Effects of cage and floor rearing system on the factors of antioxidant defense and inflammatory injury in laying ducks

Yang Zhang†, Tiantian Gu†, Yong Tian, Li Chen, Guoqin Li, Wei Zhou, Guofa Liu, Xinsheng Wu, Tao Zeng, Qi Xu, Guohong Chen* and Lizhi Lu‡

Abstract

Background: Cage-rearing in laying ducks, as a novel rearing system, not only fundamentally solves the pollution problem of the duck industry and improve bio-safety and product quality but also exhibits more benefits by implementing standardized production compared with the floor-rearing. Of course, this system also brings some welfare problems and stress injuries to layers due to lack of water environment and limited activities in the cages. However, the effects on the factors of antioxidant defense and inflammatory injury in the early cage stage are not well-understood.

Results: In this study, eighty Shaoxing layers were reared on floor and in cages from 12 weeks of age. The ducks were caged1, 2, 4, 7, and 10 days, the factors of antioxidant defense and inflammatory injury were investigated. The results showed that the caged ducks suffered liver injury to a certain extent when the ducks were just put into the cages. Analysis of antioxidant enzyme activities indicated that the different rearing system could not affect the change of antioxidant capacities, while the liver malondialdehyde (MDA) level was significant higher in the 2-d, 7-d, and 10-d ducks compared with the 1-d ducks during the change of days, while catalase (CAT) activity showed the opposite results. Additionally, quantitative real-time PCR (qRT-PCR) revealed that the relative mRNA levels of endoplasmic reticulum (ER) stress-related gene (CHOP and GRP78) were significantly upregulated in cage rearing ducks compared to that of the floor rearing ducks. Moreover, the mRNA levels of inflammatory cytokines including cyclooxygenase-2 (COX-2), nitric oxide synthase (iNOS), Interleukin 1 beta (IL-1β), Interleukin 2 (IL-2) and Interleukin 6 (IL-6), were also increased significantly in caged layers.

Conclusions: Taken together, although antioxidant defense has no obvious effect on cage stress, the stress levels of laying ducks vary greatly in the early cage stage, which not only caused liver tissue damage to some extent, but also resulted in increases in the expression of the factors of inflammatory injury. Therefore, we recommend that anti-stress agents should be added in the feed to alleviate the stress in the early cage stage.

Keywords: Cage rearing, Duck, Antioxidant enzymes, Inflammatory cytokines

Background

The market demand for ducks is increasing, and the scale of breeding must be expanded. Many resources such as land area are severely limited, and free-range methods are insufficient to meet consumer demands. Therefore, it is necessary to increase the scale of cage rearing. Previous studies reported that growing laying ducks in cages mode can increase the egg production rate and feed-egg ratio and lower the egg breakage rate with no large effects on egg quality [1–3]. However, the influence of stress on the ducks grown in cages remains unclear.

All living organisms respond to different types of environmental stressors by synthesizing oxidative stress proteins through a variety of signaling pathways, such as the endoplasmic reticulum (ER) stress response [4]. ER

* Correspondence: ghchen@yzu.edu.cn; lulizhibox@163.com
†Yang Zhang and Tiantian Gu contributed equally to this work.
1Jiangsu Key Laboratory for Animal Genetic, Breeding and Molecular Design, Yangzhou University, Yangzhou 225009, Jiangsu, China
2Institute of Animal Husbandry and Veterinary Medicine, Zhejiang Academy of Agricultural Sciences, Zhejiang 310021, Hangzhou, China
Full list of author information is available at the end of the article

