



The effect of calcium propionate supplementation on performance, meat quality, and mRNA expression of finishing steers fed a high-concentrate diet

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KEY WORDS: steers, diet, calcium propionate, performance, meat quality, mRNA expression

ABSTRACT. The effects of calcium propionate supplementation on performance, meat quality, and mRNA expression of Wagyu steers were investigated. Eighteen steers (635 ± 20 kg; 18 ± 1 month old) were randomly divided into two groups: control (CG, without calcium propionate) and experimental (CaP, 200 g calcium propionate per steer per day). All steers were reared for 51 days under the same production system and then slaughtered at a final body weight of 680 ± 18 kg. The results showed no significant differences in dry matter intake, daily gain, or feed conversion ratio between the CaP and CG groups ($P > 0.05$). The treatments did not significantly affect the pH, drip loss, cooking loss, Warner–Bratzler shear, protein, fat and ash contents in meat ($P > 0.05$). The erucic acid (C22:1) content in group CaP was significantly lower than in CG ($P < 0.05$). The content of polyunsaturated fatty acids (PUFA) in CaP showed a decreasing trend compared with CG ($P = 0.06$). The expression of genes for peroxisome proliferator-activated receptor γ (*PPARG*) and CCAAT/enhancer binding protein α (*CEBPA*), which are involved in adipogenesis, was significantly higher in group CaP than in CG ($P < 0.05$). The results indicate that supplementing calcium propionate did not affect animal performance, but changed the composition of meat fatty acids, especially PUFA and erucic acid, and could trigger upregulation of *PPARG* and *CEBPA* mRNA expression levels, which could cause long-term activation of adipogenesis. Therefore, the results of the present study point to the possibilities of improving meat quality through calcium propionate supplementation of the diet.

Received: 18 August 2014

Revised: 29 April 2015

Accepted: 15 June 2015

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Introduction

In ruminants, dietary carbohydrates are fermented by rumen microorganisms into volatile fatty acids. Propionate is the primary precursor for glucose synthesis, and can supply 90% of this sugar in these animals (Bergman, 1975). Glucose provides energy,

as well as acetyl units (a synthetic fat precursor), half of which are used for synthesis of intramuscular fat, which improves meat quality (Smith and Crouse, 1984).

Wagyu steers are famous for their genetic predisposition for intense marbling (intramuscular fat in the cross section of the *longissimus* muscle) and

the production of meat with a high percentage of oleaginous unsaturated fats (Mir et al., 1999; Wheeler et al., 2004). High marbling levels improve the palatability and acceptability of beef by affecting its composition and tenderness (Nishimura et al., 1999). Marbling is one of the determinants of beef quality in the standards binding in the USA (USDA, 1989) and Japan (JMGA, 1988). Depositing body fat, especially intramuscular fat in finishing cattle, requires high-grain diets supplying large amounts of energy. Lee-Rangel et al. (2012) found that calcium propionate could be used to replace part of the energy supplied by grain in diets for finishing lambs. Liu et al. (2009a,b) found that adding calcium propionate promoted rumen fermentation, increased feed digestibility, and maintained the energy balance of lactating dairy cows.

Numerous recent studies have suggested that propionate could act as a signal molecule inducing preadipocyte differentiation (Lee and Hossner, 2002; Xiong et al., 2004; Hong et al., 2005). Wan et al. (2009) found that supplementing propionate could increase peroxisome proliferator-activated receptor γ (*PPARG*) and CCAAT/enhancer binding protein α (*CEBPA*) mRNA expression of bovine intramuscular preadipocytes in culture. CCAAT/enhancer binding proteins, C/EBPs, including C/EBP α , C/EBP β and C/EBP δ , were the first transcription factors shown to play an important role in the process of adipocyte differentiation (Lekstrom and Xanthopoulos, 1998). PPARs have three haplotypes: PPAR α , PPAR β and PPAR γ (Issemann and Green, 1990). Many studies have found that PPAR γ was required for the differentiation of adipose tissue *in vivo* and *in vitro* (Kutoba et al., 1999; Rosen et al., 1999).

The hypothesis of this study is that calcium propionate acting as an energy source and signal molecule could improve finishing efficiency and meat quality; this would be tested by examining growth performance, meat quality, and mRNA expression of adipogenic marker genes in the *longissimus* muscle of finishing steers.

