α -glucosidase activity can thus delay carbohydrate digestion and absorption, leading to a reduction in the postprandial glucose spike (Basri et al., 2023; Wu et al., 2023). This rationale has made the inhibition of α -glucosidase a strategic target for therapeutic interventions to manage postprandial hyperglycemia (Khan et al., 2022; Zhang et al., 2022). Clinically, a number of α -glucosidase inhibitors, including acarbose and voglibose, have been employed to mitigate T2DM (Feng et al., 2024). However, chronic administration of these pharmaceuticals has its drawbacks, which urges people to seek safer and more effective α -glucosidase inhibitors (Lambrinoudaki et al., 2022; Min et al., 2024). Exploration of natural products as a repository for novel therapeutic agents has been a promising avenue (Zhang et al., 2021; Chen et al., 2022; Wang et al., 2022; Zhou et al., 2022). The active constituents



have demonstrated a diverse array of pharmacological effects, including anti-oxidant (Sun et al., 2020; Tao et al., 2022; Tang et al., 2023), anti-tumor (Chen et al., 2023; Liang et al., 2023; Song et al., 2023), anti-inflammatory (Sun et al., 2020; Wang et al., 2021; Wang et al., 2022), and anti-tissue damage (Shao et al., 2020; Qi et al., 2022; Ding et al., 2023) properties (Wang et al., 2020; Chen et al., 2022; Zang et al., 2022; He et al., 2023). Moreover, a notable advantage of natural products is their

generally lower toxicity profiles (Hao et al., 2022; Mao et al., 2022; Wang et al., 2023), which makes them a preferable source for development of α -glucosidase inhibitors.

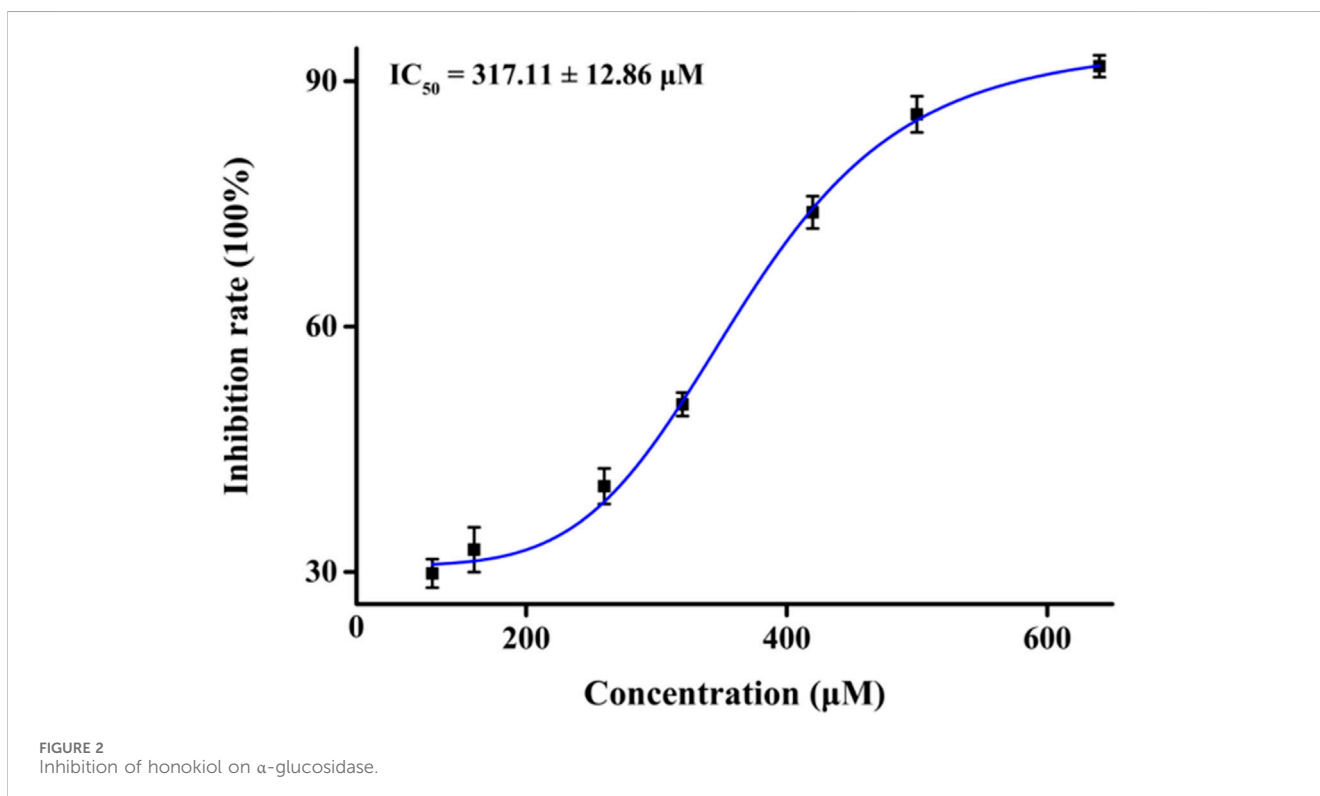
Honokiol (Figure 1), a naturally occurring compound from *Magnolia obovata* Thunb., has been recognized for its diverse medicinal properties (Zengin et al., 2017; Ma et al., 2023). As a bioactive neolignan, honokiol has demonstrated a range of activities, including anti-cancer, anti-inflammation, and anti-oxidant effects (He et al., 2015; Guo et al., 2021; Niu et al., 2021). In recent research studies, honokiol has garnered significant interest due to its ability to mitigate hyperglycemic conditions, enhance glucose uptake, and inhibit α -glucosidase activity (Bekircan et al., 2015; Pulvirenti et al., 2017; Ahmad et al., 2018). This shows the potential of honokiol as a natural α -glucosidase and hypoglycemic agent.

