

# Alleviation of lead-induced oxidative stress and immune damage by selenium in chicken bursa of Fabricius

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**Abstract** We investigated lead (Pb)-induced oxidative stress and immune damage in the chicken bursa of Fabricius (BF) and the ameliorative effect of selenium (Se). Seven-day-old male chickens were randomly divided into four groups and were provided standard diet and drinking water, Na<sub>2</sub>SeO<sub>3</sub> added to the standard diet and drinking water, standard diet and (CH<sub>3</sub>COO)<sub>2</sub>Pb added to drinking water, and Na<sub>2</sub>SeO<sub>3</sub> added to the standard diet and (CH<sub>3</sub>COO)<sub>2</sub>Pb added to drinking water for 30, 60, and 90 days. The presence of Pb inhibited total antioxidant capacity (T-AOC), glutathione peroxidase (GPx), glutathione S-transferase (GST), superoxide dismutase (SOD), and catalase (CAT) activities; decreased glutathione (GSH) content; increased malondialdehyde (MDA) and hydrogen peroxide (H<sub>2</sub>O<sub>2</sub>) contents; inhibited interleukin (IL)-2 and interferon- $\gamma$  (IFN- $\gamma$ ) messenger RNA (mRNA) expression; and increased IL-4, IL-6, IL-10, IL-12 $\beta$ , and IL-17 mRNA expression. The presence of Se relieved all of the above Pb-induced changes. There were close correlations among GSH, CAT, T-AOC, SOD, GPx, MDA, and H<sub>2</sub>O<sub>2</sub> and among IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$ . Our

data showed that Pb caused oxidative stress and immune damage in the chicken BF. Se alleviated Pb-induced oxidative stress and immune damage in the chicken BF.

**Keywords** Lead · Selenium · Chicken · Bursa of Fabricius · Cytokine · Oxidative stress

## Introduction

Widespread usage of lead (Pb) in industrial and agricultural production, such as acid battery and pigments (Mao et al. 2008), has caused environmental pollution and is a matter of international concern (Kong et al. 2015). Moreover, Pb is a toxic heavy metal that is hazardous to humans and animals. Pb impaired intelligence in children (Canfield et al. 2003), increased mortality in white sturgeon over time (Vardy et al. 2014), and induced oxidative damage in rat livers (Mohammadi et al. 2014; Mabrouk et al. 2016). Birds are also sensitive to Pb exposure. High level of Pb has been linked to the death of egret and heron chicks from Pyeongtaek, Korea (Kim and Oh 2015), and 59 terrestrial bird species are known to have ingested Pb or suffered Pb poisoning from ammunition sources (Fisher et al. 2006). Pb can inhibit the normal function of mouse splenic macrophages (Bishayi and Sengupta 2006) and impair immune system functioning (Dietert and Piepenbrink 2006). The bursa of Fabricius (BF) is the central humoral immune organ and is unique to birds (Ribatti et al. 2006). However, little is known about the effect of excess Pb on chicken BF. Therefore, we investigated the effect of Pb toxicity on the chicken BF.

Oxidative stress is an important phenomenon in Pb-induced toxicity (Wang et al. 2009). Excess Pb increased malondialdehyde (MDA) content; decreased total antioxidant capacity (T-AOC), glutathione peroxidase (GPx), and

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superoxide dismutase (SOD) activities; decreased glutathione (GSH) content; and caused oxidative stress in tilapia spleens (Dai et al. 2012). In rat livers and erythrocytes, excess Pb decreased GPx, glutathione S-transferase (GST), SOD, and catalase (CAT) activities; decreased GSH content; increased MDA and hydrogen peroxide ( $H_2O_2$ ) contents; and caused oxidative stress (Omobowale et al. 2014). Cytokines are biomarkers of heavy metal-induced immune toxicity. Cadmium (Cd) decreased interleukin (IL)-2 and interferon- $\gamma$  (IFN- $\gamma$ ) messenger RNA (mRNA) expression and suppressed immune function in chicken splenic lymphocytes (Xu et al. 2015). In oral mucosal epithelial cells, mercury induced IL-4 mRNA expression and lupus-like lesions (Seno et al. 2013). Cd induced IL-6 and inflammatory response in rat plasma (Kataranovski et al. 2009). IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$  involved in manganese (Mn)-induced toxicity in chicken splenic lymphocytes (Lu et al. 2015). We speculated that Pb poisoning would alter oxidative stress indicators and cytokines in the chicken BF. Therefore, we examined indicators of oxidative stress (T-AOC, SOD, CAT, GPx, GST, GSH, MDA, and  $H_2O_2$ ) and cytokines (IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$ ).