Material and methods

Animals and experimental design

Eighteen medium-frame Wagyu steers (635 ± 20 kg; 18 ± 1 month old) were divided into two groups in a completely randomized design and housed in a tie-stall facility. The animals were handled in strict accordance with the guidelines approved by the Animal Welfare Committee of China Agricultural University (Permit Number: DK1008). The treatments were: control group (CG, without calcium

propionate) and experimental group (CaP, 200 g calcium propionate per steer per day). The calcium propionate (99.26%) was from Dong Xin chemical plant (Teng Zhou, Shan Dong Province, China). The trial period lasted 51 days, which included a 7-day introductory period and a 44-day experimental feeding period. The adjustment period of 1 week allowed the steers to become acclimated to routine feeding and to allot time for proper diet adjustment before the experiment. During the adjustment period, calcium propionate was gradually increased in the diet. In the experimental feeding period, calcium propionate was supplemented by adding it to 15% of concentrate feed and supplied separately. Because the steers were fed individually *ad libitum*, the Ca propionate was mixed with part of the concentrate in order to make sure that 200 g Ca propionate would be eaten completely. The feeding procedure was: first, 200 g of Ca propionate mixed with part of the concentrate feed (experimental group) or the same weight of concentrate feed (control group); next the total mixed ration was given to the cattle *ad libitum* after all steers eat all feeds given in the first step. Fifteen percent concentrate was the appropriate proportion to mix with 200 g of Ca propionate.

Diets and feeding management

The basal diet met the NRC (2000) requirements for 600 kg Wagyu with a weight gain of $1.0 \text{ kg} \cdot \text{d}^{-1}$. The basal diets consisted of 25% maize stover silage and 75% concentrate (Table 1). The steers were individually fed *ad libitum* at 06:00 and 17:00 and had free access to fresh water throughout the experimental period.

Table 1. Ingredients and nutrient level of the basal diet (DM)

Ingredients	g · kg ⁻¹ DM	Nutrients level	Content
Maize stover silage	250	NE _m ² , Mcal · kg ⁻¹	2.01
Maize	523.5	NE _g ³ , Mcal · kg ⁻¹	1.36
Cottonseed meal	40	%: CP	12.59
Brewer's grains	160	NDF	40.86
Magnesium oxide	2.5	ADF	18.22
Limestone	6	EE	4.44
Sodium bicarbonate	10	Ca	0.66
Calcium phosphate	1	P	0.33
Salt	5	ash	4.51
Premix ¹	2		

¹ contained: mg: Co 198, Cu 9228, Fe 80376, I 754, Mn 58, Se 366, Zn 66350; ^{2,3} calculated based on NRC (2000); DM – dry matter, CP – crude protein, NDF – neutral detergent fibre, ADF – acid detergent fibre, EE – ether extract, NE_m – net energy for maintenance, NE_g – net energy for growth

Feed analysis

Total mixed ration (TMR) feed samples were collected once a week and dried at 65°C in an air oven, then ground through a 1-mm screen using a Wiley Mill (A.H. Thomas Co., Philadelphia, PA, USA). Ground samples were used for determining dry matter (DM), crude protein (CP), ether extract (EE), ash, calcium and phosphorus according to AOAC (1999) methods (Table 1). Neutral detergent fibre (NDF) and acid detergent fibre (ADF) analyses were carried out according to Van Soest et al. (1991).

Growth

The body weight of all steers was measured (at the beginning and at the end of the finishing period) and refusals were recorded every day to calculate the average daily gain [ADG: (final body weight – initial body weight)/days], average dry matter intake (DMI) and feed conversion ratio (FCR: DMI/ADG).

Sampling procedures

At the end of feeding trial, all experimental steers were starved for 12 h before they were loaded and transported 5 km to a commercial slaughterhouse. Efforts were made to minimize suffering during transport and slaughter. Briefly, wide ramps with 11° slopes were used for loading and unloading steers, and the transport vehicle was equipped with non-slip flooring. Pre-slaughter handling systems were used to encourage smooth movement of steers. Steers were electrically stunned before slaughtering; they were restrained in an upright position with the head held fast and the neck exposed in a suitable position for incision of the throat. The knife has a long, extremely sharp blade. After slaughter, a portion of the *longissimus* muscle (1–2 g) from the 12th to 13th rib of the carcass was sampled immediately for RNA extraction. The sample tools were soaked in 1% diethyl pyrocarbonate in water to remove RNase. The samples were placed in sterile tubes (Corning[®] cryogenic vials, cat No. 430659, Corning Inc., NY, USA), packed in gauze bags and stored in liquid nitrogen. After a 72-h post mortem period (1°C to 4°C), the *longissimus* muscle from the 12th to 13th rib was sampled to determine meat quality.

