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# [Extensive in](https://www.frontiersin.org/articles/10.3389/fimmu.2024.1404766/full)fluence of [microsporidian infection on](https://www.frontiersin.org/articles/10.3389/fimmu.2024.1404766/full) [sucrose solution consumption,](https://www.frontiersin.org/articles/10.3389/fimmu.2024.1404766/full) [antioxidant enzyme activity, cell](https://www.frontiersin.org/articles/10.3389/fimmu.2024.1404766/full) [structure, and lifespan of](https://www.frontiersin.org/articles/10.3389/fimmu.2024.1404766/full) [Asian honeybees](https://www.frontiersin.org/articles/10.3389/fimmu.2024.1404766/full)

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Apis cerana is the original host of Vairimorpha (Nosema) ceranae, a widespread fungal parasite that causes bee nosemosis, which severely threatens the health of bee colonies and the sustainable development of the apiculture industry. To evaluate the impact of V. ceranae infection on A. c. cerana workers, V. ceranae spores were purified and used to inoculate newly emerged workers to evaluate the effects of V. ceranae infection. This was followed by an in-depth investigation of V. ceranae spore load and host sucrose solution consumption. Activities of four major antioxidant enzymes (SOD, PPO, CAT, and GST) were determined. Paraffin sections of the host midgut tissue were prepared and subjected to microscopic observation. The survival rates of V. ceranae-inoculated and uninoculated workers were analyzed. The results showed that spore load gradually increased and peaked at 12 dpi. The consumption of workers in the V. ceranae-inoculated group was extremely significant higher ( $P < 0.0001$ ) than that of workers in the un-inoculated group. The results of antioxidant enzyme activity were suggestive of positive host defense via catalase (CAT) and glutathione-S-transferase (GST) in the middle stage of infection, as well as the negative fungal impact on superoxide dismutase (SOD) and polyphenol oxidase (PPO) at the whole stage of infection, reflecting the complex host-parasite interaction. Additionally, we observed a disruption in the structure of the host midgut epithelial cells. Moreover, the survival rate of workers in V. ceranaeinoculated groups was nearly always lower than that of workers in the uninoculated groups. These results demonstrate a consistent increase in spore load with the proliferation of V. ceranae, leading to persistent energetic stress and midgut epithelial cell structural damage to the host, ultimately resulting in a shortened lifespan for the host. Our findings enhance the current understanding of the interactions between A. cerana and V. ceranae as well as provide a solid basis for exploring the mechanisms underlying host response and V. ceranae infection.

#### KEYWORDS

honeybee, Apis cerana, Vairimorpha (Nosema) ceranae, host-parasite interaction, antioxidant enzyme

# 1 Introduction

Vairimorpha (Nosema) ceranae is a fungal parasite that causes nosemosis, a chronic disease that is frequently observed in bee colonies worldwide ([1](#page-10-0)). In 1996, Fries et al. ([2](#page-10-0)). first identified V. ceranae in A. cerana colonies, and established A. cerana as the original host of this parasite. Since its initial discovery in Europe in 2006, V. ceranae has rapidly spread to other parts of the world, North America, South America, Asia, Africa, and likely other regions as well, likely facilitated by the global trade of bees, bee products, and the movement of beekeeping equipment [\(3](#page-10-0)).

V. ceranae is associated with a decline in bee health, reduced colony productivity, and colony collapse disorders (CCD) [\(4,](#page-10-0) [5\)](#page-10-0). Over the past two decades, a substantial number of studies have been conducted to investigate the effect of V. ceranae infection on Apis mellifera, demonstrating that fungal invasion negatively influences various aspects of honeybee metabolism, behavior, physiology, and health ([6](#page-10-0)–[12\)](#page-10-0). For instance, studies have shown that the V. ceranae infection cause metabolic stress, upregulates genes encoding  $\alpha$ glucosidase and trehalose transport enzymes ([9\)](#page-10-0), while downregulating trehalase and Glucose-Methanol-Choline oxidoreductase genes [\(11](#page-10-0)). These alterations, as confirmed by proteomics, resulted in reduced levels of energy supply proteins ([8\)](#page-10-0). The behavioral consequences of V. ceranae infection in A. mellifera are also noteworthy, as infected bees often display increased hunger and consume more sucrose to compensate for their impaired metabolic state. However, this increased food intake does not fully alleviate the underlying nutritional and energy deficits ([6\)](#page-10-0), and accelerated lipid loss in bees suggests that lipids may be mobilized as an emergency energy source [\(12](#page-10-0)). V. ceranae infection can damage the midgut epithelial cell structure, inhibit apoptosis and immunosuppression, and shorten the lifespan of A. mellifera workers [\(7,](#page-10-0) [10\)](#page-10-0).

