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Amino acid composition, and determination and prediction of the protein digestibility of different sugarcane yeasts in broilers¹

Composição aminoacídica, determinação e predição da digestibilidade protéica de diferentes leveduras de cana para frangos

Emanuela Nataly Ribeiro Barbosa², Carlos Bôa-Viagem Rabello², Claudia Costa Lopes^{3*}, Edney Pereira Silva⁴
and Ednardo Rodrigues Freitas⁵

ABSTRACT - The aim of this study was to determine the amino acid composition and digestible protein values of different sugarcane yeasts, as well as to generate prediction equations for the apparent digestibility coefficient of dry matter (ADCDM) and crude protein (ADCCP), based on the chemical composition of the yeast and the age of the broilers. Four ileal digestion trials were carried out with 690 broilers, divided into the pre-starter, starter, growth and final phases. The birds were housed in metabolic cages and distributed in a completely randomised design, consisting of five treatments and six replications of eight, six, five and four birds respectively in each experimental phase. The treatments consisted of one reference feed and four test feeds, substituting 30% of the reference feed with four different yeasts. Yeasts two and four showed a higher percentage of amino acids. However, yeast two had the highest percentages of ADCDM, ADCCP and digestible protein: 83.64, 81.22 and 14.66% in the pre-starter phase; 78.09, 78.28 and 14.13% in the starter phase; 76.74, 74.75 and 13.49% in the growth phase; and 72.98, 71.21 and 12.85% in the final phase respectively. There was an average reduction of 0.424% in ADCDM and 0.428% in ADCCP for each day the birds aged. The following prediction equations were generated from the composition of the yeasts and age of the chickens: $ADCDM = 77.4 - 0.424*(age) - 7.51*(DM) - 45.01*(EE)$ and $- 0.424*(age) + 3.02*(CP) + 126.93*(EE) - 0.03*(GMD)$ and $ADCCP = 606.9 - 0.428*(age) - 5.8*(DM) - 0.01*(GMD)$ and $- 0.426*(age) + 2.68*(CP) + 139.7*(EE) - 0.03*(GMD)$.

Key words: Amino acids. Ileal digestibility. Digestible protein. *Saccharomyces cerevisiae*.

RESUMO - Objetivou-se determinar a composição aminoacídica e os valores de proteína digestível de diferentes leveduras de cana-de-açúcar, bem como gerar equações de predição dos coeficientes de digestibilidade aparente da matéria seca (CDAMS) e proteína bruta (CDAPB), com base na composição química da levedura e idade dos frangos de corte. Quatro ensaios de digestibilidade ileal foram realizados com 690 frangos de corte, divididos nas fases pré-inicial, inicial, crescimento e final. As aves foram alojadas em gaiolas de metabolismo e distribuídas num delineamento inteiramente casualizado constituído de cinco tratamentos e seis repetições de: oito, seis, cinco e quatro, respectivamente, para cada fase experimental. Os tratamentos foram: uma ração referência e quatro rações testes com a substituição de 30% da ração referência por quatro diferentes leveduras. As leveduras dois e quatro apresentaram maiores percentagem de aminoácidos. Entretanto, a levedura dois apresentou os maiores percentuais de CDAMS, CDAPB e de proteína digestível, estes sendo: 83,64; 81,22 e 14,66% na fase pré-inicial, 78,09; 78,28; 14,13% na inicial, 76,74; 74,75 e 13,49% na fase de crescimento e 72,98; 71,21 e 12,85% na fase final, respectivamente. Houve redução média de 0,424% no CDAMS e 0,428% no CDAPB, para cada dia de envelhecimento das aves. A partir da composição das leveduras e idade dos frangos, foram geradas as seguintes equações de predição: $CDAMS = 77,4 - 0,424*(idade) - 7,51*(MS) - 45,01*(EE)$ e $- 0,424*(idade) + 3,02*(PB) + 126,93*(EE) - 0,03*(DGM)$ e $CDAPB = 606,9 - 0,428*(idade) - 5,8*(MS) - 0,01*(DGM)$ e $- 0,426*(idade) + 2,68*(PB) + 139,7*(EE) - 0,03*(DGM)$.

Palavras-chave: Aminoácidos. Digestibilidade ileal. Proteína digestível. *Saccharomyces cerevisiae*.

