

Assessment of low amounts of meat and bone meal in the diet of laying hens by using stable isotopes

Determinação de baixa inclusão de FCOB em dietas de poedeiras por isótopos estáveis

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Abstract

The aim of this study was to assess whether the inclusion of low amounts of ingredients such as wheat bran, corn gluten and yeast in the diet of laying hens can interfere with the traceability of meat and bone meal (MBM) in eggs and its fractions (albumen and yolk). We used 256 laying hens distributed across eight treatment groups, which consisted of a diet based on corn-soybean-wheat (CSW) bran meal and other diets that had additions comprising gluten and/or (MBM) and/or yeast. To analyse the isotopic ratios ($^{13}\text{C}/^{12}\text{C}$ and $^{15}\text{N}/^{14}\text{N}$), on the 28th and 56th days, 16 eggs were randomly taken for each treatment (four for each repetition), where eight eggs (two for each repetition) were used to harvest yolk and albumen samples and the remaining eight (two for each repetition) were used to analyse the whole egg. We concluded that detection of 2.0% MBM in the egg and its fractions is possible; however, the results for treatments containing gluten and yeast were not different from those containing MBM. Therefore, the inclusion of 3.0% gluten and/or yeast does not result in different $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ isotopic values in the egg and its fractions compared to the values obtained with the addition of 2.0% MBM in the diet.

Key words: Carbon-13, corn gluten, yeast, nitrogen-15

Resumo

O objetivo foi verificar se a baixa inclusão de outros ingredientes, como farelo de trigo, glúten de milho e levedura na dieta de poedeiras, pode interferir na rastreabilidade da farinha de carne e ossos bovinos (FCOB) em ovos e suas frações (albúmen e gema). Foram utilizadas 256 galinhas poedeiras distribuídas em oito tratamentos, no qual, foram compostos de uma dieta MST a base de milho, farelo de soja e farelo de trigo e as demais dietas foram acrescidos de glúten e/ou (FCOB) e/ou levedura. Para análise da razão isotópica ($^{13}\text{C}/^{12}\text{C}$ e $^{15}\text{N}/^{14}\text{N}$), nos 28^o e 56^o dias foram tomados aleatoriamente 16 ovos por tratamento (quatro por repetição), onde oito ovos (dois por repetição) serviram para colheitas de amostras de gema e albúmen e os outros oito (dois por repetição) para ovo total. Pode-se concluir que,

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a detecção de 2,0% de FCOB no ovo e suas frações é possível, porém, os tratamentos com inclusão de glúten e levedura não foram diferentes dos tratamentos com inclusão da FCOB. Portanto, a inclusão de 3,0% de glúten e ou levedura na ração, não permite a diferenciação dos valores isotópicos de $\delta^{13}\text{C}$ e $\delta^{15}\text{N}$ do ovo e suas frações destes grupos dos que possuem 2,0% FCOB.

Palavras-chave: Carbono-13, glúten de milho, levedura, nitrogênio-15

Introduction

The use of flour containing meat, viscera and feathers as protein sources as a substitute for soy bran in the diet of monogastric animals is very common in the Brazilian poultry industry. This flour is relatively low cost and is a good source of nutrients when well processed (MOURA et al., 1994; PEREIRA et al., 1994; BELLAVER et al., 2001). However, the addition of some ingredients with high protein content may disturb the traceability of animal-derived flour in the feed of broilers as detected by stable isotopes.

According to Bellaver et al. (2001), several important markets such as the European Union and Middle East request that poultry are not fed with animal-derived products, antibiotics or growth promoting factors. Poultry grown under these restrictions are denoted organic, green and alternative (or natural), and, for this reason, the sugar-cane yeast (*Saccharomyces cerevisiae*) is widely used in animal feed due to its high protein content and nutraceutical properties.

According to Gottmann et al. (2008), the addition of high amounts of yeast and wheat bran to the feed of broilers disturbs the traceability of animal-derived flour as measured by stable isotopes. Kanayama et al. (2012) observed that the use of carbon and nitrogen isotope ratios to trace animal-derived flours in broiler carcasses may be affected by the addition of sugar-cane yeast, because, in the case of a 6% supplement, the yeast isotope signature cannot be discriminated from that of flour made from viscera and meat.

Denadai et al. (2011) detected the addition of 1.5% meat and bone meal in the albumen of eggs from poultry fed with this ingredient. In this experiment only corn, soy bran and increasing levels of MBM were used as the main components of the diets. This was to prevent any interference from the addition of more ingredients, even though it is known that other components such as wheat bran, gluten, corn gluten and yeast can be added to the diet of laying hens.