Through extensive interactions and long-term co-evolution, A. cerana and V. ceranae have evolved to coexist [\(13\)](#page-10-0). Previous studies have investigated the effects of V. ceranae infections on A. cerana ([14](#page-10-0)–[17\)](#page-10-0). Sinpoo et al. [\(14](#page-10-0)) found stronger immune responses and lower loads of Nosema apis and V. ceranae spores in workers of A. cerana compared to A. mellifera, suggesting a higher tolerance of A. cerana against microsporidian infection. Chantaphanwattana et al.

([18\)](#page-10-0) reported that V. ceranae proliferated more in A. mellifera, whereas A. cerana exhibited a stronger immune response and lower spore load, indicating a superior defense mechanism of A. cerana at the individual level. By contrasting the similarities and differences in V. ceranae infection of A. cerana and A. mellifera, a deeper understanding of the complex interactions between V. ceranae and bee hosts, as well as the factors responsible for coaptation between A. cerana and V. ceranae will be gained. In addition, novel and valuable insights from the aforementioned comparisons could guide the creation of targeted strategies to alleviate the adverse effects of fungal parasites on different bee species.

A. c. cerana is a bee species that is found in China and is widely used in beekeeping practice in several Asian countries [\(19\)](#page-10-0). Previously, our team identified the miRNA response and miRNA-regulated network of A. c. cerana workers following V. ceranae invasion [\(20](#page-10-0)), and performed a comprehensive investigation of the profiles of highly expressed genes in both the host and microsporidian [\(21](#page-10-0), [22](#page-10-0)). Recently, our group investigated the immune response of A. c. cerana workers to V. ceranae infection using a transcriptomic investigation and revealed that different cellular and humoral immune responses were used by A. c. cerana and A. mellifera ligustica workers to defend themselves against infection by V. Ceranae ([23](#page-10-0)). Moreover, this study provides an opportunity for a deeper comparison of the effect of the same fungal parasite (V. ceranae) on two different bee species (A. cerana and A. mellifera), which will enrich our understanding of bee nosemosis and facilitate the development of biological drugs for nosemosis control.

Many aspects of the effect of V. ceranae on A. cerana are yet to be explored. Pathogenic infections in insects result in the generation of reactive oxygen species (ROS) that can kill and eliminate pathogens ([24](#page-10-0)). However, excessive production of ROS in vivo induces oxidative stress, predominantly damaging various biological macromolecules, and ultimately resulting in cell death [\(25\)](#page-10-0). Antioxidant enzymes reduce the oxidative response to balance ROS production and maintain homeostasis ([26\)](#page-10-0). Superoxide dismutase (SOD) catalyzes the disproportionation of superoxide to produce hydrogen peroxide and molecular oxygen, whereas catalase (CAT) participates in the decomposition of hydrogen peroxide into water and oxygen [\(27](#page-10-0)–[29](#page-10-0)). Glutathione-S-transferase (GST) has been shown to exert function in metabolizing lipid peroxides ([27,](#page-10-0) [30](#page-10-0)). Polyphenol oxidase (PPO) is an oxidoreductase that uses molecular oxygen to catalyze the oxidation of a wide range of phenolic compounds [\(31](#page-10-0)). Insects appear to use PPO in immune responses such as melanization to physically limit the spread of microbes ([32\)](#page-10-0). Whether V. ceranae infection affects the activities of major antioxidant enzymes in A. cerana remains largely unknown.

Currently, little is known regarding the effect of V. ceranae infection on A. cerana, hindering the understanding of the mechanisms underlying the interaction between V. ceranae and A. cerana and the development of novel strategies for the control of bee nosemosis. Here, to evaluate the impact of V. ceranae infection on A. c. cerana workers, V. ceranae spores were purified and used to inoculate newly emerged workers to evaluate the effects of V. ceranae infection. This was followed by an in-depth investigation of V. ceranae spore load and host sucrose solution consumption. Activities of four major antioxidant enzymes (SOD, PPO, CAT, and GST) were determined. Paraffin sections of the host midgut tissue were prepared and subjected to microscopic observation. The survival rates of V. ceranae-inoculated and uninoculated workers were analyzed. The findings from this study provide valuable experimental evidence for V. ceranae infection of A. c. cerana workers and contribute to the dissection of the host response, microsporidian infection, and host–microsporidian interaction.