Selenium (Se), an essential trace element in human beings and animals, can alleviate the toxicity of heavy metal ions such as Pb. Se alleviated Pb-induced toxicity in the livers and brains of *Cyprinus carpio* (Ozkan-Yilmaz et al. 2014) and in the kidney of Nile tilapia (Hashish et al. 2015). Se also alleviated Pb-induced toxicity in chicken livers (Xu et al. 2016) and sword cartilage (Gao et al. 2016). Previous research from our group demonstrated that Se alleviated Pb-induced toxicity in chicken livers (Wang et al. 2015) and muscles (Jin et al. 2016). However, Pb toxicity and the alleviative effect of Se on Pb toxicity in chicken BF remain unclear. In this study, we established a combined model of Se and Pb to investigate the effect of Pb on indicators of oxidative stress (T-AOC, SOD, CAT, GPx, GST, GSH, MDA, and  $H_2O_2$ ) and cytokines (IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$ ) and to determine whether Se can prevent oxidative stress and immune injury induced by Pb in the chicken BF.

## Materials and methods

### Animal model and tissue samples

All procedures used in this study were approved by the Northeast Agricultural University Institutional Animal Care and Use Committee. One hundred eighty 1-day-old healthy Hyline male chickens were fed a standard commercial diet (containing 0.49 mg/kg Se) and drinking water for 7 days prior to experiments. Chickens were randomly divided into four groups: the control group, the Se group, the Pb group, and the Se/Pb group. The control

group was fed a standard commercial diet and drinking water. The Se group was fed  $Na_2SeO_3$  (analytical reagent grade, Tianjinzhiyuan Chemical Reagent Co. Ltd., Tianjin, China) added to the standard commercial diet at 1 mg/kg Se and drinking water. The Pb group was fed a standard commercial diet and  $(CH_3COO)_2Pb$  (analytical reagent grade, Tianjinzhiyuan Chemical Reagent Co. Ltd., Tianjin, China) was added to the drinking water at 350 mg/L Pb, according to median lethal dose ( $LD_{50}$ ) for Pb in chickens (Vengris and Mare 1974) and the need of the chicken experiment in toxicology (Klaassen 2013). The Se/Pb group was fed  $Na_2SeO_3$ , which was added to a standard commercial diet at 1 mg/kg Se, and  $(CH_3COO)_2Pb$ , which was added to the drinking water at 350 mg/L Pb. All animals were housed in the Laboratory Animal Center, College of Veterinary Medicine, Northeast Agricultural University (Harbin, China), and given feed and water ad libitum, in accordance with chicken feeding operation procedures.

On the 30th, 60th, and 90th days of the experiment, 15 chickens in each group were randomly selected for euthanasia. BF samples were immediately collected, washed with ice-cold saline, and divided into three parts. The first part was fixed with 2.5% glutaraldehyde phosphate buffered saline (*v/v*, pH 7.2) for ultrastructure observation. The second part was immediately homogenized to determine GSH, MDA, and  $H_2O_2$  contents and the activities of T-AOC, GPx, GST, SOD, and CAT. The third part was immediately frozen in liquid nitrogen and stored at  $-80^\circ C$  for later determination of IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$  mRNA expression.

### Ultrastructure

Tissue from the BF (1.0 mm  $\times$  1.0 mm  $\times$  1.0 mm) was immediately put into 2.5% glutaraldehyde phosphate buffer saline (*v/v*, pH 7.2) for 3 h at  $4^\circ C$  followed by immersion in 1% (*v/v*) osmium tetroxide for 1 h at  $4^\circ C$ . Samples were washed in phosphate buffer saline then impregnated with epoxy resin. The ultrathin sections were stained with uranyl acetate and lead citrate. Ultrastructure was observed under a transmission electron microscope.

### T-AOC, GPx, GST, SOD, and CAT activities and GSH, MDA, and $H_2O_2$ contents

T-AOC, GPx, GST, SOD, and CAT activities and GSH, MDA, and  $H_2O_2$  contents were determined using diagnostic kits produced by NanJing JianCheng Bioengineering Institute (NanJing, China) following the manufacturer's instructions.

**IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$  mRNA expression**

*Primer synthesis*

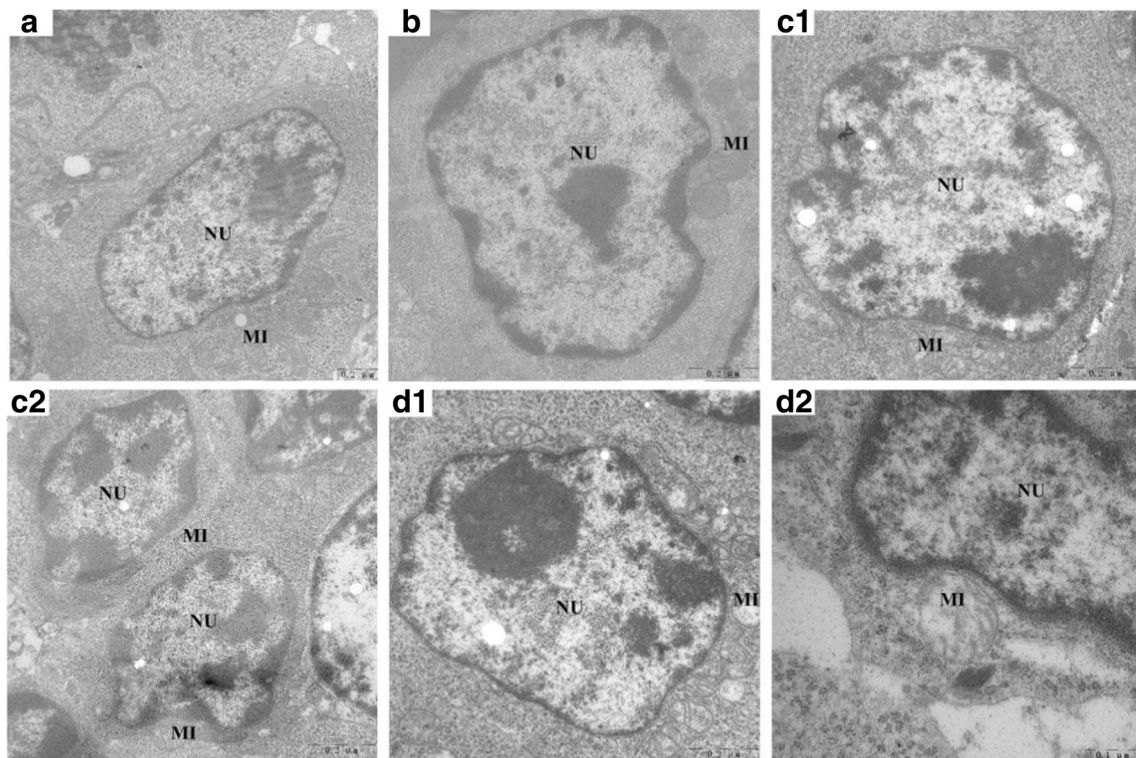
Primer sequences for  $\beta$ -actin (NM\_205518), IL-2 (NM\_204153), IL-4 (NM\_001007079), IL-6 (NM\_204628), IL-12 $\beta$  (NM\_213571), IL-17 (NM\_204460), and IFN- $\gamma$  (NM\_205149) published in GenBank were selected and synthesized by the Invitrogen Biotechnology Co. Ltd. (Shanghai, China) using the following sequences:  $\beta$ -actin forward sense: CTCTCGGCTGTGGTGGTGAA;  $\beta$ -actin reverse sense: CCGCTCTATGAAGGCTACGC; IL-2 forward sense: GAACCTCAAGAGTCTTACGGGTCTA; IL-2 reverse sense: ACAAAGTTGGTCAGTTCATGGAGA; IL-4 forward sense: GTGCCACGCTGTGCTTAC; IL-4 reverse sense: AGGAAACCTCTCCCTGGATGTC; IL-6 forward sense: AAATCCCTCCTCGCCAATCT; IL-6 reverse sense: CCCTCACGGTCTTCTCCATAAAA; IL-12 $\beta$  forward sense: TGTCTCACCTGCTATTTGCCTTAC; IL-12 $\beta$  reverse sense: CATACACATTCTCTCTAAGTTTCCACTGT; IL-17 forward sense: CATGTTGTCAGCCAGCATTCT; IL-17 reverse sense: CATCTTTTTGGGTTAGGCATCC; IFN- $\gamma$  forward sense: AAGTCATAGCGGCACATCAAAC; and IFN- $\gamma$  reverse sense: CTGGAATCTCATGTCGTTTCATCG.  $\beta$ -actin was used as an internal reference gene.

*Total RNA extraction and reverse transcription*

Total RNA was extracted from frozen BF samples using RNAiso Plus (Takara, Japan) according to the manufacturer’s protocol. The ratios of the optical densities at 260 and 280 nm (OD260/OD280) were measured using a GeneQuant 1300/100 spectrophotometer (Healthcare Bio-Sciences AB, Sweden) to determine RNA purity. The total RNA was immediately used to synthesize complementary DNA (cDNA). The cDNA was synthesized using a reverse transcriptase (RT) mix (HiGene, Harbin, China) containing 2.5  $\mu$ L of golden MLV reverse transcriptase, 6  $\mu$ L of 10 $\times$  RT buffer, 3  $\mu$ L of dNTP mixture (10 mM each), 6  $\mu$ L of total RNA, 3  $\mu$ L of 20 $\times$  oligo (dT) (25) and random primer, 1.5  $\mu$ L of RNase inhibitor (40 U/ $\mu$ L), and 38  $\mu$ L of RNase-free H<sub>2</sub>O. The RT program was 30  $^{\circ}$ C for 15 min, 55  $^{\circ}$ C for 30 min, and 80  $^{\circ}$ C for 10 min. The cDNA product generated by RT was diluted fivefold with sterile water and stored at -20  $^{\circ}$ C for real-time quantitative PCR.