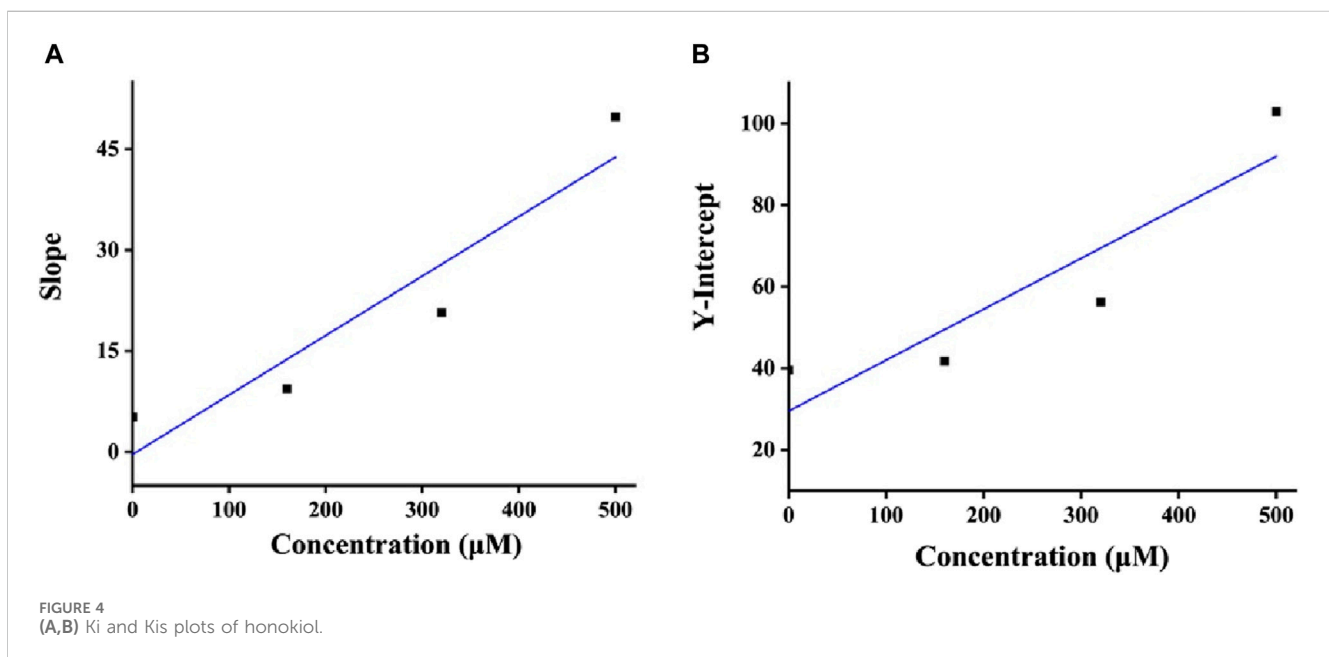
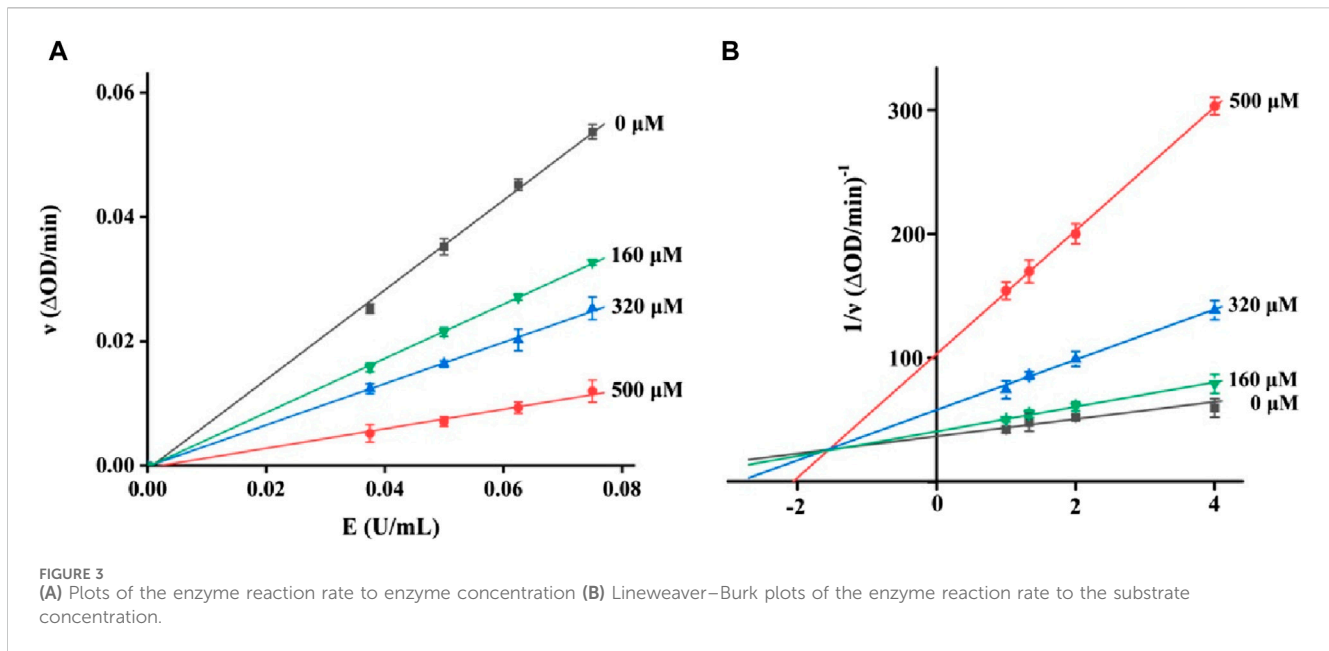
As far as we know, the detailed inhibitory effects of honokiol on α -glucosidase are still unclear. Hence, the biological activity of honokiol as an α -glucosidase inhibitor was investigated by spectroscopic methods and molecular docking.