# 2 Materials and methods

#### 2.1 Spore extraction, artificial inoculation, and honeybee preparation for bioassay experiments

A. c. cerana workers were obtained from colonies in the teaching apiary of the College of Bee Science and Biomedicine, Fujian Agricultural and Forestry University, in October 2020. There were no specific bands amplified from V. apis, V. ceranae, or several viruses, such as SBV (Sacbrood Virus), DWV (Deformed Wing Virus), IAPV (Israeli Acute Paralysis Virus), BQCV (Black Queen Cell Virus), KBV (Kashmir Bee Virus), and CBPV (Chronic Bee Paralysis Virus) based on reverse transcription-polymerase chain reaction (RT-PCR) with specific corresponding primers, that eliminated the interference of pathogens on the samples.

V. ceranae-infected A. c. cerana workers were obtained from an apiary in Minhou, Fuzhou, China. V. ceranae spores were prepared using the discontinuous density gradient Percoll method, as described by Chen et al. [\(33\)](#page-10-0). Briefly, (1) a total of 200 foragers were collected from an infected colony, the midgut tissues were dissected and then homogenized in distilled water, followed by filtration through four layers of sterile gauze and three cycles of centrifugation at 6000  $\times$  g for 5 min; (2) the supernatant was discarded as the spores remained in the sediment, and the resuspended pellet was then purified on a discontinuous Percoll gradient (Solarbio) consisting of 200 µL each of 25%, 50%, 75%, and 100% Percoll solution; (3) the spore suspension was overlaid onto the gradient and centrifuged at  $18000 \times g$  for 90 min at 4°C; (4) the spore pellet was isolated using a sterile syringe and then centrifuged again on a discontinuous Percoll gradient. Purified spores were examined by PCR using previously reported primers ([34](#page-10-0)) and then

preserved in our laboratory and at the China Common Microbial Strain Preservation and Management Center (Preservation number CGMCC: No. 28110). The spore concentration was determined using a CL kurt counter for the inoculation experiments.

Before artificial inoculation, the prepared microsporidian spores were observed microscopically and examined using PCR with specific primers for V. ceranae (F: 5'-CGGATAAAAGAGTCCGTTACC-3'; R: 5′-TGAGCAGGGTTCTAGGGAT-3′) [\(17](#page-10-0), [20\)](#page-10-0) and V. apis (F: 5′- CCATTGCCGGATAAGAGAGT-3′; R: 5′-CACGCATTGCTGC ATCATTGAC-3′), as previously described by Chen et al. [\(34](#page-10-0)). The inoculation and rearing of A. c. cerana workers were performed according to a previously established protocol [\(20\)](#page-10-0). At 24 h after emergence, members of the treatment group ( $n = 35$ ) were each immobilized and fed 5  $\mu$ L of 50% (w/v) sucrose solution containing  $1 \times 10^6$  V. ceranae spores by using a pipette, whereas members of the control group ( $n = 35$ ) were fed 5 µL of sucrose solution without V. ceranae spores. The treatment (or control) group contained six plastic cages. All workers were reared in an incubator at  $34 \pm 0.5^{\circ}$ C and 60– 70% RH (relative humidity). After 24 h, the honeybees were fed a feeder containing 4 mL of 50% (w/v) sucrose solution, and the feeders were replaced daily throughout the experiment. Each cage was carefully checked, and dead honeybees were removed daily. Three biological replicates were used in each experiment.

The prepared workers from V. ceranae- and uninoculated groups underwent spore load measurement ( $n = 3$ ), antioxidant enzyme activity examination  $(n = 3)$ , hematoxylin and eosin staining (HE) of paraffin sections ( $n = 3$ ).

## 2.2 Measurement of spore load in midguts of the workers

During the V. ceranae infection process, 1-12 days post inoculation (dpi), the spores in the midguts of the workers were extracted every 24 h based on the standard method described by Fries et al. ([35\)](#page-10-0) and Naree et al. [\(16](#page-10-0)), followed by calculation under an optical microscope (CSOIF, Shanghai, China), according to the procedure described by Cantwell ([36\)](#page-10-0). In brief, (1) after inoculation with V. ceranae, a single worker in the treatment group was collected from the cage daily, and their midgut tissues were then dissected using a clean ophthalmic forced and transferred into a sterile 1.5-mL Eppendorf (EP) tube; (2) 100 µL of sterile water was added to the EP tube, and then full grinding with an automatic high throughput tissue grinder (Meibi, Zhejiang, China) was performed; (3) using a micropipette, 20 µL of the aforementioned solution was carefully observed with a hemocytometer plate (Qiujing, Shanghai, China) under an optical microscope (CSOIF, Shanghai, China), after which spores were counted.