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INTRODUCTION

With an estimated production of 512 thousand tons in the 2013/2014 harvest (ÚNICA, 2015), products of microbial origin, such as yeasts, stand out as promising alternative ingredients in animal feed. According to Freitas *et al.* (2013), these agro-industrial by-products are now being studied, as they are generally ingredients of high biological value (RUTZ *et al.*, 2006), acquired at low cost, and readily found in certain areas of sugar cane production.

Yeast is obtained from the anaerobic fermentation of sugarcane or molasses during the process of alcohol production, collected by the centrifugation of wine or from the bottom of the fermentation vats, and then dried. According to Silva *et al.* (2008), yeast is a viable alternative in the partial replacement of protein components that make up animal feeds, such as soybean meal, due to frequent increases in the price of cereal grain and vegetable sources of protein.

According to the literature, yeast has levels of crude protein that vary from 16.85 to 40.18% (LOPES *et al.*, 2011; LOPES *et al.*, 2015, ROSTAGNO *et al.*, 2011). On the other hand to amino acid composition, Lopes *et al.* (2011) found that yeast is poor in sulphurated amino acids, such as methionine (0.26%) and cystine (0.14%); however, according to the NRC (1994), sugarcane yeast has good levels of lysine (3.23%), leucine (3.19%) and threonine (2.06%). It is a great source of complex B vitamins, carbohydrates and minerals (FLEMMING, 2005).

The availability of prediction equations, or an indirect method to determine digestibility by means of the chemical and physical parameters of feed, is important for increasing the precision of the feed-formulation process, so that it becomes possible to correct the variations inherent in obtaining ingredients (ROSTAGNO *et al.*, 2007), and allow estimates of digestible values based on the variability of feed composition.

Considering the above, the aim of this study was to determine the apparent ileal digestibility coefficients of dry matter and crude protein in different sugarcane yeasts for broilers at four different ages, as well as to generate prediction equations to estimate the values of these coefficients from the chemical and physical composition of the yeasts.

MATERIAL AND METHODS

Four yeast samples were acquired from different companies (Distilleries) in the northeast of the Brazil, and sent to the laboratory of Evonik Industries for amino

acid analysis by high performance liquid chromatography (HPLC). Samples were also sent to the Plant Chemistry and Animal Nutrition laboratories of the Universidade Federal Rural de Pernambuco - UFRPE to determine the chemical composition (Table 1) as per methodologies proposed by AOAC (1990). The geometric mean diameter (GMD) was determined following the methodology described by Zanotto and Bellaver (1996); this parameter of physical composition was analysed, as it makes it possible to correlate the granulometry of the ingredient with nutrient digestibility and animal response.

Four experimental trials were carried out in the Laboratory of Non-Ruminant Digestibility at the Department of Animal Science (UFRPE). For this purpose, 690 male Cobb 500 broilers of uniform weight were used, housed in metabolic cages (1.00 x 0.50 x 0.50 m); for each trial, the birds were distributed in a completely randomised experimental design, consisting of five treatments and six replications.

The experiment was divided into four experimental phases, each lasting eight days. In the pre-starter phase (one to eight days of age), starter phase (ten to 18 days of age), growth phase (20 to 28 days of age) and final phase (30 to 38 days of age), eight, six, five and four birds were used per experimental unit respectively.

During the experiments, the birds received water and feed *ad libitum*. The mean temperature and humidity recorded during the experimental phases were respectively: pre-starter phase 33.2 °C and 70.4%, starter phase 30.28 °C and 70.9%, growth phase 30.8 °C and 71.7%, and in the final phase 31.06 °C and 74.75%.

The treatments consisted of one reference diet (RD), based on maize and soybean meal, and another four test diets, each composed of a mixture of 70% of the RD and 30% of one of the yeasts under study, to give the following diets: Diets A - reference diet; Diets B - 70% reference feed + the addition of 30% Yeast 2; Diets C - 70% reference ration + the addition of 30% Yeast 1; Diets D - 70% reference feed + the addition of 30% Yeast 3; Diets E - 70% reference feed + the addition of 30% Yeast 4.

The reference diets (Table 2) were formulated using the feed-composition tables proposed by Rostagno *et al.* (2011). The birds were raised in a shed and received reference feed, following the recommendations of the COBB 500 management guide (2012); for each experimental period the birds were selected by weight, with a view to achieving maximum uniformity for the start of the experimental phase.