2 Results and discussion

2.1 Inhibitory activity of honokiol on α -glucosidase

First, we assessed the inhibitory activity of honokiol on α -glucosidase, as shown in Figure 2. With an increase in the honokiol concentration, the inhibition rate gradually increased, and its IC_{50} value was calculated to be $317.11 \pm 12.86 \mu\text{M}$, which was lower than that of acarbose ($IC_{50} =$





$584.51 \pm 8.56 \mu M$). The potential inhibitory activity of honokiol on α -glucosidase might make it a natural hypoglycemic agent.

2.2 Kinetic study

It is very important to clarify the inhibition mode of inhibitors against enzymes for understanding the performance of inhibitors. Hence, the kinetics of honokiol on α -glucosidase were studied. In the plots of the enzyme reaction rate to the enzyme concentration under honokiol (Figure 3A), all lines passed the origin point. This indicated

honokiol as a reversible α -glucosidase inhibitor. In the Lineweaver–Burk plots of the enzyme reaction rate to substrate concentration under honokiol (Figure 3B), all lines intersected at the second quadrant. Their slope and Y-intercept were both changed with honokiol concentration. Therefore, it is evident that honokiol was a mixed-type inhibitor.

As a mixed-type inhibitor, honokiol was determined to have an inhibition constant. The fitting plot of slope and Y-intercept versus honokiol (Figures 4A,B) yielded K_i and K_{is} values of 16.03 and 285.22 μM , respectively. The lower K_i indicated that honokiol tended to bind to substrates.

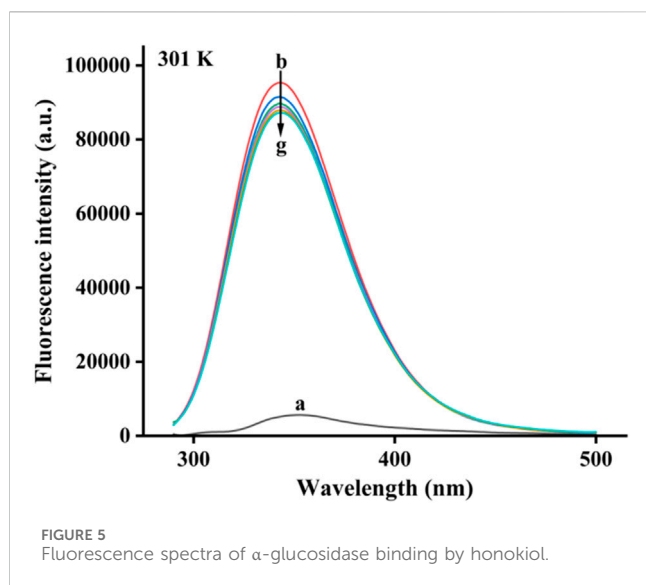


FIGURE 5
Fluorescence spectra of α -glucosidase binding by honokiol.

2.3 Fluorescence assay

Based on the fluorescence characteristics of α -glucosidase, the binding of honokiol to α -glucosidase was studied by fluorescence spectroscopy at 298K. In Figure 5, α -glucosidase presented fluorescence with a characteristic peak at 340 nm, while honokiol had very weak fluorescence at 340 nm. With continuous addition of honokiol, α -glucosidase fluorescence gradually decreased. This phenomenon indicated that there were binding interactions between honokiol and α -glucosidase, which could quench the endogenous fluorescence of α -glucosidase.

Subsequently, the binding of honokiol to α -glucosidase was further described by 3D fluorescence (Figures 6A,B). The 3D fluorescence spectra of α -glucosidase had two characteristic peaks due to intrinsic fluorophores and the backbone, which could be reduced by the addition of honokiol. This result was consistent with that of the fluorescence assay.