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stress is considered as an early or initial response of cells to stress or damage [5]. In chicken, the key regulator of the mRNA levels of the ER stress gene (GRP78) in under selenium-deficient stress was significantly elevated in the liver [6]. Additionally, the CHOP protein content is increased in pig exposed to high temperatures [7]. Oxidative stress can enhance the formation of reactive oxygen species (ROS), which induce lipid, protein, or DNA oxidation and enhance lipid peroxidation to cause oxidative injury [8]. The activities of superoxide dismutase (SOD) and catalase (CAT) and levels of malondialdehyde (MDA) are increased in the liver of broiler chickens during heat stress [9]. In the masseter muscles of psychologically stressed rats, glutathione peroxidase (GSH-Px) and CAT activities were decreased, and MDA content was increased after 3 and 5 weeks [10]. Exposure to various stress conditions can also induce inflammation. It has been reported that the level of interleukin (IL)-2 mRNA in humans is increased under cold stress [11]. Under chronic cold stress, IL-2 and IL-10 contents were increased in the spleen and bursa of Fabricius of chicken [12]. In addition, the expression of inflammatory factors such as iNOS and COX-2 are influenced by exposure to high temperature in the duck liver [13].

Cage-rearing of laying ducks, as a novel culture mode, is beneficial for improving land utilization, feed conversion and egg production rate while reducing the egg breakage rate. However, few studies have examined the influence of cage stress on laying ducks or molecular mechanism of cage stress. Thus, we compared the microstructures of the liver in ducks reared using two different breeding patterns by histopathological analysis and investigated the effect of cage stress on antioxidant enzyme activity (SOD, MDA, CAT, total antioxidant capacity (T-AOC), and GSH-Px) in the livers of ducks. Furthermore, the expression of ER stress-related genes (GRP78 and CHOP) and inflammatory factors (COX-2, iNOS, IL-1β, IL-2 and IL-6) were analyzed during cage stress. These data improve the understanding of the influence of cage stress, providing insight useful for the large-scale breeding of caged ducks.

**Methods**

**Experimental design and management**

The experimental animals were female Shaoxing ducks (*Anas platyrhynchos*) obtained from Guiliu Animal Husbandry Company (Henan, China), and were always raised on the ground before the period of trial. And then eighty 12-week-old ducks with similar body weights were randomly divided into two groups. The forty ducks were reared on the floor (RF ducks), the others were reared from the ground into the cage (RC ducks), which were in closed-end animal building. The RF ducks were fed in semi-enclosed house (0.375 m²/per) and the RC ducks were raised in alone per cage (28 × 40 × 40 cm). All ducks kept at room temperature with flowing air for adaptation. The ducks were fed ad libitum with the same commercial formula diet (Henan Jinjing Biochemical Co., Ltd., Henan, China), which mainly contained corn, soybean meal, rice bran, and wheat-middling throughout the study. The ducks were subjected to a standard light regimen of 17 h light (17 L:7D) throughout the experimental period. All experimental ducks were healthy and were not administered any antibiotic treatments during the experiment.

**Histopathologic analysis**

For conventional histopathologic analysis, the liver tissue was fixed in 4% buffered formaldehyde and then embedded in paraffin, and 4-μm thick serial sections were prepared and stained with hematoxylin and eosin according to standard protocols. The sections were analyzed under an Olympus light microscope (Tokyo, Japan) to detect evidence of injury. According to the degree of light to heavy lesions, a small amount or no lesion was considered as negative and given a score of 0; mild or small was scored as 1; moderate or medium was scored as 2; severe was scored as 3; very heavy was scored as 4.

**Antioxidant enzyme activities**

The activities of SOD (U/mL), CAT(U/mL), T-AOC(U/mL), and GSH-Px(U/mL) and the MDA (nmol/mL) level were detected using superoxide dismutase assay kit (HY-60001), catalase assay kit (HY-60015), total antioxidant capacity assay kit (HY-60021), glutathione assay kit (HY-60006), and malondialdehyde assay kit (HY-60003), respectively. The experiments were performed according to the manufacturer’s protocols.