Furthermore, to assess whether the addition of other ingredients affects the detection of MBM, Denadai et al. (2009) who used 4.5% meat flour with the addition of 3.0% wheat bran, broken rice, cotton bran, corn gluten, sugar-cane and brewers yeast concluded that the use of stable carbon and nitrogen isotopes allows their detection in eggs and its fractions (yolk and albumen) obtained from laying hens fed with an MBM-containing diet, even after the addition of other vegetal components and yeast. Therefore, the aim of this work was to assess whether the addition of other ingredients such as wheat bran, corn gluten and yeast can affect the traceability of a lower amount (2.0%) of MBM in the diet of laying hens.

Material and Methods

The experiment was performed at the Aviculture Sector of the Brotas Research and Development Unit, Department of Development Decentralization, São Paulo's Agency for Agribusiness Technology, Secretariat for Agriculture and Supply of the State of São Paulo.

Twenty-seven-week-old poultry (Bovans White strain) purchased from the Kakimoto aviary in the city of Alambari/SP (Itapetininga area) were subjected to an acclimatisation period of 28 days.

The aviary consisted of 40 metallic cages, arranged in two double rows, and a 4.0 m wide, 12.0 m long service corridor specific for egg production, with galvanized wire screens along the sides equipped with curtains, closed gables and a French tile roof with clerestories.

The cages measured 1.00 × 0.45 × 0.40 m (width × depth × height) and consisted of two internal compartments, with four hens in each, for a total of eight hens per cage. Each cage was equipped with a Plasson drinking water dispenser and a gutter feeder was placed in front of the cage. Both water and feed were available *ad libitum*, and the artificial lighting program was set as 17 hours light/day.

We used 256 laying hens randomly distributed among eight treatments, with four repetitions, considering one cage containing eight hens as one repetition.

The treatments consisted of a controlled diet based on corn-soybean-wheat bran meal, and the other diets added consisted of corn gluten and/or MBM and/or yeast according to the following specifications:

CSW: Feed based on corn-soybean-wheat;

GLUTEN: Feed based on corn-soybean-wheat + corn gluten;

MBM: Feed based on corn-soybean-wheat + 2.0% MBM;

GLUTEN + MBM: Feed based on corn-soybean-wheat + corn gluten + 2.0% FCOB;

YEAST: Feed based on corn-soybean-wheat + yeast;

YEAST + MBM: Feed based on corn-soybean-wheat + yeast + 2.0% MBM;

GLUTEN + YEAST: Feed based on corn-soybean-wheat + corn gluten + yeast;

GLUTEN + YEAST + MBM: Feed based on corn-soybean-wheat + corn gluten + yeast + 2.0% MBM.

The additions of corn gluten (3.0%) and yeast (3.0%) were determined following consultations with farmers who currently use these ingredients. Thus, the percentages of these ingredients in the feed were as close as possible to the percentages normally used in the egg industry (Brunelli et al., 2012). The amount of MBM (2.0%) added was slightly higher than the minimum detectable amount, which according to Denadai et al. (2011) is 1.5%.

The protein, calcium, phosphorus, metabolisable energy and amino acid content of the ingredients used to prepare the experimental diets were estimated according to the methods of Rostagno et al. (2005). The isotope values of these ingredients were analysed in an isotope-ratio mass spectrometer (Table 1).

Table 1. Average $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ values with the respective standard deviation ($n = 2$) of the ingredients used in the experimental diets.

Ingredients	$\delta^{13}\text{C}$	$\delta^{15}\text{N}$
Corn (1-28 days)	-12.33 ± 0.04	4.56 ± 0.18
Corn (29-56 days)	-12.30 ± 0.02	5.50 ± 0.02
Soybean, meal - 45% PC (1-28 days)	-25.83 ± 0.01	0.59 ± 0.12
Soybean, meal - 45% PC (29-56 days)	-27.75 ± 0.04	0.65 ± 0.22
Wheat, meal (1-28 days)	-27.74 ± 0.12	2.95 ± 0.15
Wheat, meal (29-56 days)	-26.78 ± 0.04	3.40 ± 0.15
Meat, meal - 40% PC	-12.94 ± 0.00	7.33 ± 0.16
Corn gluten	-14.46 ± 0.01	4.03 ± 0.09
Sugarcane yeast	-10.16 ± 0.02	4.49 ± 0.02

Two distinct lots of corn-soybean-wheat bran meals were purchased so that the experimental conditions were as close as possible to those in the aviary, that is, continuous delivery of ingredients obtained from several producers, likely leading to variation in the isotope signature, especially for the $\delta^{15}\text{N}$ (‰) values, due to fertilization and soil conditions. Two rounds of each diet were supplied, one to feed hens between the 1st and 28th days with

the first lot of corn-soybean-wheat bran meal, and a second round with the second lot of ingredients between the 29th and 56th days.