#### 2.3 Detection of host average sucrose solution consumption

According to the method described in Section 2.1, separate experimental workers were prepared for the purpose of statistical analysis of sucrose solution consumption and survival rates. The daily average consumption of sucrose solution by each worker in

<span id="page-3-0"></span>the V. ceranae- and uninoculated groups was analyzed using a previously described method ([37](#page-10-0), [38\)](#page-10-0). The feeder containing 4 mL of sucrose solution was weighed and the weight was recorded as A. To avoid errors due to repeated changes in feeders, the feeder was weighed every 24 h thereafter and the weight was recorded as B. The sucrose solution in the feeders was replaced daily. The numbers of survivors in both V. ceranae- and un-inoculated groups were counted. The average sucrose solution consumption per worker per day was calculated according to the following formula: (B-A)/ number of survivors in the V. ceranae-inoculated group (or uninoculated group). Each biological replicate was performed three times,  $n = 35$ .

## 2.4 Examination of the antioxidant enzyme activity

At 1–12 dpi, three workers were randomly selected from V. ceranae- and un-inoculated groups, the midgut tissues were carefully dissected and then added into a sterile EP tube containing 180 µL of saline followed by fully ground with a high-throughput tissue grinder (MEIBI, Hangzhou, China), respectively. The grinding fluid was centrifugation at 5000  $\times$  g for 5 min, and the supernatant was then transferred to a sterile EP tube, stored at −80°C. Following the corresponding instructions, the activities of SOD and PPO were examined using the Insect SOD ELISA Kit (MLBIO, Shanghai, China) and Insect PPO ELISA Kit (MLBIO, Shanghai, China), respectively, whereas those of CAT and GST were determined using the Insect CAT ELISA Kit (MLBIO, Shanghai, China) and Insect GST ELISA Kit (MLBIO, Shanghai, China), respectively. The experiment included three biological replicas. Specific antioxidant enzyme activity was expressed as units of enzyme activity per milligram of protein.

#### 2.5 Paraffin sectioning, HE staining, and microscopic observation of midgut tissues of workers

At 7, 8, 9, and 10 dpi, three workers were randomly selected from V. ceranae- and uninoculated groups, and the midgut tissues were carefully dissected. Next, according to the method described by Xing et al. [\(23\)](#page-10-0), midgut tissue from one of the workers was fixed with 4% paraformaldehyde (PFA). By using an embedding center (Junjie, Wuhan, China) and a microtome (Leica, Nussloch, Germany), paraffin sections of midgut tissues were stained with HE staining by Shanghai Sangon Biological Engineering Co. Ltd., and then detected under an optical microscope with digital camera (SOPTOP, Shanghai, China).

#### 2.6 Survival rate of workers

Using the same experimental group as in Section 2.3, before replacing the sucrose solution every 24 hours, the survival status of workers in both the V. ceranae-inoculated group and the uninoculated group was recorded, and dead ones were removed. This recording was continued until 20 dpi. Subsequently, the daily survival rates for the both groups were calculated on basis of the formula: (number of living worker bees in the treatment or control group)/105.

## 2.7 Statistics

Statistical analysis of the data was performed using SPSS software. 21 (IBM, Armonk, NY, USA) and GraphPad Prism 6.0 software (GraphPad, San Diego, CA, USA). To analyze significant differences among multiple groups of data, either One-way ANOVA (followed by Tukey's post-hoc test), Two-way ANOVA (followed by Bonferroni's post-hoc test), multiple t tests was performed on the spore load and average sucrose solution consumption, and activity of SOD, PPO, GST and CAT using the SPSS software. Data are presented as the mean ± standard deviation (SD). Tukey's test and the significant difference letter marker method were used to compare the number of spore loads between the 2 d. The log-rank (Mantel–Cox) test was used to analyze the host survival rate.

# 3 Results

#### 3.1 Dynamics of V. ceranae spore load in the midguts of A. c. cerana workers

After discontinuous Percoll gradient centrifugation, oval spores with high refraction were observed using an optical microscope (OM) (Figure 1A). In addition, the expected fragment (76 bp) could



<span id="page-4-0"></span>be amplified from the purified spores with specific primers for V. ceranae but could not be amplified from sterile water and spores with *V. apis* specific primers [\(Figure 1B](#page-3-0)). These results confirmed that the prepared spores were V. ceranae spores.