**RNA extraction and quantitative real-time PCR (qRT-PCR)**

Total RNA was extracted from the liver tissue using Trizol (TAKARA, Dalian, China). The RNA was resuspended in RNase-free water, and the concentration and purity were measured using a NanoDrop Spectrophotometer (NanoDrop...
Technologies, Wilmington, DE, USA). RNA (1 μg) was mixed with gDNase (Toyobo, Osaka, Japan) for reverse transcription. The process included an initial step at 37 °C for 5 min, followed by incubation at 37 °C for 15 min, 50 °C for 5 min, and 98 °C for 5 min; cDNA was stored at −80 °C for qRT-PCR. The primers used are listed in Additional file 1: Table S1. The qRT-PCR assay was carried out on a LightCycler 96 (Roche, Basel, Switzerland) with samples containing 10 μL SYBR Green Master Mix, 0.4 μL forward/reverse primer, 2 μL cDNA template and 7.2 μL RNA-free water (Vazyme, Nanjing, China). Reaction conditions included one cycle at 95 °C for 30 s, 40 cycles of 95 °C for 10 s and 60 °C for 30 s. Experiments detecting all genes were performed in triplicate and expression levels were assessed relative to duck β-actin as an internal standard. Relative gene expression levels were calculated using the 2^{−ΔΔCt} method.

Statistical analysis
The comparisons among different rearing systems and different days in the cage were fulfilled using two-way ANOVA in SPSS 13.0 software (SPSS, Inc., Chicago, IL, USA). P level below 0.05 were considered to indicate statistical significance. All data were analyzed using GraphPad Prism 5.0 software (GraphPad, Inc., La Jolla, CA, USA) and the results are presented as the means ± S.E.

Results
Histopathology of the liver
To determine the changes in liver tissue after cage stress, histological analysis was performed. The results indicated that RF ducks showed normal histological structure (Fig. 1a1-e1), while the livers of ducks reared in battery cages displayed some tissue injury corresponding to the time of cage stress (Fig. 1a2-e2). After 1 and 2 days of stress, the RC ducks showed severe liver injury, infiltration of inflammatory cells, and exudation of blood cells compared to the RF ducks (Fig. 1a2 and b2), indicating variable cellular vacuolization and hydropic degeneration in the liver in the early days of cage stress exposure. As the time of cage stress increased, liver injury got better after the 4-d, 7-d, and 10-d cage stress period (Fig. 1c2, d2, and e2).

Fig. 1
Histopathology of the liver. Hematoxylin and eosin staining of liver sections in the floor ducks (a1-e1) and cage ducks (a2-e2): a, b, c, d, e represented the days in the cage (1, 2, 4, 7, and 10 d, respectively). Black arrow represented cellular vacuolization
Antioxidant enzyme activity
To further characterize the effect of cage stress on antioxidant capacity of Shaoxing ducks, the SOD, CAT, T-AOC, and GSH-PX activities and MDA level were measured. We observed that rearing systems did not cause significant changes on the activity of MDA, SOD, CAT, T-AOC, and GSH-PX (Fig. 2). On the other hand, the MDA level appeared a significant increase at 2 days and then showed a gently trend due to the change of days, besides, the MDA level was significant higher in the 2-d, 7-d, and 10-d ducks compared with the 1-d ducks during the change of days (Additional file 2: Table S2). While catalase (CAT) activity showed the opposite results that the CAT activity showed a great decreasing trend and then got a peaked significantly at 4 days.

Expression of the liver CHOP and GRP78 in Shaoxing ducks
To evaluate the expression of ER stress response marker genes in the liver of Shaoxing ducks after cage stress, CHOP and GRP78 gene levels were measured by qRT-PCR. There was a significant interaction between factors (rearing system and days in the cage) for the CHOP and GRP78 level (Fig. 3). We found that CHOP mRNA was significantly upregulated in the RC ducks after 1 and 2 days of cage rearing, particularly in the RC ducks after 2 days, while the CHOP mRNA in RF ducks showed a gentle process (Fig. 3a). And comparing with the RF ducks, the CHOP mRNA expression in RC ducks was significantly increased in 1d and 2d. The mRNA expression of the GRP78 gene was significantly increased after 1 and 10 days of captivity, comparing with RF ducks (Fig. 3b).