Meat quality and fatty acid analysis

The pH of the *longissimus* muscle was measured directly using a pH meter (Eutech Instruments, pH Spear, Thermo Fisher Scientific Inc., Shanghai, China). One steak was cut into 15 mm × 15 mm × 30 mm and then suspended at 4°C for 48 h to calculate drip loss. Another steak was vacuum-

packed in a polyethylene bag and heated at 80°C until the internal temperature reached 70°C to calculate the cooking loss percentage, then divided into six 1 cm-diameter round strips for Warner–Bratzler shear force measurement (Salter Brecknell, Model 2356X Manhattan, KS, USA). The meat samples were lyophilized and pulverized for the determination of chemical composition, including moisture, protein, fat and ash according to AOAC (1999) methods. The chemical composition was determined after removing the connective tissue, subcutaneous fat and intermuscular fat. Therefore, the fat content was considered to reflect the intramuscular fat.

Fatty acids were determined by the use of a gas chromatography and, therefore, were first transformed into fatty acid methyl esters (FAMES). FAMES were prepared according to O'Fallon et al. (2007) with small modifications because of the high fatty acid content in Wagyu meat. The changes were in the weight of the meat samples and duration of manual shaking to completely extract fatty acids. The successive steps were: 1 ml of C13:0 internal standard (0.5 mg C13:0 · ml⁻¹ MeOH), 0.7 ml 10N KOH in water, and 5.3 ml MeOH were added to tubes containing 0.2 g of meat sample. After vortexing the tubes were incubated at 55°C in a water bath. The tubes were then cooled to below room temperature in a cold tap water bath and 0.58 ml 12N H₂SO₄ in water was added. The tubes were vortexed again and incubated at 55°C for 1.5 h with vigorous manual shaking for 5 s every 10 min with precipitated K₂SO₄ appearing. After FAME synthesis, the tubes were cooled in a cold tap water bath. Then 3.0 ml of hexane was added and the tubes were vortexed for 8 min and centrifuged for 5 min at 3000 rpm.

The hexane layer containing the FAMES was transferred into 2 ml gas chromatography (GC) vials. The vials were capped and stored at –20°C until GC analysis. The fatty acid composition of the FAMES was determined according to the O'Fallon et al. (2007) method. Fatty acids were identified by comparing their retention times to those of methylated fatty acid standards (Sigma, Sigma-Aldrich Shanghai Trading Co. Ltd., Shanghai, China). Fatty acid concentrations were expressed as mg · g⁻¹ of sample. The fatty acid composition of FAMES was determined by capillary GC on a SPTM-2560, 100 m × 0.25 mm × 0.20 m capillary column (Supelco, Bellefonte, PA, USA) installed on a Agilent 6890 GC (Agilent Technologies, Santa Clara, CA, USA). The initial oven temperature of 140°C was held for 5 min, subsequently increased to 240°C at a rate of 4°C · min⁻¹, then held for 20 min. Helium was used as the carrier gas at a flow rate of 0.5 ml · min⁻¹, and the column head pressure was

280 kPa. The injector and the detector were set at 250°C and 260°C, respectively. The split ratio was 30:1.

Total RNA extract and relative quantitative real-time polymerase chain reaction

Total RNA was extracted using the TRIzol (Biotek Corporation, Beijing, China) reagent. The concentration and the OD260 and OD280 values of the extracted RNA were determined using an ultraviolet spectrophotometer Nanodrop ND-1000 (Nanodrop Technologies, Inc., Willmington, DE, USA). The quality of the total extracted RNA was checked by agarose gel electrophoresis.