The spore counting results suggested that the spore load of V. ceranae in the midguts of workers presented an overall elevation trend (Figure 2). In detail, as compared with initial spore load (1  $\times$ 10<sup>6</sup>), the *V. ceranae* spore load in the midguts of the workers continuously increased from 1 dpi  $(3.5 \pm 0.87 \times 10^6)$  to 4 dpi (52.83)  $\pm$  2.75  $\times$  10<sup>6</sup>), from 5 dpi (72  $\pm$  13.08  $\times$  10<sup>6</sup>) to 8 dpi (48.53  $\pm$  11.533  $\times$  10<sup>7</sup>), and from 9 dpi (42.68  $\pm$  9.42  $\times$  10<sup>7</sup>) to 12 dpi (10.51  $\pm$  0.83  $\times$ 10<sup>8</sup>) (Figure 2); however, the spore load was decreased from 8 to 9 dpi. In addition, the quantity of spores reached the peak at 12 dpi (Figure 2).

## 3.2 Influence of V. ceranae infection on average sucrose solution consumption of A. c. cerana workers

The average sucrose consumption of V. ceranae-inoculated and un-inoculated workers was  $0.0308 \pm 0.0080$  g/d and  $0.0371 \pm 0.0107$ g/d, respectively. The consumption of workers in the V. ceranaeinoculated group was extremely significant higher (P<0.0001) than that of workers in the un-inoculated group, significant difference in the average sucrose solution consumption between the two groups was observed at 5 ( $P = 0.0244$ ), 8 dpi ( $P = 0.0005$ ), 10 dpi ( $P =$ 0.0443), 11 dpi (P = 0.0497), 12 dpi (P = 0.0243), 13 dpi (P = 0.0286) and 20 dpi ( $P = 0.0018$ ) ([Figure 3](#page-5-0)).

## 3.3 Impact of V. ceranae infection on SOD and PPO activities in A. c. cerana midguts of the workers

Compared to that in the uninoculated group, the SOD activity in the *V. ceranae*-inoculated group was decreased at  $2 (P = 0.2736) - 10$  dpi  $(P = 0.0399)$ , with a significant difference observed at 5 dpi  $(P = 0.0399)$ 0.0097), 6 dpi (P = 0.0434), 9 dpi (P = 0.0015), and 10 dpi [\(Figure 4A](#page-5-0)). The PPO activity was consistently lower in the V. ceranae-inoculated group in comparison to that in the un-inoculated group at  $2(P =$ 0.2249)–12 ( $P = 0.5391$ ) dpi, although no statistically significant difference was observed between these two groups [\(Figure 4B](#page-5-0)).

## 3.4 Impact of V. ceranae infection on CAT and GST activities in the midguts of A. c. cerana workers

CAT activity was reduced in the V. ceranae-inoculated group in comparison with that in the un-inoculated group at  $1 (P = 0.9209)$ – 3 dpi ( $P = 0.0981$ ), [\(Figure 5A\)](#page-6-0); conversely, then was elevated in the *V. ceranae*-inoculated group at 4 ( $P = 0.1596$ )–7 ( $P = 0.1954$ ) dpi; CAT activity was again downregulated in the host midgut following *V. ceranae* inoculation at 8 ( $P = 0.3183$ )–12 dpi ( $P = 0.4700$ ), with a significant difference observed at 10 dpi ( $P = 0.0478$ ) ([Figure 5A\)](#page-6-0). Additionally, GST activity was up-regulated at  $5 (P = 0.5171) - 6$  dpi  $(P = 0.3692)$  and then reduced at 7  $(P = 0.8196)$ –10 dpi  $(P = 0.0070)$ , followed by an increase at 11 dpi ( $P = 0.1391$ ), with a significant difference observed at 9 dpi ( $P = 0.0229$ ), and 10 dpi ( $P =$ 0.0070) [\(Figure 5B](#page-6-0)).



<span id="page-5-0"></span>

## 3.5 Influence of V. ceranae infection on midgut epithelial cell structure of A. c. cerana workers

Microscopic observation of paraffin sections revealed darkly stained microsporidia in the midgut epithelial cells of V. ceranaeinoculated workers at 7–10 dpi [\(Figures 6C,](#page-7-0) D, G, H, K, L O, P), whereas no V. ceranae was detected in the midgut epithelial cells of uninoculated workers [\(Figures 6A,](#page-7-0) B, E, F, I, J, M, N). The number of V. ceranae spores in host cells gradually increased with increasing infection time ([Figures 6C,](#page-7-0) D, G, H, K, L O, P).