The diets (Table 2) were formulated to meet the nutritional requirements of poultry according to Rostagno et al. (2005) and balanced so that the contents were isoenergetic, isoproteic, isocalcic, isophosphoric and isoaminoacidic for methionine + cystine and lysine.

Table 2. Percentage of the ingredients used and calculated nutritional levels of the experimental diets.

Ingredients	Diets (%)						
	MST	Gluten	FCOB	Gluten + FCOB	Yeast	Yeast + FCOB	Gluten + Yeast + FCOB
Corn, grain	58.87	61.33	60.28	59.61	58.19	60.99	60.64
Soybean, meal (45%PC)	27.35	22.93	25.14	22.83	25.06	18.60	20.63
Wheat, meal	1.00	1.00	1.00	1.00	1.00	1.00	1.00
Meat, meal (45%PC)	0.00	0.00	2.00	2.00	0.00	2.00	0.00
Corn gluten	0.00	3.00	0.00	0.00	0.00	3.00	3.00
Sugarcane yeast	0.00	0.00	0.00	3.00	3.00	3.00	3.00
Soybean, oil	1.90	0.84	1.37	1.37	1.90	0.67	0.84
Limestone	8.70	8.72	8.54	8.55	8.70	8.56	8.72
Dicalcium phosphate	1.53	1.55	1.02	1.00	1.51	1.02	1.54
DL-Methionine	0.15	0.13	0.15	0.14	0.14	0.13	0.13
L-Lysine	0.00	0.00	0.00	0.00	0.00	0.00	0.00
Choline chloride	0.05	0.05	0.05	0.05	0.05	0.05	0.05
Salt	0.30	0.30	0.30	0.30	0.30	0.30	0.30
Vitamin-supplement ¹	0.10	0.10	0.10	0.10	0.10	0.10	0.10
Mineral-supplement ²	0.05	0.05	0.05	0.05	0.05	0.05	0.05
Caulim	0.00	0.00	0.00	0.00	0.00	0.53	0.00
TOTAL	100.00	100.00	100.00	100.00	100.00	100.00	100.00
Calculated Nutritional Levels							
Metabolizable energy (kcal/kg)	2799.95	2800.14	2800.08	2800.00	2799.99	2800.10	2800.10
Crude protein (%)	17.50	17.50	17.51	17.50	17.50	17.50	17.50
Calcium (%)	3.80	3.80	3.80	3.80	3.80	3.80	3.80
Available phosphorus (%)	0.38	0.38	0.38	0.38	0.38	0.38	0.38
Crude fiber (%)	2.59	2.43	2.50	2.38	2.47	2.21	2.21
Methionine (%)	0.43	0.42	0.43	0.42	0.42	0.43	0.43
Methionine + Cystine (%)	0.71	0.72	0.71	0.70	0.70	0.71	0.71
Lisina (%)	0.91	0.82	0.89	0.92	0.93	0.83	0.83

¹- Vitamin supplement per kg feed: Vitamin A: 7.000 UI; Vitamin D3: 2.000 UI; Vitamin E: 5 mg; Vitamin K3: 1.6 mg; Vitamin B2: 3 mg; Vitamin B12: 8 µg; Calcium pantothenate: 5 mg; Niacin: 20 mg; Antioxidant: 15 mg; ²- Mineral supplement per kg feed: Copper: 8 mg; Iron: 50 mg; Iodine: 1.2 mg; Manganese: 70 mg; Zinc: 50 mg; Selenium: 0.2 mg.

Immediately before the experiment was initiated, 16 eggs were sampled, of which eight were used to measure the average isotope signature of the egg ($\delta^{13}\text{C} = -17.97 \pm 0.20$ and $\delta^{15}\text{N} = 5.21 \pm 0.08$) and eight for the isotope signatures of the yolk ($\delta^{13}\text{C} = -17.99 \pm 0.11$ and $\delta^{15}\text{N} = 4.87 \pm 0.15$) and albumen ($\delta^{13}\text{C} = -17.96 \pm 0.10$ and $\delta^{15}\text{N} = 5.28 \pm 0.12$).

The laying hens received the experimental diet for 56 days, to substitute the carbon and nitrogen in the tissues responsible for egg production with those of the new diet. On days 28 and 56^t, 16 eggs per treatment (four for each repetition) were randomly taken, of which eight eggs (two per repetition) were used to harvest yolk and albumen samples, and the remaining eight (two per repetition) were used for total egg sampling. The purpose of performing two samplings (one on day 28 and the other on day 56) was to determine the best day of isotope substitution in the eggs.