Expression of liver inflammatory cytokines in Shaoxing ducks
To determine the effect of cage stress on the mRNA levels of inflammatory genes in the liver. The significant interaction between factors (rearing system and days in the cage) for the COX-2, iNOS, IL-1β, IL-2 and IL-6 mRNA level was observed (Fig. 4). The iNOS mRNA expression in RF ducks appeared a gentle tendency while the iNOS mRNA level in RC ducks performed the tendency that first decreasing and then rising, especially in 10 days of cage stress. Besides, the iNOS mRNA levels in the RC ducks were significantly higher than that in the RF ducks, particularly in the 1-d, 4-d and 10-d (Fig. 4a). Meanwhile, the expression of COX-2 mRNA showed similar tendency with iNOS whether in the RF ducks or the RC ducks. The mRNA expression of COX-2 was significantly increased in the 1-d, 7-d, and 10-d cage stress group (Fig. 4b). Furthermore, the mRNA level of IL-1β was gradually upregulated, reaching a peak after 7 days in the RC ducks and the IL-1β mRNA levels in the RC ducks were significantly higher than that in the RF ducks, particularly in the 4-d and 7-d RC ducks (Fig. 4c). The IL-2 mRNA level was higher in the RC ducks than that in RF ducks, and was particularly significant in 1-d, 7-d and 10-d RC ducks. IL-2 mRNA levels showed a decreasing trend and then gradually increased during cage stress.
stress (Fig. 4d). Finally, the IL-6 mRNA expression in RC ducks showed no significant change until 10-d cage stress and IL-6 mRNA levels were significantly higher in the 10-d RC ducks than in the RF ducks (Fig. 4e).

**Discussion**

Environmental stress is experienced by most animals and can induce various responses involving the balance of the oxidant/anti-oxidant system, as well as cause oxidative damage to several tissues by altering the enzymatic and non-enzymatic antioxidant status and enhancing ROS production [14, 15]. The anti-oxidative enzyme system, including SOD, CAT, and GSH-Px, plays an important role in the first line of antioxidant defense and MDA acts as a general biomarker for biological oxidative stress [16]. Previous studies showed that the MDA level were upregulated significantly in chickens exposed to high temperatures [17]. Other studies demonstrated that the MDA content was increased by chronic heat stress [18]. While in the present study, the activities of anti-oxidative enzymes in different rearing systems did not change significantly, and the result was inconsistent with...
other studies [17, 18], which indicates that the body of ducks has a certain degree of adaptability to the external environment and the balance of the oxidant and antioxidant systems was cooperated to repair the ability of scavenge free radicals [19]. However, histological analysis of Shaoxing ducks revealed severe liver injury, infiltration of inflammatory cells, and exudation of blood cells, particularly in the 1-d and 2-d RC ducks. These results indicate that in the early cage stress period, short-term stress stimulation was sufficient to cause stress damage to tissues of the body [15]. The liver, as the most abundant organ of protein metabolism, is closely related to protein processing and also the target of attack in the endoplasmic reticulum [20]. Once the endoplasmic reticulum is attacked, the unfolded protein reaction in the endoplasmic reticulum is intensified, eventually leading to hepatocytes in the endoplasmic reticulum stress state, aggravating cell damage [1]. Interestingly, after 4 days of stress test, the liver of caged ducks got better, which indicated that the cage stress caused some damage to the liver, but did not exceed the range of adaptation of the body. Therefore, after 4 days of stress, the liver tissue structure has a tendency to return to normal due to self-regulation of the body.

A recent study has been revealed that oxidative stress induces continuous ER stress by interfering with oxidation of the internal environment of the ER, and thus the ER stress response may occur downstream of oxidative stress [6]. To determine the effects of cage rearing on ER stress in ducks, we investigated the expression levels of ER stress-related signaling molecules including CHOP and GRP78 in the liver tissues of normal and caged ducks. A previous study showed that GRP78 expression was significantly upregulated in the liver during cage stress, indicating that cage rearing induces the ER stress response. Although rearing systems did not cause significant changes on the activity of anti-oxidative enzymes, the caged ducks suffered liver injury to a certain extent. These results suggest that cage stress could lead to tissue damage and increase the expression levels of inflammatory factors during the cage stress period, which may provide useful information for performing cage rearing of egg-laying ducks on a large scale.