In addition, the columnar and goblet cells were closely arranged, the epithelial cells were orderly arranged, the periesophageal membrane was clearly visible [\(Figures 6A,](#page-7-0) B, E, F, I, J, M, N), the outlines of columnar and goblet cells of V. ceranae-inoculated workers were arranged loosely and became unclear, and the periesophageal membrane was invisible ([Figures 6C,](#page-7-0) D, G, H, K, L [O,](#page-7-0) P). Moreover, as the infection progressed, the epithelial cells fell off the midgut wall and entered the intestinal lumen, together with the nuclei, cytoplasm, and various types of V. ceranae spores [\(Figure 6\)](#page-7-0).

#### 3.6 Influence of V. ceranae infection on lifespan of A. c. cerana workers

The survival rate statistics indicated that the survival rates of workers in both the V. ceranae-inoculated and uninoculated groups at 1–5 dpi were high, and there was little difference between the groups [\(Figure 7\)](#page-8-0). Survival rates in both groups began to decrease at 5 dpi ([Figure 7\)](#page-8-0). At 5–11 dpi, the survival rate of workers in the V. ceranae-inoculated group was lower than that in the un-inoculated group, whereas the survival rate of workers in the V. ceranaeinoculated group was significantly lower than that in the uninoculated group at 11–20 dpi [\(Figure 7\)](#page-8-0).



Detection of SOD (A) and PPO (B) activities in midguts of V. ceranae- and un-inoculated workers ( $n=3$ ). Data were showed as mean  $\pm$  SD, that analyzed using the Two-way ANOVA (Bonferroni's post-hoc test), multiple t tests; ns indicates nonsignificance, \* indicates  $P < 0.05$ , \*\* indicates  $P < 0.01$ .

<span id="page-6-0"></span>

# 4 Discussion

There is an ongoing debate and uncertainty surrounding the precise lifecycle duration of V. ceranae across different hosts. In A. mellifera, all intracellular stages of the life cycle of V. ceranae were observed at 3 dpi inside the worker's epithelial cells using transmission electron microscopy, suggesting that the time for V. ceranae to complete the developmental cycle was 3 d ([39](#page-10-0)). Based on other lines heterologous lepidopteran cell line IPL-LD-65Y in vitro, Gisder et al. reported that the life cycle of N. apis, the sister species of V. ceranae, is approximately 4 d ([40](#page-10-0)). Although A. cerana is the original host of V. ceranae, its life cycle of V. ceranae in the midgut epithelium cells of A. cerana is unknown. Here, we observed a continuous increase in spore load from 1–12 dpi ([Figure 2\)](#page-4-0), indicative of sustained proliferation of V. ceranae within host cells. Additionally, the number of spores increased 10-fold every 4 d [\(Figure 2\)](#page-4-0), suggesting that the life cycle of V. ceranae in the midgut epithelial cells of A. cerana workers was approximately 4 d, which differs from reports on A. mellifera. Intriguingly, we observed the presence of spores  $(3.5 \pm 0.87 \times 10^6)$  as early as 1 dpi. Similar phenomenon was observed in previous studies, namely that by 3 h post inoculation, mature spores can be observed in the ventricular lumen of infected A. mellifera workers [\(39\)](#page-10-0). We hypothesized that the detected V. ceranae spores at 1 dpi contained spores that did not infect the bee host cells or empty spores. However, further studies are needed to verify this hypothesis.

Nosema spp. have lost mitochondria during evolution, and thus must acquire energetic substrates directly from hosts for maintenance, growth, and reproduction ([41,](#page-10-0) [42](#page-10-0)). Intestinal lesions caused by V. ceranae proliferation may decrease the absorptive capacity of honeybees and cause starvation symptoms, such as impoverishment of hypopharyngeal protein secretions ([8\)](#page-10-0). Chronic nutrient and energy stress suppress the innate immune system [\(43\)](#page-10-0). After V. ceranae infection, workers of A. mellifera consumed a significantly higher amount of sucrose solution over the 24-h period tested ([6\)](#page-10-0). Here, we observed that the sucrose solution consumption of V. ceranae-inoculated workers was nearly always higher than that of uninoculated workers from 1–20 dpi, which indicated that the

workers attempted to compensate for the imposed energy stress by consuming more sucrose solution. This is consistent with the results of a previous study on A. mellifera workers infected with V. ceranae ([44](#page-10-0)). In summary, these results demonstrate that V. ceranae infection results in energy stress in both A. mellifera and A. cerana workers. In addition, the chronic stress caused by V. ceranae infection is likely to negatively influence foraging behavior, nutritional balance, immune response, and other important aspects of A. cerana workers, thus deserving more attention.

