



# Effect of dexamethasone on gene expression of cannabinoid receptor type 1 and adenosine monophosphate-activated protein kinase in the hypothalamus of broilers (*Gallus domesticus*)

Linglian Kong<sup>a</sup>, Chuanpi Xiao<sup>a,b</sup>, Hai Lin<sup>a</sup>, Johan Buyse<sup>c</sup>, Xianlei Li<sup>a</sup>, Zhigang Song<sup>a,\*</sup>

<sup>a</sup> Department of Animal Science, Shandong Agricultural University, Taian, Shandong 271018, China

<sup>b</sup> Precision Livestock and Nutrition Unit, Gembloux Agro-BioTech, University of Liège, Gembloux 5030, Belgium

<sup>c</sup> Division Animal and Human Health Engineering, Department of Biosystems, KU Leuven, Kasteelpark Arenberg 30, Leuven 3001, Belgium

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## ABSTRACT

Hypothalamic neural circuits play a critical role in integrating peripheral signals and conveying information about energy and nutrient status. We detected cannabinoid receptor type 1 (CB1) distribution in the hypothalamus, liver, duodenum, jejunum, and ileum among 7- and 35-day-old broilers. The effects of dexamethasone (DEX) on CB1 gene expression were evaluated by *in vitro* and *in vivo* experiments on glucocorticoid receptor (GR) and adenosine monophosphate-activated protein kinase (AMPK) in the hypothalamus of broilers. *In vitro*, hypothalamic cells from 17-day-old broiler embryos were incubated with either 0.1% dimethyl sulfoxide or DEX (100 nmol/mL) for 1 h. In the *in vivo* study, 28-day-old broilers were injected with DEX for 24 h or 72 h. Results showed that CB1 was mainly expressed in the hypothalamus, and 72 h DEX treatment increased the expression. One-day treatment of broilers with DEX did not change the hypothalamic CB1 gene expression. Moreover, DEX treatment for 24 h and 72 h increased the mRNA level of hypothalamic AMPK $\alpha$ 2 and GR. However, no differences were observed on the gene expression of CB1, GR, and AMPK $\alpha$ 2 in hypothalamic cells with DEX-treated for 1 h. In conclusion, CB1 is mainly expressed in the hypothalamus of broilers; 72-h DEX exposure can regulate the CB1 system and AMPK signaling pathway of the broiler hypothalamus.

## 1. Introduction

In poultry species, as in all animals, hypothalamic neural circuits play a critical role in integrating peripheral signals and conveying information about energy and nutrient status. Stressful conditions increase the activity of the hypothalamic-pituitary-adrenal (HPA) axis (Vandenborne et al., 2005). Glucocorticoids (GCs), the end products of the HPA axis, exert feedback effects at the hypothalamus and pituitary levels and counter-regulatory hormones by influencing carbohydrate, lipid, and protein metabolism. Corticotropin-releasing hormone is the primary feedback target of glucocorticoids at the hypothalamic level (Kretz et al., 1999). Dietary supplementation and subcutaneous injection of corticosterone can reduce the expression of the Corticotropin-releasing hormone gene in the hypothalamus; meanwhile, corticosterone can stimulate the food intake of broilers under high-energy diet (Song et al., 2011a).

Many of changes observed during GCs excess correspond to

metabolic steps regulated by adenosine monophosphate-activated protein kinase (AMPK). Previous studies suggested that AMPK regulates cellular and systemic energy homeostasis and acts as a sensor of energy status in broilers (Song et al., 2012). In poultry, other candidates that target neuroendocrine energy balance mechanisms are currently under investigation. The endogenous cannabinoid is an important signaling molecule in the neuroendocrine control of homeostasis and is involved in the regulation of stress response, energy metabolism, and other processes (Di et al., 2005). Endogenous cannabinoids bind to cannabinoid receptor type 1 (CB1) receptors on presynaptic axon terminals and inhibit the release of  $\gamma$ -aminobutyric acid and glutamate. Moreover,  $\gamma$ -aminobutyric acid, the major inhibitory neurotransmitter in the central nervous system (CNS), can relieve stress (Freund et al., 2003). In this context, the endogenous cannabinoid system represents a new and promising research target in poultry.