*Real-time quantitative PCR*

Real-time quantitative PCR was performed using a LightCycler@96 (Roche, Switzerland) according to the manufacturer’s instructions. Reactions were performed in a 10- $\mu$ L mixture including 1  $\mu$ L of diluted cDNA, 0.3  $\mu$ L of forward



**Fig. 1** The ultrastructure of BF tissues. **a** the control group, **b** the Se group, **c** the Se/Pb group, **d** the Pb group. Magnification: **a**  $\times$ 15,000; **b**, **c1**, **c2**, **d1**  $\times$ 25,000; **d2**  $\times$ 40,000

primer (10  $\mu$ M), 0.3  $\mu$ L of reverse primer (10  $\mu$ M), 5  $\mu$ L of SYBR green master, and 3.4  $\mu$ L of PCR-grade water. The PCR program was 95  $^{\circ}$ C for 10 min, followed by 40 cycles of 95  $^{\circ}$ C for 15 s and 60  $^{\circ}$ C for 60 s, 95  $^{\circ}$ C for 10 s, 65  $^{\circ}$ C for 60 s, and 97  $^{\circ}$ C for 1 s. The melting curve showed a single peak for each PCR product. The experiment was repeated three times for each sample. The relative abundance of mRNA was calculated according to the Pfaffl method (Pfaffl 2001).

## Statistical analysis

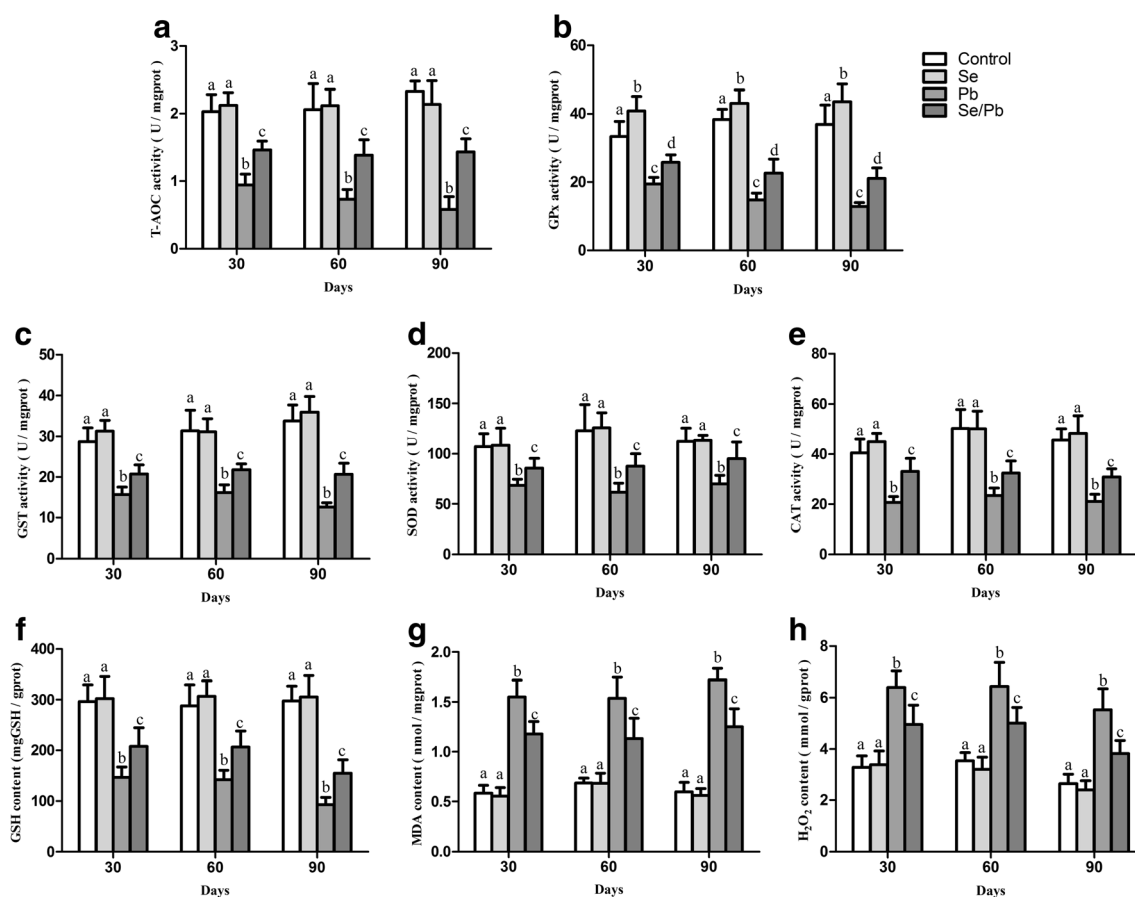
All data analyses were performed using SPSS (version 17; SPSS Inc., Chicago, IL, USA) for Windows. Statistical comparisons were performed using one-way ANOVA and verified by non-parametric Kruskal-Wallis and Mann-Whitney  $U$  tests. Different lowercase letters indicate significance among all groups at the same time point, with  $P < 0.05$ . Data are expressed as the mean  $\pm$  standard deviation. Pearson's  $r$  was used to measure the linear correlation among multiple

variables. Principal component analysis (PCA) was used to define the most important parameters, which were also used as key factors for individual variations using the Statistics 6.0 program (version 19, SPSS Inc., Chicago, IL, USA). All variables were standardized using z-scores and then calculated using the squared Euclidean distance. Finally, hierarchical clustering was performed with a standardized dataset using Ward's method.