Insects maintain homeostasis in the body by neutralizing ROS and xenobiotics through the activation of various metabolic pathways that help protect cellular components from oxidative stress ([26](#page-10-0), [45](#page-10-0)). The antioxidant enzyme system, comprising SOD, PPO, CAT, and GST, is a critical physiological system for resisting oxidative stress in insects ([46](#page-11-0), [47\)](#page-11-0). After infection with pathogens, the insect immune system stimulates the production of ROS, including superoxide radicals, to kill the invaders [\(24](#page-10-0)). CAT efficiently breaks down toxic hydrogen peroxide into harmless water and oxygen molecules ([48\)](#page-11-0). GST protects cells from oxidative stress by conjugating glutathione to ROS and other electrophilic compounds [\(49\)](#page-11-0). It has been previously reported that GST is linked to oxidative stress response in the midgut of Drosophila melanogaster larvae [\(50\)](#page-11-0). In the present study, CAT activity in the V. ceranae-inoculated group increased significantly at 5, 6, and 7 dpi (Figure 5A), whereas GST activity was significantly elevated at 6 dpi (Figure 5B). We inferred that the host enhanced the levels of CAT and GST activities in response to V. ceranaecaused oxidative stress in the middle stage of the infection process, thereby maintaining homeostasis of the internal environment. SOD catalyzes the dismutation of superoxide radicals into hydrogen peroxide and oxygen, thereby regulating ROS levels and preventing oxidative damage to cells [\(51](#page-11-0)). A significant decline in the activities of both SOD and GST was detected at 10 dpi, and CAT activity was reduced at both 9 and 10 dpi ([Figures 4A,](#page-5-0) 5). These results suggest that these three critical antioxidant enzymes are inhibited in the host midgut. This phenomenon may be attributed to the escalating spore load in the late infection stage, leading to a

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#### FIGURE 6

Microscopic observation of paraffin midgut sections of un-inoculated and V. ceranae-inoculated A. c. cerana workers. (A, B) Midgut tissue of uninoculated worker at 7 dpi without spores; (C, D) Midgut tissue of inoculated worker at 7 dpi with V, ceranae spores; (E, F) Midgut tissue of un-inoculated worker at 8 dpi without spores; (G, H) Midgut tissue of inoculated worker at 8 dpi with V. ceranae spores; (I, J) Midgut tissue of uninoculated worker at 9 dpi without spores; (K, L) Midgut tissue of inoculated worker at 9 dpi with V. ceranae spores; (M, N) Midgut tissue of uninoculated worker at 10 dpi without spores; (O, P) Midgut tissue of inoculated worker at 10 dpi with V. ceranae spores. (A, C, E, G, I, K, M, O) were microscopic fields under 200x amplification, whereas (B, D, F, H, J, N, P) were microscopic fields under 400x amplification. Black dashed boxes show the selected region for observation under 200× amplification.

prolonged intensification of V. ceranae stress in the host and consequently suppressing the activities of SOD, CAT, and GST. PPO is involved in the melanization process, a common immune response in insects that involves the production of dark and pigmented structures that can trap and kill invading pathogens or parasites. In addition to its role in melanization, it simultaneously induces cellular and humoral immunity [\(52\)](#page-11-0). Roberts and Hughes ([53](#page-11-0)) found a negative correlation between PPO levels in the midguts

of A. mellifera workers and V. ceranae infection intensity. Intriguingly, although the level of PPO activity in the V. ceranaeinoculated group was always lower than that in the uninoculated group, there was no significant difference between the groups ([Figure 4B\)](#page-5-0), indicating that PPO activity in the host midguts was not affected by fungal invasion. It is inferred that, as the host of origin of V. ceranae, A. cerana workers potentially evolved a strategy to counteract the negative influence of microsporidians

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on PPO activity. Further studies are required to explore these underlying mechanisms. Insects exhibit dynamic changes in GST and CAT in response to pathogen infection. For instance, in the larvae of Holotrichia parallela, following joint infection of entomopathogenic nematode and Bacillus thuringiensis, there was an initial increase in GST and CAT activity during the early stage, followed by a decrease in activity in the later stage ([54](#page-11-0)). In this study, the contrasting alterations observed in GST and CAT levels between the middle and late stages of V. ceranae infection could potentially be ascribed to the insects' initial augmentation of antioxidant enzyme activity as a mechanism of defense against the fungal invasion. Conversely, a decline in enzyme activity may occur in the later stage as a result of energy depletion or other physiological adjustments.