Cannabinoids are natural lipophilic products from the flower of *Cannabis sativa* (Di Marzo et al., 2004). The two known types of

\* Corresponding authors.

E-mail addresses: [zhigangs@sdau.edu.cn](mailto:zhigangs@sdau.edu.cn), [245777398@qq.com](mailto:245777398@qq.com) (Z. Song).

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cannabinoid receptors, CB1 (Matsuda et al., 1990) and CB2 (Munro et al., 1993), have been identified. CB2 is predominately found in cells of the immune system, and it is only present in microglia within the brain (Núñez et al., 2004). With mounting evidence of more, CB1 is a G protein-coupled cannabinoid receptor (Matsuda et al., 1990), widely distributed in the CNS (Terry et al., 2010; Tsou et al., 1998). Besides, CB1 is highly abundant in the brain, involved in synaptic signaling (Stincic and Hyson, 2008), and can present in the metabolically relevant peripheral organs, such as gastrointestinal tissues, liver, pancreas, muscle, adipose tissue, and adrenal gland (Breivogel and Childers, 1998; Pagotto et al., 2006). Several studies suggest that CB1 activation shows high efficiency and leads to profound biological effects by modulating the activity of a wide variety of hypothalamic neuronal populations. GCs act rapidly, probably through membrane receptors, on corticotropin-releasing-factor-synthesizing cells to release endocannabinoids. These act locally at presynaptic axons via cannabinoid receptors to reduce excitatory glutamatergic input, and therefore induce fast feedback inhibition of corticotropin-releasing factor, by GCs (Dallman, 2003). Co-application of GCs and a CB1 inverse agonist block the GC-induced suppression of HPA axis responses to acute stress (Evanson et al., 2010), suggested GCs act nongenetically to provide rapid feedback inhibition of the paraventricular nucleus of hypothalamus by GC actions at membrane-associated GR and subsequent dendritic synthesis and release of CBs (Myers et al., 2012). Furthermore, in mammals, GCs can increase the number of endocannabinoids and stimulate AMPK activity in the hypothalamus (Di et al., 2003). However, there are few studies about GCs effects on the cannabinoid system in broilers. It is hoped that future research will give evidence for an interaction between GCs and food intake, possibly mediated by endocannabinoid secretion from target neurons in the hypothalamus (Dallman, 2003). Therefore, it is hypothesized that there is CB1 in broilers and the effect of exogenous injection of DEX on hypothalamic gene expression was studied. Specifically, we investigated CB1 distribution in the hypothalamus, liver, and intestine of broilers 7- and 35-days post-hatching. Dexamethasone (DEX), a synthetic GC, was employed to induce the hyper-GC status in the present study (Cai et al., 2011). We also examined the effects of DEX on the mRNA levels of AMPK, CB1, and GR in the hypothermic cell culture prepared from broiler embryos. We then elucidated the 24 h and 72 h effects of DEX administration to broilers on the mRNA levels of hypothalamic CB1, GR, and AMPK.

## 2. Materials and methods

### 2.1. Animal care, birds, and experimental design

The study was approved by Shandong Agricultural University and carried out following the Guidelines for Experimental Animals of the Ministry of Science and Technology (19,881,114, Beijing, People's Republic of China). Forty-eight 1-day-old male Arbor Acres broilers ( $n = 48$ ) were obtained from a local hatchery (Processing Co., Ltd. Shandong Bao Farming). All birds were placed in raised-floor pens and raised in environmentally controlled rooms under standard conditions, and free access to feed and water during the breeding period. The experimental diets met the National Research Council (1994) requirements. The composition and nutrient levels of the diet are shown in Table 1. The temperature was 35 °C initially, which was gradually reduced by 2 °C per week according to the age of broilers, until it was decreased to 24 °C. The birds were given 23 h of light and 1 h of darkness in the first week, followed by 16 h of light and 8 h of darkness for subsequent research.