## Results

### Ultrastructure

Observation of the ultrastructure of chicken BF on the 90th day revealed that cells in the control group (Fig. 1a) and in the Se group (Fig. 1b) were normal, with smooth rounded nucleolus (NU), normal chromatin, and intact mitochondria (MI) with normal cristae. Cells in the Se/Pb group had swollen mitochondria with degenerated and disordered cristae



**Fig. 2** The effect of Pb on T-AOC, SOD, CAT, GPx, and GST activities; GSH, MDA, and  $H_2O_2$  contents; and antagonistic effect of Se on Pb in chicken BF. Forty-five chickens consisted of 5 replicate pens, with each pen containing 9 chickens. Three chickens were randomly selected from

each pen on the 30th, 60th, and 90th days. One chicken each pen was measured. Bars represent mean  $\pm$  SD ( $n = 5$ ). Bars with different lowercase letters in different groups at the same time point are significantly different ( $P < 0.05$ )

(Fig. 1c1, 1c2). Cells in the Pb group had swollen mitochondria with even more pronounced cristae degeneration and vacuoles (Fig. 1d1, 1d2).

**T-AOC, GPx, GST, SOD, and CAT activities and GSH, MDA, and H<sub>2</sub>O<sub>2</sub> contents**

Measuring T-AOC, GPx, GST, SOD, and CAT activities and GSH, MDA, and H<sub>2</sub>O<sub>2</sub> contents revealed no significant differences ( $P > 0.05$ ) in T-AOC (Fig. 2a), GST (Fig. 2c), SOD (Fig. 2d), and CAT (Fig. 2e) activities and the contents of GSH (Fig. 2f), MDA (Fig. 2g), and H<sub>2</sub>O<sub>2</sub> (Fig. 2h) between the control group and the Se group on the 30th, 60th, and 90th days of the study. However, GPx (Fig. 2b) activity in the Se group was significantly higher ( $P < 0.05$ ) than that in the control group. T-AOC, GPx, GST, SOD, and CAT activities and GSH content in the Pb group were significantly lower ( $P < 0.05$ ) than those in the control, Se, and Se/Pb groups on the 30th, 60th, and 90th days. T-AOC, GPx, GST, SOD, and CAT activities and GSH content in the Se/Pb group were significantly lower ( $P < 0.05$ ) than those in the control and Se groups on the 30th, 60th, and 90th days. MDA and H<sub>2</sub>O<sub>2</sub> contents in the Pb group were significantly higher ( $P < 0.05$ ) than those in the control, Se, and Se/Pb groups on the 30th, 60th, and 90th days. MDA and H<sub>2</sub>O<sub>2</sub> contents in the Se/Pb group were significantly higher ( $P < 0.05$ ) than those in the control and Se groups on the 30th, 60th, and 90th days.

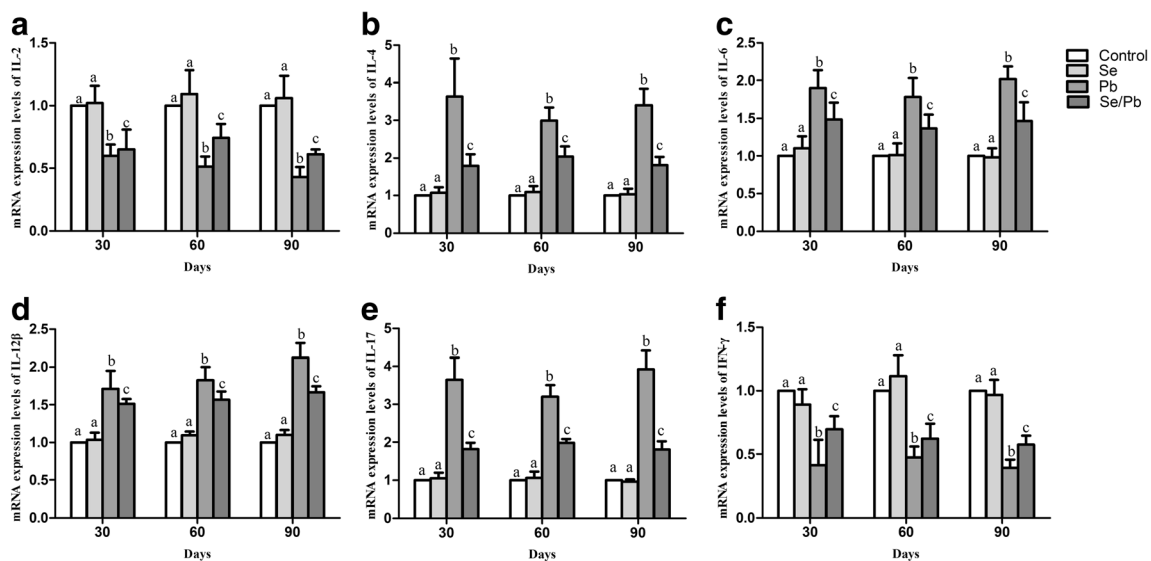
**IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$  mRNA expression**