In a previous study, Higes et al. [\(39](#page-10-0)) detected that, 7 d after V. ceranae infection, the majority of epithelial cells showed evidence of

degeneration, such as the presence of vacuoles in the cytoplasm or a brightly stained nucleus. Few unaltered cells were also observed, and either at the tips or at the bottom of the epithelial folds, there were cells containing V. ceranae intracellular stages. Here, we observed that the number of V. ceranae in the A. cerana worker midgut epithelial cells increased from 7 to 10 dpi [\(Figure 6](#page-7-0)), which is consistent with the findings of Higes et al. ([39\)](#page-10-0) Collectively, these results offer a solid basis for further investigations of V. ceranae infection in A. cerana workers, such as functional studies on fungal virulence factor-associated genes and the development of an RNAi-based control strategy.

V. ceranae infection damaged the structure of midgut epithelial cells in A. mellifera workers ([7](#page-10-0)). Based on microscopic observations by optical microscopy and transmission electron microscopy, Fries et al. ([2\)](#page-10-0) first reported that V. ceranaeinfected ventriculus cells of A. cerana workers were shed into or burst open into the gut lumen where mature spores were released,



and described the size of the spores. Darkly stained microsporidia were clearly detected in the midgut epithelial cells of V. ceranaeinoculated workers at 7–10 dpi [\(Figure 6](#page-7-0)). Additionally, the boundaries of the midgut epithelial cells in V. ceranaeinoculated workers were blurred, and the nuclei almost disappeared, with nucleic acid substances dispersed around the cytoplasm [\(Figure 6\)](#page-7-0). These results suggest that V. ceranae infection also causes serious damage to the structure of the midgut epithelial cells of A. c. cerana workers. Collectively, the findings from previous studies ([2](#page-10-0)) and this study offer experimental evidence for the damage caused by V. ceranae infection to the structure of A. c. cerana worker midgut epithelial cells, laying the foundation for further investigation of the pathogenesis of bee nosemosis.

Paris et al. ([55](#page-11-0)) assessed the survival rate and eating behavior of A. mellifera workers infected with V. ceranae for 1–22 d, and the results showed that the survival rate of infected workers was significantly lower than that of uninfected workers. We previously observed that the mortality rate of A. mellifera workers inoculated with V. ceranae spores gradually increased over time, and that the mortality rate of V. ceranae-inoculated workers was significantly higher than that of uninoculated workers at 7 and 10 dpi [\(33](#page-10-0)). It has been documented that V. ceranae infection significantly increased the mortality rate of A. cerana workers ([56\)](#page-11-0). In this study, the survival rates of workers in both the V. ceranae- and un-inoculated groups were high at 1–5 dpi ([Figure 7\)](#page-8-0), implying that V. ceranae had little influence on host longevity at the early stage of infection. However, at 5–20 dpi [\(Figure 7\)](#page-8-0), the survival rate of V. ceranaeinoculated workers was always lower than that of uninoculated workers, and the difference was significant at 11–20 dpi, indicating that V. ceranae infection shortened the lifespan of A. cerana workers, similar to the findings of Huang et al. ([56\)](#page-11-0). In conclusion, V. ceranae can negatively affect the midgut epithelial cell structure and lifespan of A. c. cerana workers during infection.

Collectively, the findings of the present study indicate that the spore load continuously increases as V. ceranae proliferates in the midgut of A. c. cerana workers, giving rise to continuous energetic stress for the host, apparent damage to the host midgut epithelial cell structure, induction of CAT and GST activities in the host midgut at the middle stage, and inhibition of the activities of SOD, CAT, and GST at the late stage of infection, which ultimately shortens the host lifespan [\(Figure 8](#page-8-0)).

#### 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 author.

## Ethics statement

The animal study was approved by Laboratory Animals Ethics and Welfare Committee of College of Animal Sciences, Fujian Agriculture and Forestry University (PZCASFAFU24018). The

study was conducted in accordance with the local legislation and institutional requirements.

# Author contributions

XF: Methodology, Software, Validation, Writing – original draft. HaZ: Methodology, Software, Validation, Writing – original draft. HeZ: Conceptualization, Methodology, Software, Validation, Writing – original draft. SD: Software, Validation, Writing – original draft. JQ: Data curation, Methodology, Writing – original draft. YS: Formal analysis, Writing – original draft. KL: Formal analysis, Writing – original draft. HJ: Data curation, Writing – original draft. YW: Data curation, Writing – original draft. YL: Data curation, Writing – original draft. DZ: Data curation, Visualization, Writing – original draft. ZF: Formal analysis, Methodology, Visualization, Writing – review & editing. DC: Conceptualization, Funding acquisition, Project administration, Writing – review & editing. RG: Conceptualization, Funding acquisition, Project administration, Supervision, Writing – review & editing.

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## Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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