### 2.2. CB1 distribution

To investigate the distribution of CB1 in broiler tissues, eight birds with similar body weight (BW, 7 days:  $154 \pm 4.3$  g, 35 days:  $1540 \pm 11.7$  g) from each group were selected in 7 and 35 days of age. The tissue samples of the hypothalamus, liver, duodenum, jejunum, and ileum

**Table 1**

Composition and nutrient levels of the diet used in this study (% air-dry basis).

Item	Content, %
<b>Ingredients</b>	
Corn	52.194
Soybean meal	38.926
Soybean oil	5.129
Limestone	1.463
CaHPO <sub>4</sub>	0.80
NaCl	0.30
L-Lys • H <sub>2</sub> SO <sub>4</sub>	0.267
DL-Methionine	0.302
L-Threonine	0.099
Phytase (5000 IU/g)	0.02
Vitamins premix <sup>1</sup>	0.20
Mineral premix <sup>2</sup>	0.20
Choline chloride	0.10
Total	100.00
<b>Nutrient levels<sup>3</sup></b>	
CP	21.50
ME (MJ/kg)	12.81
Ca	0.95
Nonphytate phosphorus	0.44
Lysine	1.19
Methionine	0.59
Cysteine + Methionine	0.87
Threonine	0.76
Tryptophan	0.22

<sup>1,2</sup> Vitamin premix and mineral premix provided the following per kg of diet: VA 9000 IU, VD<sub>3</sub> 2000 IU, VE 11.0 IU, VK 1.00 mg, thiamine 1.20 mg, riboflavin 5.80 mg, niacin 66.0 mg, pantothenic acid 10.0 mg, pyridoxine 2.60 mg, biotin 0.20 mg, folic acid 0.70 mg, VB<sub>12</sub> 0.012 mg, Mn 100 mg, Zn 75.0 mg, Fe 80.0 mg, I 0.65 mg, Cu 8.00 mg, Se 0.35 mg. <sup>3</sup> Nutrient levels were calculated values.

were collected, immediately cooled in liquid nitrogen, and stored at -80 °C for further analysis. The sampling method was in accordance with our previous research (Lei et al., 2013).

### 2.3. DEX treatment

*In vivo* and *in vitro* experiments were designed to detect the effect of DEX on the gene expression of CB1, GR, and AMPK $\alpha$ 2 in hypothalamus. At 28 days of age, twenty-four birds were assigned to two groups of twelve birds, which were randomly subjected to one of the following treatments for 24 h or 72 h: subcutaneous injection of DEX (2 mg/kg BW) or sham-treated control (injection of saline). At the end of the experimental period (for each injection time), six birds were sacrificed from each group by exsanguination. Thereafter, 1 to 2 g of the sample was obtained from each hypothalamus according to our previous research (Hu et al., 2019), cooled in liquid nitrogen, and stored at -80 °C for further analysis.

*In vitro* treatment: hypothalamic cell culture and treatment. Fertile eggs (Arbor Acres, 41st week in the laying cycle) were obtained from a maternal flock in a commercial hatchery (Jinan SAIS Poultry). The eggs were incubated under standard conditions at the Department of Animal Sciences of Shandong Agricultural University. Neuron-enriched hypothalamic cell culture was prepared from birds at day 17 of incubation. The hypothalamus was obtained by following a previously described method (Song et al., 2012). The dissociated hypothalamic tissue was mechanically triturated. The cells were plated in six-well culture plate with a final density of  $1.5 \times 10^6$  cells/well containing minimum essential medium (MEM) Earl salts (Invitrogen, Carlsbad, CA, USA) with 5% horse serum (HS, Invitrogen), 5% fetal calf serum (Biological Industries), B-27 supplement (1 lg/mL, Invitrogen), 0.6% glucose, gentamicin (20 lg/mL), and 2 mM glutamine (Invitrogen). The plates were precoated 1 day before the experiments with 0.01% poly-L-lysine solution (Sigma-Aldrich, St Louis, MO, USA). On the fourth day of *in vitro*

experiments, the medium was switched to MEM supplemented with 10% HS, 20 mg/mL 50-fluoro-20-deoxyuridine, and 50 mg/mL uridine (Sigma-Aldrich) to limit glial growth. Finally, the medium was changed to non-serum medium on seventh day of *in vitro* experiments, and cells were incubated for 3 days at 37 °C in a humidified incubator containing 5% CO<sub>2</sub>. The hypothalamic cells were treated with 100 nmol/mL DEX (dissolved in non-serum medium with 0.1% dimethyl sulfoxide, DMSO) for 1 h, or received with non-serum medium contained 0.1% DMSO as a control. There are six replicates in each group, with one well of cells in six-well culture plate for each replicate.