The cDNA was produced from 2 µg RNA using a super RT Kit PR6601 (Biotek Corporation, Beijing, China) and the protocol recommended by the manufacturer. The primer sequences of the target genes, *PPARG*, *CEBPA*, and a house-keeping gene, *GAPDH* (glyceraldehyde-3-phosphate dehydrogenase), were designed using Primer Express 2.0 software (Applied Biosystems, Stockholm, Sweden). The primer sequences were: *PPARG*, forward 5'-CGACCAACTGAACCCAGA-3' and reverse 5'-AGCGGGAAGGACTTTATGT-3'; *CEBPA*, forward 5'-CCCATAAAGCCAGCACT-3' and reverse 5'-CCACCTTCACGCAGAACA-3'; *GAPDH*, forward 5'-GGTGATGCTGGTGCTGAGT-3' and reverse 5'-ATGATGACCCTCTTGCG-3'. Quantitative real-time PCR (qPCR) was used to analyse the expression of *PPARG* (accession number: BC116098.1), *CEBPA* (accession number: BC149006.1), and the housekeeping gene, *GAPDH* (accession number: NM_001034034.1), which was used as an endogenous control (Smith et al., 2012). The qPCR reaction system had the following parameters: 1 µl of the cDNA template, 10 µl of 2 × SYBR real-time PCR premixture PR7002, (BioTeke, Beijing, China), 0.4 µl of the forward primer, 0.4 µl of reverse primer, and 8.2 µl to 20 µl of ddH₂O. The reaction liquid was transferred into a 96-well thermal sealing ring. The BIO 584BR detection system (Bio-Rad Co., Ltd., CA, USA) was used with the thermal cycling variables recommended by the manufacturer: one cycle at 94°C for 5 min, 45 cycles of denaturation at 94°C for 15 s, 60°C for 15 s and for annealing extension at 72°C for 15 s. The last extension step was performed at 72°C for 10 min.

A standard curve was generated separately for the housekeeping gene and each target gene. Five standard solutions differing in concentration were made by 10-fold dilutions. To reduce the error between the qPCR assays, every qPCR assay was used to make a standard curve. The results of *PPARG/*

GAPDH and *CEBPA/GAPDH* were expressed by -Lg10 copy number.

Statistical analysis

The effects of dietary calcium propionate supplementation on growth performance, meat quality, and gene expression in the *longissimus* muscle of finishing steers fed a high-energy diet were evaluated using ANOVA of the generalized linear model (GLM) procedures of SAS 9.0 (SAS Institute Inc., 2002). The differences between groups were analysed in terms of least square group means using a t-test. Differences were considered significant at $P < 0.05$, and not significant at $P > 0.1$.

Results and discussion

Effect of dietary calcium propionate supplementation on performance

The results of the present study show that dietary calcium propionate supplementation did not affect dry matter intake (DMI), average daily weight gain (ADG), or feed conversion ratio (FCR) during the experimental periods ($P > 0.05$; Table 2). In the rumen, calcium propionate dissociates into calcium and propionate ions; the latter is a precursor for glucose synthesis in the liver (Aiello et al., 1989). Propionate participates in feed intake by regulating the liver neural system (Anil and Forbes, 1988). However, the results of current studies indicate that calcium propionate supplementation has inconsistent effects on dry matter intake. Calcium propionate supplementation (200 g · day⁻¹) did not affect feed intake in the present study (Table 2), which is consistent with the results in dairy cattle (Beem, 2003; DeFrain et al., 2004; Liu et al., 2009b) and lambs (Lee-Rangel et al., 2012). However, many inconsistent effects of calcium propionate supplementation on dry matter intake have also been reported. McNamara and Valdez (2005) stated that calcium

Table 2. Effect of dietary calcium propionate supplementation on growth performance

Indices	Treatments		SEM	P
	CG	CaP		
Initial body weight, kg	634.2	637.1	22.36	0.93
Final body weight, kg	676.3	682.8	24.18	0.85
DMI, kg · day ⁻¹	9.26	9.56	0.24	0.90
ADG, kg · day ⁻¹	0.82	0.90	0.09	0.58
FCR	11.29	10.62	1.33	0.61

CaP – experimental group, CG – control group, DMI – dry matter intake, ADG – average daily gain, FCR – feed conversion ratio

propionate supplementation ($125 \text{ g} \cdot \text{day}^{-1}$) increased the DM intake of dairy cattle by 11% prepartum and by 13% post partum. Similarly, Villalba et al. (1996) found that supplementing calcium propionate at $8.3 \text{ g} \cdot \text{day}^{-1}$ increased the feed intake of lambs fed poor quality straw, whereas supplementation at $16.6 \text{ g} \cdot \text{day}^{-1}$ decreased feed intake. Nevertheless, some authors also observed that calcium propionate supplementation depressed feed intake (Rigout et al., 2003; Bradford and Allen, 2007). Therefore, we can suppose that inconsistent effects of calcium propionate supplementation on dry matter intake may depend on the nutrition level of basic diets and the doses of calcium propionate supplementation.