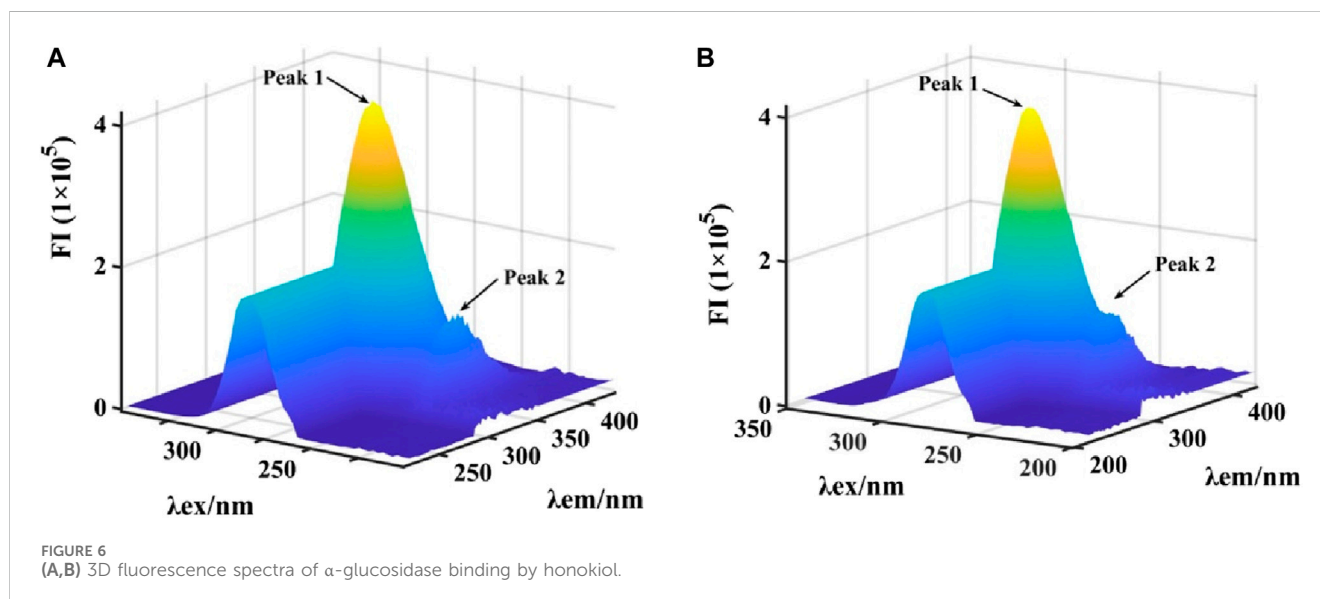


FIGURE 6
(A,B) 3D fluorescence spectra of α -glucosidase binding by honokiol.

2.4 CD spectra

CD spectra were investigated to evaluate the specific effects of honokiol on α -glucosidase structure. α -Glucosidase showed its own unique CD spectra at 210–222 nm (Figure 7). Honokiol treatment changed the CD spectra of α -glucosidase (Figure 7), which further indicated the binding of honokiol to α -glucosidase. The conformational changes in α -glucosidase were obtained from CD spectral data and showed that honokiol treatment resulted in changes in the α -glucosidase secondary structure content (Table 1). This might be the reason for the inhibition of honokiol on α -glucosidase.

2.5 Molecular docking

The docking interaction of honokiol with α -glucosidase was simulated. In a 3D view of docking (Figure 8A), honokiol was bound into the α -glucosidase active pocket, with a binding energy of -4.9 kcal/mol, presumably binding to amino acid residues in the pocket. Further analysis (Figure 8B) found that honokiol formed hydrogen bonds with GLU-276 (2.5 Å), ASP-349 (2.0 Å), and ASP214 (1.9 Å). Moreover, honokiol formed hydrophobic bonds with TYR-71, TYR-313, LEU-437, ARG-312, and PHE300. These main interactions between honokiol and α -glucosidase might be the reason for honokiol's inhibitory effect on α -glucosidase activity.