As shown in Fig. 3, there were no significant differences ( $P > 0.05$ ) of IL-2 (Fig. 3a), IL-4 (Fig. 3b), IL-6 (Fig. 3c), IL-12 $\beta$  (Fig. 3d), IL-17 (Fig. 3e), and IFN- $\gamma$  (Fig. 3f) mRNA expression between the control group and the Se group on the 30th, 60th, and 90th days in the experiment. IL-2 and IFN- $\gamma$  mRNA expression in the Pb group was significantly lower ( $P < 0.05$ ) than that in the control, Se, and Se/Pb groups on the 30th, 60th, and 90th days. IL-2 and IFN- $\gamma$  mRNA expression in the Se/Pb group was significantly lower ( $P < 0.05$ ) than that in the control and Se groups on the 30th, 60th, and 90th days. IL-4, IL-6, IL-12 $\beta$ , and IL-17 mRNA expression in the Pb group was significantly higher ( $P < 0.05$ ) than that in the control, Se, and Se/Pb groups on the 30th, 60th, and 90th days. IL-4, IL-6, IL-12 $\beta$ , and IL-17 mRNA expression in the Se/Pb group was significantly higher ( $P < 0.05$ ) than that in the control and Se groups on the 30th, 60th, and 90th days.

**Chemometrics**

*Correlation coefficient analysis*

Pearson’s  $r$  correlation coefficient analysis (Table 1) indicated significant positive correlations among CAT, GPx, GSH, SOD, and T-AOC (at the 0.01 level); among IL-4, IL-6, IL-12 $\beta$ , and IL-17 (at the 0.01 level);



**Fig. 3** The effect of Pb on IL-2, IL-4, IL-6, IL-12 $\beta$ , IL-17, and IFN- $\gamma$  mRNA expression and antagonistic effect of Se on Pb in chicken BF. Forty-five chickens consisted of five replicate pens, with each pen containing nine chickens. Three chickens were randomly selected from

each pen on the 30th, 60th, and 90th days. One chicken each pen was measured. Bars represent mean  $\pm$  SD ( $n = 5$ ). Bars with different lowercase letters in different groups at the same time point are significantly different ( $P < 0.05$ )

**Table 1** Pearson's *r* correlation coefficient matrix among antioxidant factors and cytokines measured in the chicken BF

Factor	CAT	GPx	GSH	H <sub>2</sub> O <sub>2</sub>	MDA	SOD	T-AOC	IL-2	IL-4	IL-6	IL-12β	IL-17
GPx	0.950#	1.000										
GSH	0.949#	0.964#	1.000									
H <sub>2</sub> O <sub>2</sub>	-0.870#	-0.804#	-0.852#	1.000								
MDA	-0.921#	-0.912#	-0.968#	0.922#	1.000							
SOD	0.954#	0.903#	0.895#	-0.876#	-0.878#	1.000						
T-AOC	0.954#	0.941#	0.969#	-0.900#	-0.955#	0.938#	1.000					
IL-2	0.119	0.243	0.127	-0.009	-0.100	0.068	0.151	1.000				
IL-4	-0.105	-0.200	-0.159	0.215	0.242	-0.091	-0.213	-0.692#	1.000			
IL-6	-0.088	-0.193	-0.141	0.171	0.218	-0.065	-0.176	-0.725#	0.958#	1.000		
IL-12β	-0.321	-0.445#	-0.386*	0.279	0.387*	-0.295	-0.406*	-0.777#	0.849#	0.888#	1.000	
IL-17	-0.161	-0.249	-0.188	0.219	0.236	-0.160	-0.258	-0.728#	0.942#	0.910#	0.907#	1.000
IFN-γ	0.187	0.315	0.229	-0.040	-0.167	0.163	0.272	0.719#	-0.713#	-0.699#	-0.856#	-0.851#

\**P* < 0.05; #*P* < 0.01

between H<sub>2</sub>O<sub>2</sub> and MDA (at the 0.01 level); between IL-2 and IFN-γ (at the 0.01 level); and between GSH and IL-12β (at the 0.05 level). There were significant negative correlations between MDA and IL-12β (at the 0.05 level); between H<sub>2</sub>O<sub>2</sub> and CAT, GPx, GSH, SOD, and T-AOC (at the 0.01 level); between MDA and CAT, GPx, GSH, SOD, and T-AOC (at the 0.01 level); between IL-2 and IL-4, IL-6, IL-12β, and IL-17 (at the 0.01 level); and between IFN-γ and IL-4, IL-6, IL-12β, and IL-17 (at the 0.01 level). The results in Table 1 described the relationships among these factors and clearly revealed significant correlations among antioxidant factors and among cytokines.