The results of the present investigation showed that calcium propionate supplementation did not affect average daily weight gain (ADG) during the experimental periods ($P > 0.05$). Similarity, Lee-Rangel et al. (2012) also found that supplementing $10 \text{ g} \cdot \text{kg}^{-1}$ calcium propionate to lambs fed 55% and 65% concentrate did not affect the ADG. Whitney et al. (2000) observed a greater ADG in calves that was associated with increased ruminal propionate production. The propionate concentrate was not significantly different between the treatments (CG: 20.84 vs CaP 21.06 $\text{mmol} \cdot \text{l}^{-1}$; $P = 0.91$; data is not showed in manuscript). Therefore, no effect of calcium propionate on ADG in the present study was expected because the high concentrate diets supply so much ruminal propionate that adding Ca propionate does not significantly increase ruminal propionate concentration.

Effect of dietary calcium propionate supplementation on meat quality

As shown in Table 3, calcium propionate supplementation did not affect the variables related to meat quality ($P > 0.05$). Supplementation of this compound to a high-energy diet for finishing steers significantly decreased ($P < 0.05$) the erucic acid content and led to a decreasing tendency in the PUFA content ($0.05 < P < 0.1$) compared with that of the control group (Table 4). No significant difference was found in the other indicators of meat fatty acid composition between the treatments ($P > 0.05$; Table 4).

Previous studies reported lower PUFA contents in grain-fed beef compared with grass-fed beef (Yang et al., 2002; Noci et al., 2005). Jiang et al. (2013) also found that the PUFA content of beef fed a finishing diet at 70% concentrate was higher than those fed at 85% concentrate. These results are probably caused by differences between treatments in PUFA biohydrogenation at the ruminal level.

Table 3. Effect of dietary calcium propionate supplementation on meat quality

Indices	Treatments		SEM	P
	CG	CaP		
pH	5.12	5.12	0.31	0.85
Drip loss, %	13.98	14.14	0.59	0.85
Cooking loss, %	25.31	25.91	1.21	0.72
Warner–Bratzler shear, WBS, kg	4.18	4.82	0.48	0.35
Water, %	69.11	68.82	1.4	0.89
Protein/DM, %	61.06	60.67	2.541	0.91
Fat/DM, %	34.84	36.41	2.697	0.68
Ash/DM, %	3.00	2.95	0.155	0.82

CaP, CG – see Table 2

Table 4. Effect of dietary calcium propionate supplementation on meat fatty acids composition

Fatty acid, $\text{mg} \cdot \text{g}^{-1}$	Treatments		SEM	P
	CG	CaP		
C12:0 Lauric	0.16	0.15	0.0208	0.74
C14:0 Myristic	7.30	7.53	0.784	0.84
C14:1 Myristoleic	1.98	2.15	0.3371	0.73
C15:0 Pentadecanoic	0.77	0.74	0.0683	0.74
C16:0 Palmitic	69.22	67.93	5.1565	0.86
C16:1 Palmitoleic	10.32	9.99	1.0002	0.82
C17:0 Heptadecanoic	2.02	1.94	0.1664	0.76
C17:1 Heptadecenoic	1.41	1.36	0.0985	0.76
C18:0 Stearic acid	33.83	30.12	2.7171	0.35
C18:1t <i>trans</i> -Octadecenoic	108.05	106.16	8.3472	0.87
C18:1c Oleic	108.33	114.53	10.1621	0.66
C18:2t <i>trans</i> -Linoleic acid	0.20	0.16	0.0396	0.51
C18:2c Linoleic	4.44	4.03	0.2074	0.18
C20:0 Arachidic	0.24	0.22	0.0209	0.50
C20:1n9 Eicosenoic	0.49	0.46	0.0419	0.63
C18:3n3 α -Linolenic	0.30	0.21	0.0396	0.14
C22:1n9 Erucic	0.35	0.30	0.0157	0.03
C22:2 Docosadienoic	0.52	0.48	0.031	0.40
TFA	348.47	349.93	22.6210	0.65
SFA	113.55	108.63	8.2144	0.68
MUFA	230.93	234.96	16.1964	0.99
PUFA	5.45	4.88	0.2171	0.06
Omega-3	0.30	0.21	0.0396	0.14
Omega-6	4.64	4.19	0.2186	0.15
Omega-9	0.84	0.76	0.0414	0.20