3 Materials and methods

3.1 α -Glucosidase inhibitory activity

α -Glucosidase was dissolved in PBS (pH 6.8), and honokiol was dissolved in DMSO. Honokiol solution and α -glucosidase solution were mixed and incubated for an appropriate time, and then a certain amount of substrate p-nitrophenyl- α -D-galactopyranoside

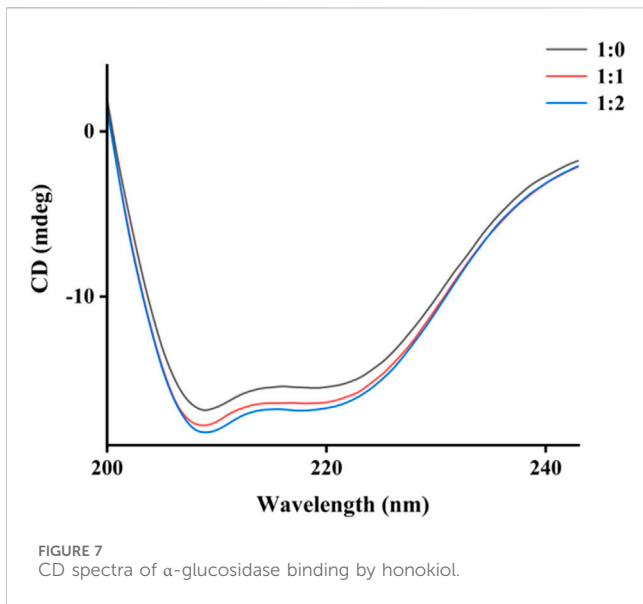


FIGURE 7
CD spectra of α -glucosidase binding by honokiol.

(pNPG) was added. Then, the absorbance of the solution at 405 nm was determined. Then, the α -glucosidase inhibitory effect of honokiol was obtained (Xu et al., 2020; Ali et al., 2023).

3.2 Inhibition kinetics

The test procedure for inhibition kinetics followed the same protocol as the α -glucosidase inhibition assay. For enzyme kinetics, the absorbance of a mixture with different concentrations of honokiol and α -glucosidase was recorded. For substrate kinetics, the absorbance of the mixture with different concentrations of honokiol and substrate was recorded (Kaur et al., 2021).

3.3 Fluorescence

Fluorescence measurements of α -glucosidase were conducted at an excitation wavelength of 280 nm (Wu et al., 2024). Then, honokiol was added step by step, and the corresponding fluorescence of the mixture was recorded.

3.4 3D fluorescence

3D fluorescence spectra of α -glucosidase with/without honokiol were recorded. The concentration of α -glucosidase was 0.1 mg/mL. Honokiol (0.25 μ M) was added to α -glucosidase to prepare their mixture.

TABLE 1 Conformational changes in α -glucosidase by honokiol.

[Enzyme]: [Honokiol]	α -Helix (%)	β -Sheet (%)	β -Turn (%)	Random coil (%)
1: 0	16.7	37.6	20.0	56.2
1: 1	17.0	36.5	19.9	59.4
1: 2	17.2	36.1	19.8	61.5