The yolks were separated manually from the albumen and the total egg samples (yolk + albumen) were homogenised using a mixer. Once properly identified, the samples were placed in plastic bags and frozen at -20°C for subsequent processing.

The samples were prepared at the Stable Isotope Centre of the UNESP Biosciences Institute, Botucatu Campus.

The egg, yolk and albumen samples were dried in a ventilation incubator at 56°C for 48 h. Then,

$$\delta_{(\text{sample, standard})} = [(R_{\text{sample}} - R_{\text{standard}})/R_{\text{standard}}] \times 1000 \quad (1)$$

Where R represents the ratio between the less abundant and more abundant isotopes, in particular $^{13}\text{C}:^{12}\text{C}$ and $^{15}\text{N}:^{14}\text{N}$. Dimensionless.

The isotope data obtained were subjected to multivariate analysis of variance (MANOVA) and principal component analysis (PCA) with the statistics software Minitab 16 (2010), as confirmed by Sartori et al. (2012) who showed the need to analyse the isotope pair.

egg and yolk samples were subjected to lipid extraction in ethylic ether in a Soxhlet instrument for 4 h. All samples were ground in a cryogenic mill (*Spex-modo 6750 freezer/mill*, Metuchen, EUA) at -196°C for 5 min, with rotation at 920 rpm to obtain a homogenous material, was a fine powder.

To determine the carbon and nitrogen isotope ratios of the samples (albumen, egg and yolk), 50-60 μg material was weighed inside tin capsules on a high-sensitivity analytical scale to ascertain the $\delta^{13}\text{C}$, and 500-600 μg for the $\delta^{15}\text{N}$ measurement.

For the feed, the utilized amount of material was 60-70 μg and 2100-2200 μg , respectively. To determine the isotope composition of more naturally abundant samples, we used an isotope-ratio mass spectrometer (*Delta V Advantage Isotope Ratio MS*, Thermo Scientific, Germany) coupled to an Elemental Analyser (*Flash 2000 Organic Elemental Analyzer EA for IRMS*), in which, in the presence of oxygen (O_2) and copper oxide (CuO), the sample is quantitatively burned to obtain CO_2 . The gases produced were then separated on a gas chromatography column and analysed using a mass spectrometer.

The isotope ratio values were expressed as the delta (δ) value relative to the Pee Dee Belemnite (PDB) international standards for ^{13}C and to atmospheric nitrogen for ^{15}N , according to the following general equation (1):

Isotopic fractionation was calculated according to Hobson and Clark (1992) by using the difference between the δ of the tissue and the diet (equation 2), both for $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$:

$$\Delta = \delta_{\text{tissue}} - \delta_{\text{diet}} \quad (2)$$

The negative value in the unit delta (Δ) indicates that the tissue is low in ^{13}C compared to the diet (Hobson and Clark, 1992).

The calculated fractionations were subjected to MANOVA using the statistics package of the Minitab 16 (2010) software to verify the differences in isotope incorporation between the sampling days (28th and 56th days).

Results and Discussion

The results of the isotope analysis ($\delta^{13}\text{C}$ and $\delta^{15}\text{N}$) of the feed utilised in this study are shown in Table 3. It was possible to observe a difference in the values of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ between the diets formulated for feed in the periods ranging from 1 to 28 and 29 to 56 experimental days, resulting from isotope variation in the feed ingredients (Table 1), since two food lots were used. This difference between evaluation days was also seen in eggs and their fractions (Table 3), showing that, isotopically, the animal is consuming, with a variation of around 2‰ for ^{13}C and 3‰ for ^{15}N (DeNIRO; EPSTEIN, 1978).

Furthermore, due to the inclusion of corn-soybean-wheat (CSW), enrichment in the ^{13}C and ^{15}N isotope values in the diets, as well as in the eggs and fractions, was seen. This observation is consistent with other experiments in poultry (ARAUJO et al., 2011; OLIVEIRA et al., 2010; DENADAI et al., 2009; DENADAI et al., 2008; GOTTMANN et al., 2008; MÓRI et al., 2008; CARRIJO et al., 2006).

In the same way that the $\delta^{13}\text{C}$ value varies, $\delta^{15}\text{N}$ in the feed also varies according to the percentage composition of the ingredients of the diet. The isotope signature of nitrogen in plants

varies depending on how it is fixed. The soybean has a $\delta^{15}\text{N}$ value close to the standard value of atmospheric N_2 ($\delta^{15}\text{N} \cong 0.0 \pm 1.0\text{‰}$), as they fix nitrogen from the air and have a $\delta^{15}\text{N}$ fractionation value around one unit, being usually below 1.003‰ (KOHL; SHEARER, 1980; HANDLEY; RAVEN, 1992; WERNER; SCHMIDT, 2002). On the other hand, the $\delta^{15}\text{N}$ value in plants that are not able to fix atmospheric N_2 , which is the case for corn, greatly depend on the abundance of this isotope in the soil, as well as that added to fertilizers (CHOI et al., 2002).