### Principal component analysis

The results of the PCA are shown in Tables 2 and 3 and Fig. 4. Parameter determination was based on ordination plots, corresponding to the first and second principal components as 56.067 and 33.637%, respectively (Tables 2 and 3). In addition, Fig. 4 clearly indicated that GSH, CAT, T-AOC, SOD, GPx, MDA, and H<sub>2</sub>O<sub>2</sub> were closest to each other in the principal component (PC) 1 matrix, meaning that their relationships were closer based on PC 1. Among these antioxidant factors, GSH, CAT, T-AOC, SOD, and GPx had clearly opposite relationships with MDA and H<sub>2</sub>O<sub>2</sub> in PC 1 (Table 3 and Fig. 4). Moreover, Table 3 and Fig. 4 show that IL-4, IL-6, IL-

**Table 2** Component matrix in chicken BF

Component	Initial eigenvalues			Extraction sums of squared		
	Total	% of variance	Cumulative %	Total	% of variance	Cumulative %
1	7.289	56.067	56.067	7.289	56.067	56.067
2	4.373	33.637	89.704	4.373	33.637	89.704
3	0.572	4.402	94.107			
4	0.312	2.403	96.509			
5	0.176	1.353	97.862			
6	0.096	0.741	98.603			
7	0.063	0.484	99.087			
8	0.037	0.288	99.375			
9	0.028	0.219	99.594			
10	0.025	0.195	99.789			
11	0.014	0.108	99.897			
12	0.009	0.069	99.966			
13	0.004	0.034	100.000			

**Table 3** The correlation coefficients of the former two principal components in chicken BF

Element	PC1	PC2
CAT	0.979	-0.065
GPx	0.947	-0.190
GSH	0.973	-0.102
H <sub>2</sub> O <sub>2</sub>	-0.919	0.053
MDA	-0.966	0.123
SOD	0.957	-0.038
T-AOC	0.976	-0.152
IL-2	0.026	-0.842
IL-4	-0.068	0.935
IL-6	-0.041	0.942
IL-12β	-0.267	0.932
IL-17	-0.104	0.962
IFN-γ	0.115	-0.868
Explained variance (%)	56.067	33.637

IL-12β, and IL-17 were positively correlated with PC 2, contrary to IL-2 and IFN-γ.

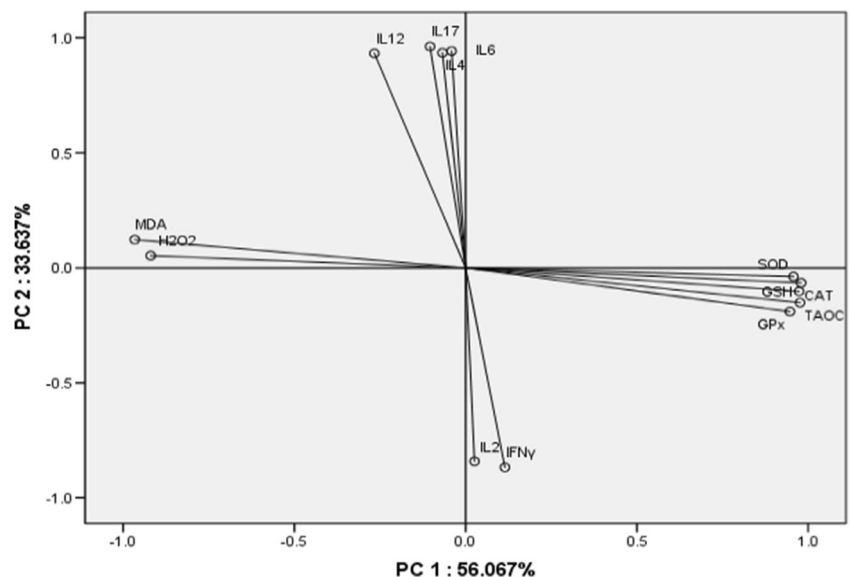
**Discussion**

Organisms maintain their homeostasis through antioxidant enzyme system, such as T-AOC, GPx, GST, SOD, and CAT, and non-enzymatic antioxidant system, such as GSH. Pb can damage mitochondria and the antioxidant enzyme system and reduce the antioxidant capacity of organisms (Venkareddy and Muralidhara 2015; Mabrouk et al. 2016). In rats, excess Pb

reduced the levels of GPx and GSH in livers (Liu et al. 2011) and kidneys (Liu et al. 2011; Lakshmi et al. 2013); increased MDA content in livers and kidneys (Lakshmi et al. 2013); and caused oxidative stress. Ni et al. (2004) found that Pb increased H<sub>2</sub>O<sub>2</sub> content and caused oxidative stress in human coronary artery endothelial cells and vascular smooth muscle cells. However, little is known about the effect of excess Pb on T-AOC, GPx, GST, SOD, and CAT activities and the contents of GSH, MDA, and H<sub>2</sub>O<sub>2</sub> in chicken BF. Therefore, we studied the effect of excess Pb on oxidative stress factors in the chicken BF. We found that Pb decreased T-AOC, GPx, GST, SOD, and CAT activities; decreased GSH content; and increased MDA and H<sub>2</sub>O<sub>2</sub> contents. Our results suggested that excess Pb caused oxidative stress in the chicken BF. Previous studies have shown similar mechanisms. Excess Mn decreased GPx activity, increased MDA content, and caused oxidative stress in chicken splenic lymphocytes (Zhu et al. 2016). Pb increased MDA content and reduced GPx, SOD, and CAT activities in worker erythrocytes (Chinde et al. 2014). Liu et al. (2013) showed that excess Mn reduced T-AOC, GPx, and SOD activities; increased MDA content; and caused oxidative stress in chicken BF.