CaP, CG – see Table 2; TFA – total fatty acids, SFA – saturated fatty acids, MUFA – monounsaturated fatty acids, PUFA – polyunsaturated fatty acids

Neat et al. (1981) reported that the development of the increased capacity for chain shortening and oxidation of erucic acid in the liver of rats fed diets high in erucic acid (C22:1) coincides in time with the development of an increased peroxisomal capacity for fatty acid oxidation. Osmundsen et al. (1979) found that incubation of peroxisomes isolated from hepatocytes with erucoyl-CoA leads to the formation of the same shortened products. So the increased mRNA expression of *PPARG* in group CaP in the

present study could be hypothesized to reflect a higher peroxisomal capacity for shortening and oxidation of erucic acid in the CaP group, thus leading to lower erucic acid levels in meat compared with the CG.

Effect of dietary calcium propionate supplementation on the expression of mRNA of adipogenesis genes in muscle

Marbling is an important factor determining beef quality. Duarte et al. (2013) found that increased mRNA expression of *PPARG* and *CEBPA* resulted in the intramuscular fat (IMF) content being greater in Wagyu muscle than in Angus muscle. Transcriptional remodelling leads to the activation of a series of adipocyte-related genes that cause adipocyte differentiation. Transcription factors, such as PPAR γ and C/EBP α , participate in preadipocyte differentiation and interact to generate fully mature adipocytes (Evan and Ormond, 2006). C/EBP α is a transactivator of PPAR γ , and both transcription regulators act together to promote adipogenesis (Wu et al., 1999).

In the present study, calcium propionate supplementation of a high-energy diet significantly increased the mRNA expression of both *PPARG* and *CEBPA* in the experimental group (CaP) compared with the control group, CG ($P < 0.05$; Figure 1). Wan et al. (2009) also found that propionate significantly increased the expression of *PPARG* and *CEBPA* mRNA of cultured fat cells of the bovine *longissimus dorsi* muscle. In addition, Moisés et al. (2014) found that high-starch diets could trigger up-regulation of the PPAR γ signalling network in *longissimus lumborum* muscle tissue. Therefore, adding calcium propionate could trigger upregulation of the *PPARG* and *CEBPA* mRNA expression levels, which could cause long-term activation of adipogenesis.

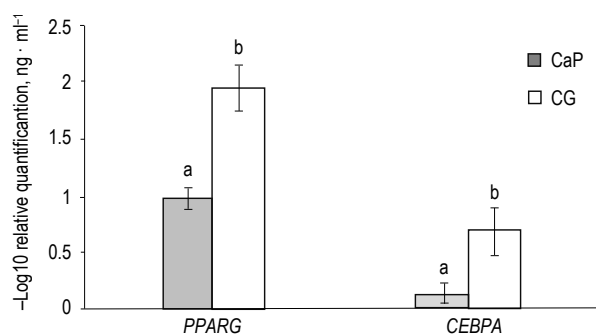


Figure 1. Effect of dietary calcium propionate supplementation on peroxisome proliferator-activated receptor γ (*PPARG*), CCAAT/enhancer binding protein α (*CEBPA*) mRNA relative expression (CaP – experimental group; CG – control group); ^{ab} columns of different groups (CaP, CG) with different letter above differ significantly ($P \leq 0.05$) from each other examined gene, separately

The intramuscular fat content was not, however, significantly different between treatments ($P > 0.05$; Table 3) mainly because the quantity of calcium propionate supplementation and the duration of the experiment were not sufficient to achieve significant effects based on the high-concentrate diets used in the present study. This is in accordance with the study of Mach et al. (2009), who found that supplementing another energy substance, glycerine, to Holstein bulls fed high-concentrate diets also had no effects.

Conclusions

The results indicated that calcium propionate supplementation could improve beef quality though increasing the mRNA expression of adipogenic genes.

Acknowledgements

This study was supported by grants from the National Natural Science Foundation of China (No. 31372335) and the Earmarked Fund for Modern Agro-Industry Technology Research System (Beef Cattle and Yaks, CARS-38) and The Beijing municipal commission of education project (201202910411136).

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