#### 2.4. Total RNA extraction and real-time PCR

Total RNA was isolated by the guanidinium isothiocyanate method with Trizol reagent (Invitrogen, San Diego, CA). The concentration and purity of each RNA sample were detected using a NanoDrop photometer (ND-2000, Thermo Scientific, Wilmington, USA). The integrity of RNA was detected by 1% agarose gel electrophoresis. Reverse transcription of 1 µg total RNA was performed using PrimeScript™ RT reagent Kit with gDNA Eraser (RR047A, TaKaRa, Dalian, China). RT-PCR was performed to determine the gene expression using TB Green Premix Ex Taq (RR820A, TaKaRa) in ABI 7500 Real-Time PCR Systems (Thermo Scientific). The reaction program included the following: pre-denaturation at 95 °C for 10 s, then denaturation at 95 °C for 5 s for a total of 40 cycles, and finally annealing and extension at 60 °C for 40 s. Each reaction well was repeated 3 times, and the primer sequence is shown in Table 2. The amplification efficiency of primers was calculated with a standard curve. The specificity of the amplified product was verified by the melting curve. The geometric mean of the expression of  $\beta$ -actin and glyceraldehyde-3-phosphate dehydrogenase (*GAPDH*) was used to normalize the expression of the target gene. Relative gene expression levels of each target gene were analyzed by the 2<sup>-ΔΔCt</sup> method.

#### 2.5. Statistical analysis

Data were presented as mean ± SD. All data were checked for normality with Shapiro–Wilk test (95% confidence level). Statistical differences between groups were analyzed with one-way ANOVA with Tukey's multiple comparisons. SPSS software (SPSS 26.0, SPSS, Chicago, USA) was used. Differences were considered significantly different at  $P < 0.05$ .

**Table 2**  
Gene-specific primers used for broilers gene expression analysis.

Gene <sup>a</sup>	Association no.	Primer sequences	Amplicon size (bp)
<i>CB1</i>	NM_001038652.1	F: GGCTGTTTCCTTATGCCCT R: GCTTGGCCTTTTGTGCTACC	262
<i>GR</i>	NM_001037826.1	F: AACCTGCTCTGGCTGACTTCTC R: CCCATCACTTCCGATCTGTTT	121
<i>AMPKα2</i>	NM_001039605.1	F: GGGACTGAAACCAGAGAACG R: ACAGAGGAGGCATAGAGGAG	215
$\beta$ -actin	NM_205518.1	F: CAGACATCAGGGTGTGATGG R: TCAGGGGCTACTCTCAGCTC	183
<i>GAPDH</i>	NM_204305.1	F: GCCCAGAACATCATCCCA R: CGGCAGGTCCAGGTCAACA	137

<sup>a</sup> Abbreviations: *CB1*, cannabinoid receptor type 1; *GR*, glucocorticoid receptor; *AMPKα2*, AMP-activated protein kinase α2;  $\beta$ -actin, beta-actin; *GAPDH*, glyceraldehyde phosphate dehydrogenase.

### 3. Results

#### 3.1. Distribution of hypothalamic and peripheral *CB1* gene expression

As shown in Fig. 1, the *CB1* expressed both in the hypothalamus, liver, duodenum, jejunum, and ileum of 7- and 35-day-old birds. The *CB1* mRNA level in the hypothalamus was significantly higher than those in the liver, duodenum, jejunum, and ileum for 7- and 35-day-old broilers ( $P < 0.001$ ; Fig. 1A and B).