Conclusions
In summary, we provide insight into the influence of the early stress of captivity on Shaoxing duck. Expression analysis showed that the CHOP and GRP78 genes were significantly upregulated in the liver during cage stress, indicating that cage rearing induces the ER stress response. Although rearing systems did not cause significant changes on the activity of anti-oxidative enzymes, the caged ducks suffered liver injury to a certain extent. These results suggest that cage stress could lead to tissue damage and increase the expression levels of inflammatory factors during the cage stress period, which may provide useful information for performing cage rearing of egg-laying ducks on a large scale.

Supplementary information
Supplementary information accompanies this paper at https://doi.org/10.1186/s12863-019-0806-0.

Additional file 1: Table S1. The primers of expressed genes detected in the study.

Additional file 2: Table S2. Main effect of rearing system and days in the cage on SOD, MDA, CAT, GSH-PX and T-AOC activity.

Abbreviations
CAT: Catalase; COX-2: Cyclooxygenase-2; ER: Endoplasmic reticulum; ES: Environmental stress; GSH-Px: glutathione peroxidase; IL: Interleukin; IL-1β: Interleukin 1 beta; IL-2: Interleukin 2; IL-6: Interleukin 6; iNOS: Nitric oxide synthase; MDA: Malondialdehyde; qRT-PCR: Quantitative real-time PCR; RC ducks: Reared in the cage; RF ducks: Reared on the floor; ROS: Reactive oxygen species; SOD: Superoxide dismutase; T-AOC: Total antioxidant capacity; UPR: Unfolded protein response

Acknowledgements
We are sincerely grateful to Prof Dr. Chen Guohong, Dr. Li Zhilu and all the teachers in our research team for their valuable guidance in the course of this research work. To all the members of the research team, we do appreciate the moral support and the immense support we received during the period of this research work.

Authors’ contributions
TG, GC, LL and QX conceived of the study, and participated in its design and coordination. YZ, TG, WZ and GL carried out RNA purification and quantitative RT-PCR. TG, LC and XW carried out antioxidant enzyme activity analysis and histopathologic experiment. YT, WZ, LC, GQL, TZ and XW participated in the design of the study and performed the statistical analysis. YZ, TG and YT contributed to the manuscript preparation. YZ, TG, LL and GC interpreted the results and contributed to edit the manuscript. All authors read and approved the final manuscript.
Funding
This study was supported by the National Natural Science Foundation of China (31402066), the National Key Research and Development Project (2018YFD05051040), the Research Program of Zhejiang Basic Public Welfare (LGN18CT170033), the Natural Science Foundation of Zhejiang Province (LQ17C170003) and the Natural Science Foundation of China (13172106, 31572385). These funding bodies played no role in the design of the study and collection, analysis, and interpretation of data and in writing the manuscript.

Availability of data and materials
The data sets supporting the results of this article are included within the article and its additional file.

Ethics approval and consent to participate
All protocols used in the present study were approved by the Institutional Animal Care and Use Committee of Yangzhou University (approval number: 151–2014). The Administration of Affairs Concerning Experimental Animals (Yangzhou University, China, 2012) and the Standards for the Administration of Experimental Practices (Jiangsu, China, 2008) were established to perform the procedures. The ducks were obtained from the Guiliu Animal Husbandry Company (Henan, China). We also confirm that all efforts were made to reduce the pain.

Consent for publication
Not applicable.

Competing interests
The authors declare that they have no competing interests.

Author details
1. Jiangsu Key Laboratory for Animal Genetic, Breeding and Molecular Design, Yangzhou University, Yangzhou 225009, Jiangsu, China. 2. Institute of Animal Husbandry and Veterinary Medicine, Zhejiang Academy of Agricultural Sciences, Zhejiang 310021, Hangzhou, China. 3. Key Laboratory of Information Traceability for Agricultural Products, Ministry of Agriculture of China, Hangzhou 310021, Zhejiang, China. 4. Guiliu Animal Husbandry Company, Zhoukou 450000, Henan, China. 5. Key Laboratory of Animal Genetics, Breeding and Molecular Design of Jiangsu Province, Yangzhou University, Yangzhou 225009, PR, China.

Received: 27 December 2018 Accepted: 22 December 2019
Published online: 30 December 2019

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