The isotope fractionation factor for tissue - diet was calculated in an attempt to find the best day, after changing the diet, to collect the samples for traceability analysis. The isotope fractionation factor of the eggs and their fractions (Table 4) was different between the collection days (1-28 days and 29-56 days). The difference may have been caused by changing the diet on day 28 and not by the lack of time to allow a new isotopic signature, since Denadai et al. (2006) analysed the isotopic turnover in 120-week-old laying hens and found that the values of total isotopic exchange of $\delta^{13}\text{C}$ varied between 29 and 35 days, depending on the treatment.

As the analysed tissues had low isotope fractionation, they were a good reflection of the food consumed. Hobson and Clark (1992) determined the fractionation tissue - diet of the egg components from different bird species and found them to vary by -1.0 to 0.2‰ in the lipid-free yolk and between 0.8 and 1.6‰ in the albumen, results that are similar to those found in the present study.

Table 3. Average $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ isotope values (average + SD) of diets (n = 2), eggs, yolks, and albumens (n = 8).

Treatments	Diet		Eggs		Yolks		Albumens	
	$\delta^{13}\text{C}$	$\delta^{15}\text{N}$	$\delta^{13}\text{C}, \%$	$\delta^{15}\text{N}, \%$	$\delta^{13}\text{C}, \%$	$\delta^{15}\text{N}, \%$	$\delta^{13}\text{C}, \%$	$\delta^{15}\text{N}, \%$
	28 days							
MST	-17.02 ± 0.03	1.91 ± 0.17	-18.59 ± 0.13	5.05 ± 0.27	-18.96 ± 0.14	4.89 ± 0.18	-18.24 ± 0.10	5.06 ± 0.09
GLUTEN	-15.93 ± 0.11	2.69 ± 0.19	-17.19 ± 0.11	4.78 ± 0.13	-17.54 ± 0.19	4.99 ± 0.15	-17.08 ± 0.12	5.35 ± 0.19
YEAST	-16.27 ± 0.07	2.44 ± 0.08	-17.86 ± 0.13	5.05 ± 0.13	-18.57 ± 0.14	5.82 ± 0.07	-17.82 ± 0.16	5.28 ± 0.12
GLUTEN + YEAST	-16.40 ± 0.18	2.75 ± 0.06	-16.68 ± 0.16	5.17 ± 0.13	-16.10 ± 0.19	4.91 ± 0.16	-16.87 ± 0.12	5.58 ± 0.12
FCOB	-16.50 ± 0.10	2.17 ± 0.03	-18.25 ± 0.12	4.84 ± 0.19	-18.47 ± 0.17	5.44 ± 0.18	-17.98 ± 0.09	5.40 ± 0.09
GLUTEN + FCOB	-15.25 ± 0.03	2.76 ± 0.03	-16.86 ± 0.11	5.63 ± 0.21	-17.31 ± 0.09	5.07 ± 0.18	-16.95 ± 0.10	5.56 ± 0.15
YEAST + FCOB	-15.86 ± 0.09	2.72 ± 0.24	-17.23 ± 0.14	5.24 ± 0.18	-18.09 ± 0.12	5.34 ± 0.08	-17.43 ± 0.11	5.49 ± 0.08
GLUTEN + YEAST + FCOB	-15.63 ± 0.04	2.82 ± 0.06	-15.99 ± 0.22	5.63 ± 0.19	-17.33 ± 0.19	5.66 ± 0.14	-16.34 ± 0.10	5.42 ± 0.14
	56 days							
CONTROL	-17.70 ± 0.11	1.96 ± 0.13	-18.68 ± 0.11	4.60 ± 0.09	-19.26 ± 0.16	5.45 ± 0.10	-18.30 ± 0.21	4.33 ± 0.12
GLUTEN	-16.34 ± 0.09	2.47 ± 0.13	-17.37 ± 0.15	5.17 ± 0.18	-17.91 ± 0.12	5.77 ± 0.12	-17.06 ± 0.13	4.80 ± 0.10
YEAST	-16.53 ± 0.05	2.34 ± 0.04	-18.31 ± 0.11	4.99 ± 0.09	-18.63 ± 0.12	5.75 ± 0.14	-18.10 ± 0.13	4.53 ± 0.08
GLUTEN + YEAST	-15.97 ± 0.15	2.47 ± 0.11	-17.31 ± 0.17	5.20 ± 0.09	-17.58 ± 0.18	5.95 ± 0.21	-16.99 ± 0.15	4.87 ± 0.12
FCOB	-16.74 ± 0.15	2.62 ± 0.15	-18.04 ± 0.12	5.14 ± 0.12	-18.86 ± 0.16	5.97 ± 0.09	-18.34 ± 0.14	4.92 ± 0.09
GLUTEN + FCOB	-16.01 ± 0.03	2.90 ± 0.08	-16.86 ± 0.18	5.52 ± 0.14	-17.58 ± 0.13	6.15 ± 0.05	-16.84 ± 0.21	5.09 ± 0.08
YEAST + FCOB	-15.95 ± 0.04	3.23 ± 0.08	-17.98 ± 0.24	5.23 ± 0.09	-18.10 ± 0.13	6.17 ± 0.23	-17.61 ± 0.14	4.99 ± 0.15
GLUTEN + YEAST + FCOB	-15.32 ± 0.16	3.06 ± 0.03	-16.65 ± 0.19	5.51 ± 0.20	-17.00 ± 0.18	6.96 ± 0.11	-16.42 ± 0.10	5.05 ± 0.19