#### 3.2. *In vivo* and *in vitro* DEX treatment and hypothalamic *CB1*, *AMPKα2*, and *GR* gene expression

One- and three-days DEX treatments increased the hypothalamic *AMPKα2* and *GR* mRNA levels ( $P = 0.001$ ,  $P = 0.0007$ ;  $P = 0.046$ ,  $P = 0.04$ , respectively; Fig. 2A, B, C, and D). The mRNA levels of *CB1* did not change in the hypothalamus of 24 h DEX-treated birds ( $P = 0.505$ ; Fig. 2E). However, 72 h DEX treatment increased the hypothalamic *CB1* gene expression ( $P = 0.001$ ; Fig. 2F). Treatment of hypothalamic cells with DEX did not alter the gene expression of *AMPKα2*, *GR*, and *CB1* ( $P = 0.581$ ,  $P = 0.455$ ,  $P = 0.293$ , respectively; Fig. 3A, B, and C).

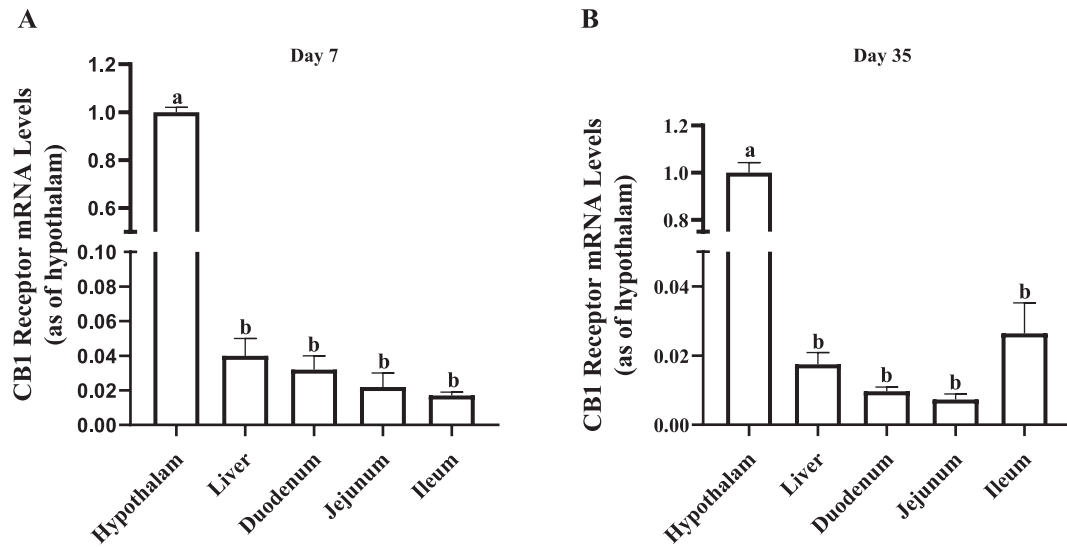
### 4. Discussion

#### 4.1. Distribution of *CB1* in 7- and 35-day-old broiler tissues

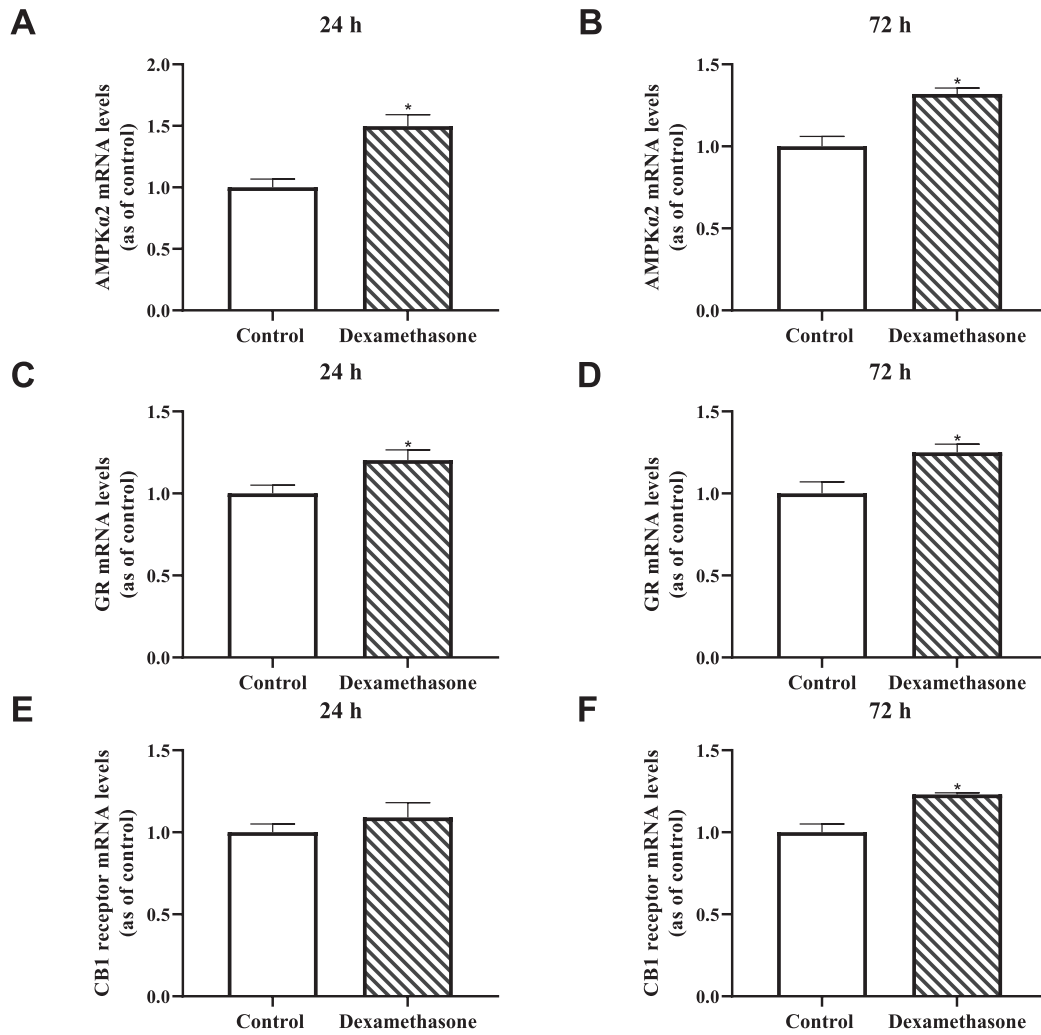
Previous reported indicated that *CB1* mRNA is present in the CNS (Terry et al., 2010; Tsou et al., 1998) and in peripheral tissues, including adipose tissues, liver, and myocardium, of mammalian species (Butler and Korbonsits, 2009). Furthermore, *CB1* is expressed by neurons throughout the body, including the brain and spinal cord, autonomic, and enteric nervous systems (Núñez et al., 2004). However, *CB1* in broilers has not been studied sufficiently (Soderstrom and Johnson, 2000). In the current study, *CB1* was expressed in the peripheral tissue (liver and small intestine) of broilers, which was consistent with previous findings. Compared with the hypothalamus, the expression of *CB1* in the liver, duodenum, jejunum, and ileum is significantly reduced, which was consistent with previous reports that *CB1* is mainly expressed at the end of the cerebellum axon, and this distribution pattern has been confirmed throughout the brain (Pickel et al., 2012). It is reported that *CB1* are heterogeneously expressed within subregions of the hypothalamus; in particular, *CB1* are present at greater density within the paraventricular nucleus of hypothalamus relative to other hypothalamic regions (Wittmann et al., 2007). However, compared to other brain regions, *CB1* density in the hypothalamus is lower (Herkenham et al., 1991) in spite of the fact that cannabinoids have many effects on hypothalamic-mediated processes (Fernández-Ruiz et al., 1997). Besides, there is evidence for *CB1* expression at all levels of the HPA axis (Hillard et al., 2016). In birds, the expression pattern of *CB1* mRNA appears to be highly conserved across species in key regions such as parts of the cerebellum and forebrain, and displayed high levels of expression in a number of regions that are known to be involved in visual processing throughout the brain (Stincic and Hyson, 2008).