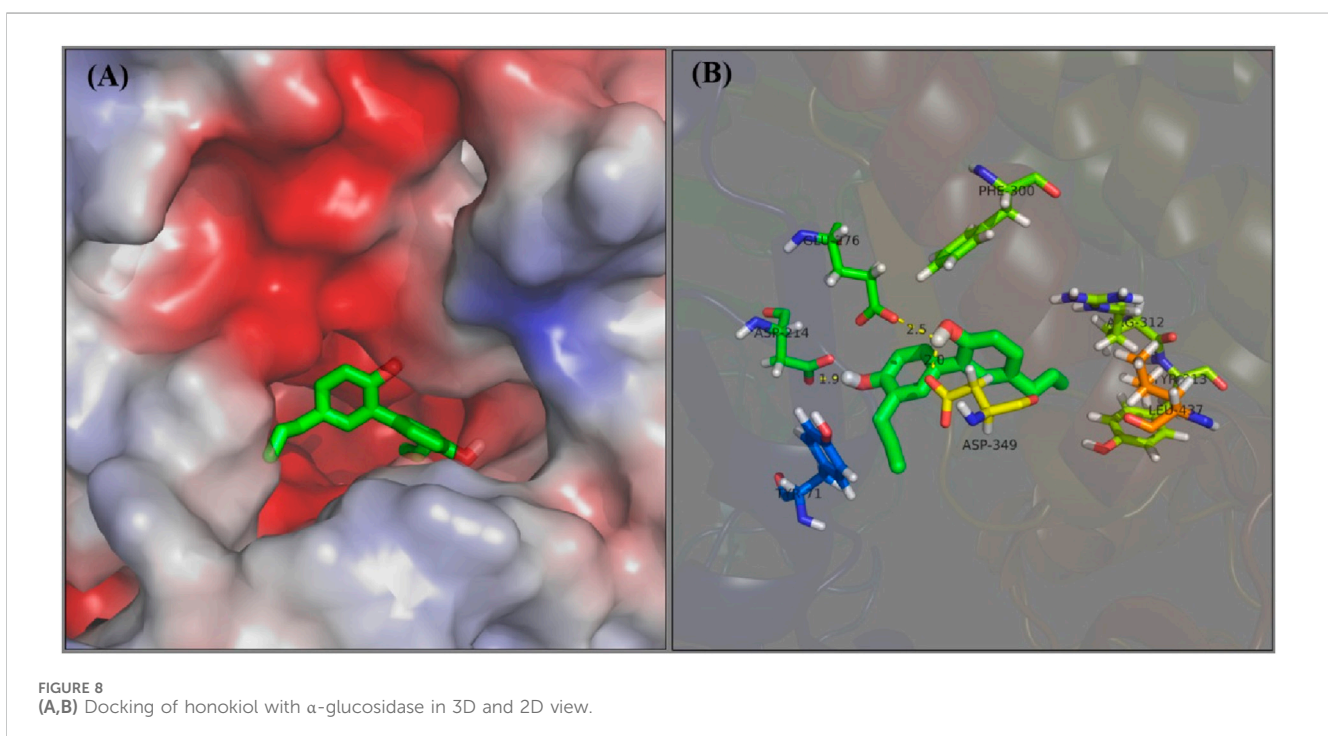


FIGURE 8
(A,B) Docking of honokiol with α -glucosidase in 3D and 2D view.

3.5 CD spectra

CD spectra of α -glucosidase with/without honokiol were also recorded. The concentration of α -glucosidase was 0.1 mg/mL. Honokiol was added to α -glucosidase to prepare their mixture. The data were analyzed using CDNN software (Li et al., 2024).

3.6 Molecular docking

The docking of honokiol with α -glucosidase was conducted using SYBYL (Deng et al., 2022; Patil et al., 2022). After being imported into the software, the honokiol structure was hydro-treated and charge-treated. Then, the homology model of α -glucosidase was also prepared by hydro-treating and charge-treating. Due to the absence of ligands in the protein, the active pocket of α -glucosidase was produced. Then, the docking of honokiol with α -glucosidase was performed in the default mode.

3.7 Statistical analysis

All data were presented as mean \pm SD. One-way ANOVA was performed to evaluate the differences between the groups (Zhao et al., 2017; Zhang et al., 2021; Zheng et al., 2021; Sheng et al., 2023). $p < 0.05$ was considered significant.

4 Conclusion

As a naturally occurring compound from *Magnolia obovata* Thunb., honokiol was ascertained for its anti- α -glucosidase activity and inhibition mechanism. So we designed experiments to clarify these properties. Activity assays showed that honokiol was a reversible mixed-type inhibitor of α -glucosidase, and its IC_{50} value was $317.11 \pm 12.86 \mu\text{M}$. Fluorescence, 3D fluorescence, and CD spectra investigations indicated that the binding of honokiol to α -glucosidase caused a reduction in α -glucosidase activity. Docking

simulated the detailed interactions between honokiol and α -glucosidase.

Data availability statement

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

Author contributions

HZ: investigation, project administration, writing—original draft, and writing—review and editing. XZ: investigation, project administration, and writing—original draft.

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Conflict of interest

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