Oxidative stress has been shown to impair the immune function of chickens (Zhang et al. 2012). Pb is toxic to immune function in humans and other species (Dietert et al. 2004). Heavy metals can change cytokines and cause immune toxicity in organisms. Excess Mn decreased IL-2 mRNA expression and induced immune suppression in chicken BF (Liu et al. 2012b) and splenic lymphocytes (Lu et al. 2015). Hemdan et al. (2005) reported that Pb induced IL-4, inhibited IFN-γ, and caused immune toxicity in human peripheral mononuclear blood cells. Excess Pb increased IL-6 and caused inflammation

**Fig. 4** Ordination diagram of PCA for the parameters that were measured in the chicken BF



in rat kidneys (Liu et al. 2012a). In rats, Pb poisoning during late pregnancy increased IL-12 and caused immune toxicity in the splenic cells of their offspring (Bunn et al. 2001). Cd increased IL-17 mRNA expression and induced immune suppression in rat spleens (Demenesku et al. 2014). Pb reduced IFN- $\gamma$  and caused immune suppression in rat T cells (Fang et al. 2012). Similar results were obtained in our study. We found that Pb increased IL-4, IL-6, IL-12 $\beta$ , and IL-17 mRNA expression and decreased IL-2 and INF- $\gamma$  mRNA expression in the chicken BF. Our results suggested that excess Pb caused immune toxicity in the chicken BF. Moreover, our morphological examination revealed that excess Pb caused ultrastructural changes and immune damage in the chicken BF. Liu et al. (2013) also found that excess Mn damaged the structure of chicken BF.

Se is involved in a variety of physiological processes in the form of selenoproteins (Yao et al. 2013). In our study, Se increased GPx activity in the chicken BF. Se can counteract Cd-induced oxidative stress and enhance immunity (Zwolak and Zaporowska 2012). Se intake improved immune function in adults (Broome et al. 2004) and antagonized the toxic effect of Cd and arsenic. Se ameliorated Cd-induced SOD and GPx activities and oxidative stress in chicken testes (Li et al. 2010) and livers (Li et al. 2013). In rats, Se ameliorated changes in MDA and IL-6 levels, oxidative stress, and hepatotoxicity induced by arsenic (Shafik and El Batsh 2016). Se also ameliorated the Cd-induced decrease in GPx, SOD, and CAT activities; GSH and IFN- $\gamma$  concentrations; increase in MDA and IL-6 concentration; oxidative stress; and immune suppression in rat livers and kidneys (El-Boshy et al. 2015). Se alleviated Cd-induced reticulum damage (Zhao et al. 2014); changes in IL-2, IL-4, IL-17, and IFN- $\gamma$  mRNA expression; and immune toxicity in chicken splenic lymphocytes (Xu et al. 2015). Se improved fish immune response induced by copper toxicity (Abdel-Tawwab et al. 2007). However, little is known about whether Se can alleviate Pb-induced immune toxicity. Our study showed that Se alleviated Pb-induced changes in indicators of oxidative stress (T-AOC, SOD, CAT, GPx, GST, GSH, MDA, and H<sub>2</sub>O<sub>2</sub>) and cytokines (IL-2, IFN- $\gamma$ , IL-4, IL-6, IL-12 $\beta$ , and IL-17), suggesting that Se alleviated Pb-induced oxidative stress and immune toxicity in the chicken BF.

In summary, our data showed that Pb caused mitochondrial swelling with cristae degeneration and vacuoles; inhibited T-AOC, GPx, GST, SOD, and CAT activities; decreased GSH content; increased MDA and H<sub>2</sub>O<sub>2</sub> contents; inhibited IL-2 and IFN- $\gamma$  mRNA expression; and induced IL-4, IL-6, IL-10, IL-12 $\beta$ , and IL-17 mRNA expression. Se alleviated Pb-induced changes to ultrastructure and to all of the above factors. Our data demonstrated that Pb caused oxidative stress and immune damage in the

chicken BF. Se alleviated Pb-induced oxidative stress and immune damage in the chicken BF. Our research suggested that the mechanism for Pb-induced injury to the BF of chickens is as follows: excess Pb induces oxidative stress, then alters biomacromolecule functions of immune cells and the structure of BF, and finally influences cytokine mRNA expression in the chicken BF.

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#### Compliance with ethical standards

**Conflict of interest** The authors declare that they have no conflicts of interest.

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