#### 4.2. Effects of DEX-treatment *in vivo* and *in vitro* on *CB1*, *GR* and *AMPKα2* gene expression in broiler hypothalamus

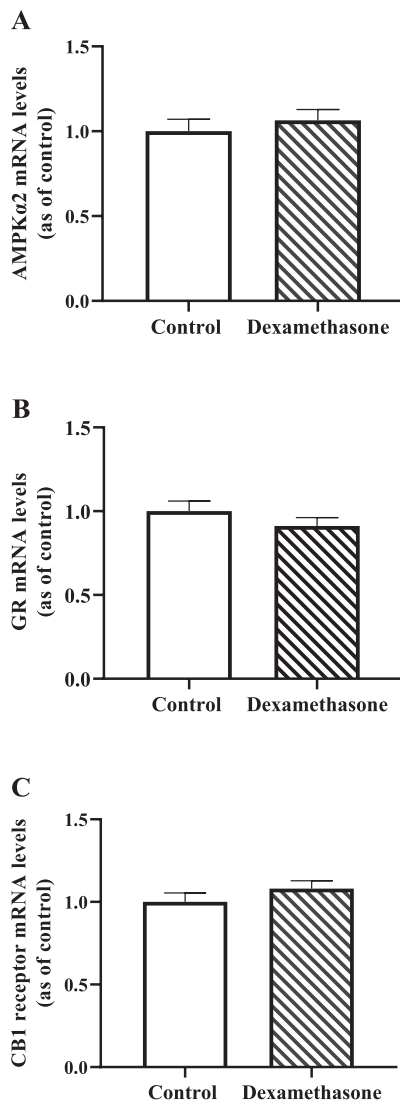
Although *CB1* is highly expressed through the brain, the regulatory effects of GCs on *CB1* expression are not ubiquitous throughout the CNS. In rodents, exogenous GCs cause different *CB1* expression responses in diverse tissues (Hill et al., 2005; Wang et al., 2007). In the current study, the mRNA levels of *CB1* did not change in the hypothalamus of 24 h DEX-treated birds, however, treated with DEX for 72 h showed an increased hypothalamic *CB1* gene expression. Consistent with the present findings, increased *CB1* expression was reported after 1 week of Glucocorticoid treatment in flushed bone and bone marrow cells of male



**Fig. 1.** Cannabinoid receptor type 1 mRNA levels in the hypothalamus, liver, duodenum, ileum, and jejunum of 7 (A) - and 35 (B) -days-old broilers. Values are means  $\pm$  SEM ( $n = 8$ ); <sup>a, b</sup> means with different letters differ significantly ( $P < 0.05$ ).



**Fig. 2.** Effect of 24 h (A) and 72 h (B) dexamethasone treatments on the hypothalamic AMPKα2 mRNA levels (*in vivo*); Effect of 24 h (C) and 72 h (D) dexamethasone treatments on the hypothalamic GR mRNA levels (*in vivo*); Effect of 24 h (E) and 72 h (F) DEX treatments on the hypothalamic CBI mRNA levels (*in vivo*). Values are means  $\pm$  SEM ( $n = 6$ ); asterisks indicate values that are significantly different between the control and DEX-treated group ( $* P < 0.05$ ).



**Fig. 3.** A: Effect of dexamethasone treatment on AMPK $\alpha$ 2 mRNA levels of the hypothalamic cells in broiler embryos (*in vitro*); B: Effect of dexamethasone treatment on GR mRNA levels of the hypothalamic cells in broiler embryos (*in vitro*); C: Effect of dexamethasone treatment on CB1 mRNA levels of the hypothalamic cells in broiler embryos (*in vitro*). Values are means  $\pm$  SEM ( $n = 6$ ).

rats (Ko et al., 2012). Treatment of cultured adipocytes with DEX for 24 h increased CB1 expression, suggesting that the effect of high GCs is mediated by enhanced CB1 expression (Pereira et al., 2014). Long-term corticosteroid treatment reduced the density of CB1 binding sites in the hippocampus and amygdala (Bowles et al., 2012). Repeated corticosteroid treatment resulted in a GR-dependent decrease in the CB1 protein expression in primary sensory neurons (Hong et al., 2011).