Table 4. Isotope fractionation factor between eggs, yolk, and albumen and diet, in laying hens subjected to experimental diets for 1–28 days and 29–56 days.

Treatments	Fracionamento (Δ)					
	Egg - diet		Yolk - diet		Albumen - diet	
	^{13}C	^{15}N	^{13}C	^{15}N	^{13}C	^{15}N
	28 days					
MST	-1.57	3.14	-1.94	2.98	-1.22	3.15
GLUTEN	-1.27	2.09	-1.62	2.3	-1.15	2.66
YEAST	-1.59	2.61	-2.30	2.47	-1.55	2.84
GLUTEN + YEAST	-0.28	2.42	0.29	3.24	-0.47	2.84
FCOB	-1.75	2.67	-1.97	2.90	-1.48	3.23
GLUTEN + FCOB	-1.61	2.87	-2.06	2.58	-1.70	2.80
YEAST + FCOB	-1.37	2.52	-2.24	2.43	-1.57	2.77
GLUTEN + YEAST + FCOB	-0.37	2.81	-1.7	3.29	-0.72	2.6
	56 days					
MST	-0.53	2.65	-1.12	3.49	-0.15	2.37
GLUTEN	-0.99	2.70	-1.53	3.31	-0.68	2.34
YEAST	-1.78	2.65	-2.10	3.42	-1.57	2.19
GLUTEN + YEAST	-1.34	2.73	-1.61	3.48	-1.02	2.40
FCOB	-1.30	2.52	-2.12	3.35	-1.60	2.30
GLUTEN + FCOB	-0.85	2.62	-1.58	3.25	-0.83	2.19
YEAST + FCOB	-2.04	2.00	-2.15	2.94	-1.67	1.76
GLUTEN + YEAST + FCOB	-1.34	2.45	-1.69	3.91	-1.10	2.43

In the egg, the effect of YEAST treatment was significant according to MANOVA at $p < 0.05$, and the effects of the other treatments and fractions were significant at $p < 0.01$.

The average values of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ found in eggs (yolk + albumen), yolk and albumen on day 28 and 56 were statistically analysed and plotted in the graphs shown in Figures 1, 2 and 3, respectively.

The analysis of egg factors on day 28 (Figure 1A) indicated that the CSW treatment was in the same response group as the MBM treatment, and the latter was in the same group as the YEAST treatment. The group supplied with GLUTEN, YEAST + MBM, GLUTEN + MBM, and GLUTEN + YEAST integrated the undefined response groups. However, only the treatment consisting of MBM + GLUTEN + YEAST did not overlap with the other treatments, forming a response group with its own samples. Therefore, in this period, three large response groups were formed.

On day 56 (Figure 1B), only the CSW treatment and its respective samples formed a response

group, without being influenced by other groups. The remaining response groups showed an overlap between the treatments. Consequently, a difference between the groups was observed, thus forming three groups: the CSW group, treatments with addition of MBM and/or yeast (MBM, YEAST, YEAST + MBM, and GLUTEN + YEAST), and the treatments with inclusion of meat and bone meal and/or gluten (GLUTEN, MBM + GLUTEN, and GLUTEN + YEAST + MBM).