The endocrine response to physiological and psychological stress is mainly mediated by GCs by activating the HPA axis, which is released from the adrenal cortex. Glucocorticoid signaling is responsible for many physiological and psychological changes after stress, including increased food consumption, suppression of the immune system, and changes in memory and mood through effects on GRs (Hillard et al., 2016). This phenomenon was described in our early studies, wherein immune stress significantly reduced the food intake of broilers (Song et al., 2011b). Corticosterone treatment exhibits an immunosuppressive effect on the innate immune system of broilers, and such an effect could be enhanced by high dietary energy consumption (Yuan et al., 2008). A high-energy diet inhibits the appetite of broilers and reduces the gene

expression of GR in the hypothalamus (Hu et al., 2019). Several lines of evidence suggest that ligand-dependent GR activation mediates the role of GCs, which are expressed in almost every cell type. Activated GR, as a transcription factor, controls the expression level of target genes and regulates intracellular signaling pathways (Hafezi and Nouhi, 2002; Karin and Chang, 2001). Glucocorticoid receptor activation can increase CB1 expression. For example, cannabinoid receptor type 1 upregulation in the spinal cord of rats with compressive nerve injury depends on GR gene expression (Wang et al., 2007). The glucocorticoid-induced CB1 expression in mouse osteoblasts increased through GR-dependent regulation of transcription and translation (Wu et al., 2011). After long-term treatment with GCs, CB1 expression in rat bone was upregulated (Ko et al., 2012). In the present study, the change in the CB1 mRNA level of cultured broiler embryo hypothalamic cells were not significant. However, cannabinoid receptor type 1 may be related to GR gene expression in the DEX-exposed broiler hypothalamus. An implication of this result is the possible increase in CB1 mRNA expression due to increased GR mRNA expression. These mechanisms may differ because of factors, such as DEX treatment time.

Many of the changes observed during excessive GC levels correspond to the metabolic steps regulated by AMPK (Scerif et al., 2013). The effect of GCs on AMPK activity is tissue-dependent (Wang et al., 2010). AMPK in broilers is a regulator of cellular and systemic energy homeostasis and acts as a sensor of energy states (Song et al., 2012). The AMPK signal in the hypothalamus is related to control of food intake (Minokoshi et al., 2004). This finding is under the present observation that fasting-induced AMPK activation in the hypothalamus of broilers (Song et al., 2012). Meanwhile, activated AMPK stimulates neuropeptide Y gene expression and food intake, whereas inactivated AMPK causes anorexia in broilers (Liu et al., 2014). Liu et al. (2015) proposed the potential role of the hypothalamic rapamycin pathway target in appetite-related gene regulation by GCs; however, this finding should be further verified. Glucocorticoids participate in appetite control (Yuan et al., 2008). Previous studies suggested that hypothalamic CB1 activation is a powerful orexigenic signal and acts as a key determinant of energy balance in mice (Cardinal et al., 2012). In mammalian species, cannabinoid receptor type 1 is required for the hypothalamic AMPK effects of GCs (Scerif et al., 2013). In the present study, the gene expression of AMPK $\alpha$ 2 subunit and CB1 in 72 h DEX-treated broilers was significantly increased. CB1 that mediates AMPK regulation by GCs is one possible explanation, consistent with a previous report (Scerif et al., 2013). Different physiological and pathological stresses, such as glucose deprivation, ischemia, and hypoxia, leading to activation of AMPK (Kahn et al., 2005). Once AMPK is activated, it switches off anabolic pathways, such as fatty acid, triglyceride, and cholesterol synthesis, in favor of catabolic pathways, such as glycolysis and fatty acid oxidation, to restore energy balance.

In conclusion, CB1 is mainly expressed in the hypothalamus of broilers; long-term DEX exposure can regulate the CB1 system and AMPK signaling pathway of broiler hypothalamus. Moreover, hypothalamic cannabinoid signaling could be increased under conditions associated with hypersecretion of glucocorticoids, such as stress.

#### Conflicts of interest

We declare no conflict of interest with any financial organization regarding the material discussed in the manuscript.

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