The analysis of yolk collected on day 28 (Figure 2A) resulted in the formation of four response groups. The first and second groups were formed by samples from the treatment groups receiving GLUTEN + YEAST + MBM and GLUTEN + YEAST, respectively. The third response group was CSW, YEAST, MBM and YEAST + MBM; the treatments with GLUTEN + MBM and GLUTEN formed the fourth and last response group. It is

possible that the addition of 3% gluten prevented the detection of 2.5% MBM. However, Denadai et al. (2011), when using MBM in the feed of laying hens, were able to detect the addition of 3.0% MBM in the yolk, however a lower level was detected in the albumen (1.5% MBM).

On day 56, the yolk analysis showed more uniform grouping (Figure 2B), in which the treatments CSW and MBM + GLUTEN + YEAST and their respective samples formed response groups with no interference from other treatments. The remaining groups showed the influence of more than one treatment.

For the albumen collected on day 28 (Figure 3A), the treatments CSW and GLUTEN + YEAST + MBM were not influenced by other treatments, forming a response group only with their respective samples. The remaining treatments formed undefined response groups, with significant interference between each other.

However, on day 56 (Figure 3B), CSW and MBM did not interfere with other treatments, forming a response group with their respective samples. The response group that had received YEAST was influenced by CSW treatment; however, GLUTEN treatment overlapped with those including MBM, thus preventing their discrimination. This is consistent with Denadai et al. (2009), who affirmed that the addition of C_4 , corn gluten, and yeast led to higher enrichment in the feed, causing the isotope values of the eggs and their fractions to be even more different from the values for the CSW group.

However, Kanayama et al. (2012) observed that the inclusion of 6% sugar-cane yeast in broiler feed containing corn and soybean might interfere with the traceability of animal-derived flours in the case of carbon- and nitrogen-stable isotopes. The authors

have also discussed studies involving the detection of 4% to 6% yeast inclusion in the feed.

Gottmann et al. (2008) showed that the traceability of animal-derived flours in the pectoral muscle of the broilers, determined by stable isotopes, was disturbed by the presence of yeast and wheat bran.

The analysis of principal components in the albumen fraction, both at 28 and 56 days, led to the formation of more defined response groups, corroborating the findings of Denadai et al. (2008, 2009), who confirmed that albumen comprised the fraction that allowed the best detection of MBM additions to plant-derived components and yeast.

However, both in the egg and in its fractions (albumen and yolk), each group was more defined at 56 days. The discrepancy in the results between the evaluation days suggests that feeding for only 28 days may not be sufficient for the complete integration of the diet isotope signature into eggs. This is not consistent with the results of Carrijo et al. (2000), who concluded that a maximum of 28 days is sufficient to alter the isotope signatures in eggs from laying hens when using two distinct dietary components (C_3 and C_4 grains). Analysing the pectoral muscle of 42-day-old broilers, Gottmann et al. (2008) detected the inclusion of 1.5% viscera flour using discriminating multivariate statistical analysis, while Oliveira et al. (2010) detected the inclusion of 8% viscera flour in the feed of 42-day-old broilers by MANOVA analysis.

The results obtained with the inclusion of 3.0% corn gluten and 3.0% yeast were not different from those obtained with the inclusion of 2.0% meat and bone meal in the egg and its fractions (albumen and yolk), possibly due to the low percentage of the addition (2.0%), even though the percentage used by the industry is between 4.5% and 5.0%.

Figure 1. Graph showing the ^{13}C and ^{15}N principal components of egg after 28 (A) and 56 experimental days (B).

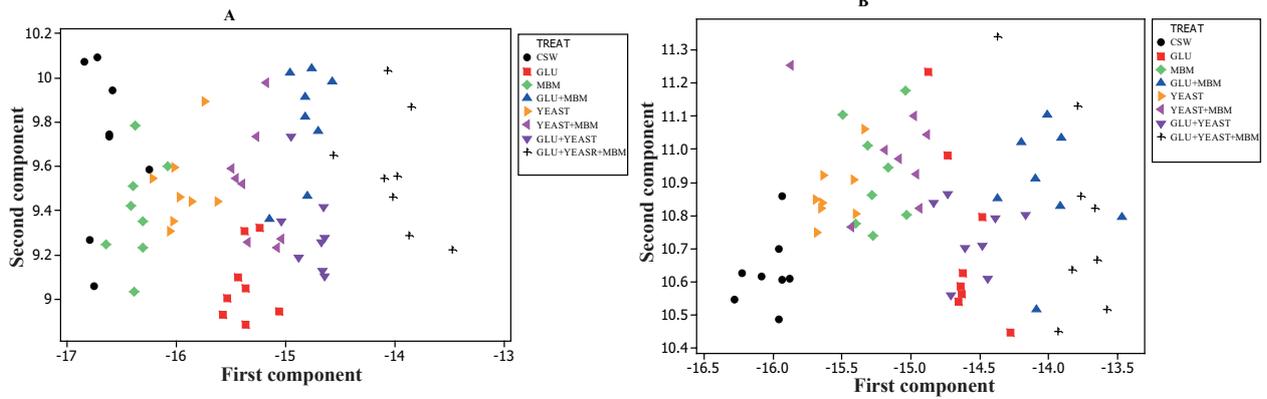


Figure 2. Graph showing the ^{13}C and ^{15}N principal components of yolk after 28 (A) and 56 experimental days (B).

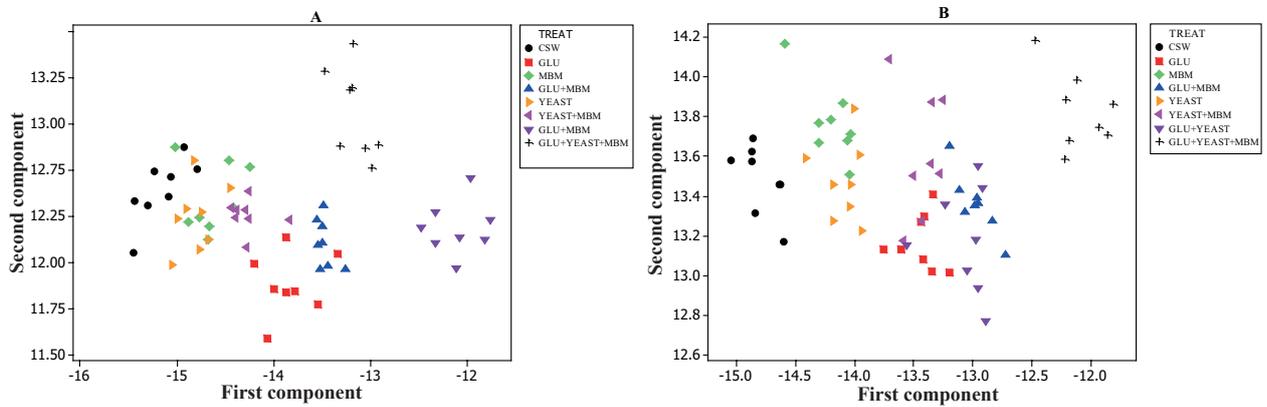
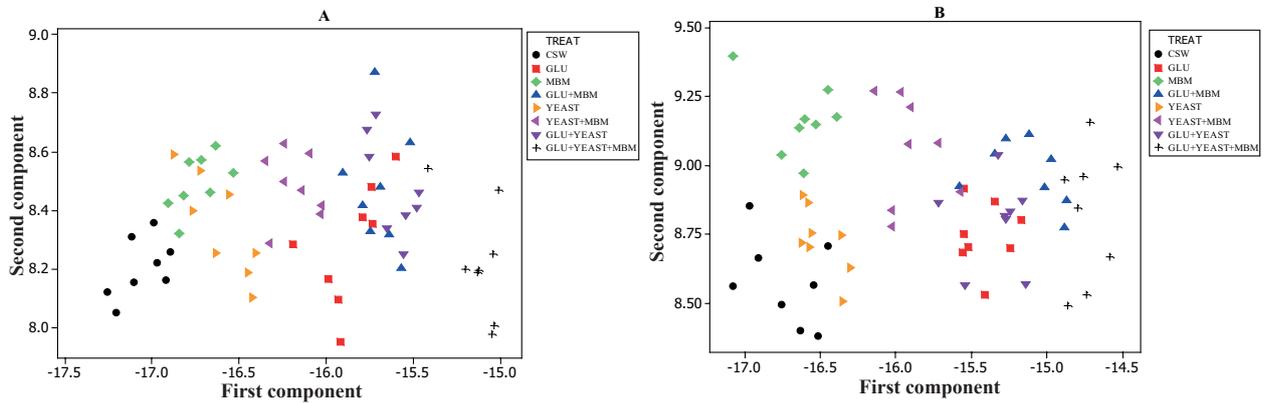


Figure 3. Graph showing the ^{13}C and ^{15}N principal components of albumen after 28 (A) and 56 experimental days (B).



Conclusions

The detection of 2.0% MBM in the egg and its fractions (albumen and yolk) was possible. However, the effect of corn gluten and yeast addition to the diet of laying hens was not different from that obtained with the addition of MBM.

The addition of 3.0% corn gluten and/or yeast to the diet of laying hens did not allow for the differentiation of the isotope values of $\delta^{13}\text{C}$ and $\delta^{15}\text{N}$ in the egg and its fractions (albumen and yolk) for the groups in which 2.0% MBM was included.

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