During the last four days of each experimental period, 1% chromium oxide (Cr₂O₃) was added to the feed

as an indicator. At the end of each period, the birds were sacrificed by cervical dislocation to collect the contents of the ileal segment. Two hours before sacrificing the animals, feed intake was stimulated so that more material could be present in the ileum. The ileum was exposed after abdominal incision and a segment, approximately 18 cm long, ending 1.5 cm from the ileo-cecal-colic junction, was clamped and its contents delicately removed with the help of tweezers, placed in plastic tubes and stored in a freezer at -20 °C.

At the end of all the experimental periods, the collected materials were unfrozen, homogenised, placed in properly identified Petri dishes, weighed, refrozen and then vacuum-dried at -54 °C for 48 hours in a lyophiliser.

The pre-dried samples were then ground in a ball mill and sent to the laboratory to determine dry matter, crude protein and chromium, as per the methodology described by AOAC (1990). From results of the analyses, calculations were made to determine the apparent digestibility coefficient of the dry matter (ADCDM), crude protein (ADCCP) and digestible protein (PD), using the following formulae:

(1) Indigestibility Factor:

$$(IF) = \text{Diet indicator} / \text{Digesta indicator} \quad (1)$$

(2) Apparent digestibility coefficient of dry matter:

$$ADCDM (\%) = 100 - (IF * 100) \quad (2)$$

(3) Apparent digestibility coefficient of crude protein:

$$ADCCP (\%) = CP_{\text{diet}} - (CP_{\text{digesta}} * IF) / CP_{\text{diet}} * 100 \quad (3)$$

(4) Digestible protein:

$$DP = CP_{(\text{diet}/\text{feed})} * ADCCP_{(\text{diet}/\text{feed})} / 100 \quad (4)$$

The values determined for the apparent ileal digestibility coefficients of the dry matter (ADCDM), crude protein (ADCCP) and digestible protein (DP) of the feed and the sugarcane yeast in the experimental trials were submitted to analysis of variance, and the mean values compared by Tukey's test at 5% probability.

The chemical and physical composition of the yeasts, i.e. dry matter, crude protein, ether extract, mineral matter, neutral detergent fibre, crude energy and geometric mean diameter, as well as the variable, age of the birds, were also submitted to multiple regression analysis, these values being the independent variables, and the values for ADCDM and ADCCP, the dependent variables, to verify those variables that best correlated with the values for ADCDM and ADCCP, thereby generating prediction equations for the apparent and corrected apparent metabolisable energy for the yeasts by means of the Indirect Elimination Stepwise (Backward) method during the experimental phases.

The coefficients of determination (R^2) and the Akaike information criterion (AIC) (Akaike, 1974) were used to verify the good fit of the prediction equations to the data. The statistical analyses were carried out using the Statistical Analysis Systems 9.1.3 statistical software package (SAS INSTITUTE, 2008).

RESULTS AND DISCUSSION

The values for the amino acid composition of the four sugarcane yeasts under evaluation are shown in Table 3.

Table 1 - Physicochemical and energetic composition of the sugarcane yeasts used in the experimental trial, expressed as dry matter

Nutrient	Yeast			
	1	2	3	4
Dry Matter, %	90.71	88.84	89.99	90.47
Crude Protein, %	17.11	18.05	14.94	21.12
Mineral Matter, %	7.54	8.05	9.1	9.0
Ether Extract, %	0.37	0.44	0.49	0.30
NDF ¹ , %	6.86	7.65	3.02	12.13
Gross Energy, kcal/kg	4.214	4.268	4.011	4.095
GMD ² , µm	661	695	933	616

¹ NDF-Neutral Detergent Fibre; ² GMD - Geometric Mean Diameter *Expressed as dry matter

Table 2 - Percentage composition and nutritional values of the experimental diets

Ingredient	Experimental phase			
	Pre-starter	Starter	Growth	Final
Maize	55.405	56.371	60.283	64.522
Soybean meal 45%	37.470	35.778	31.741	27.704
Soybean oil	2.427	3.680	4.259	4.202
Dicalcium phosphate	1.941	1.835	1.689	1.537
Calcitic Limestone	0.921	0.888	0.842	0.801
Common salt	0.517	0.502	0.477	0.452
L- lysine HCL (78.8%)	0.374	0.203	0.209	0.264
DL- Methionine (99%)	0.193	0.0825	0.244	0.239
L- Threonine (98.5%)	0.148	0.0580	0.0515	0.070
Vitamin-mineral supplement ¹	0.600	0.600	-	-
Vitamin supplement ²	-	-	0.100	0.100
Mineral supplement ³	-	-	0.100	0.100
Total	100.0	100.0	100.0	100.0
Energetic and nutritional composition				
Metabolisable energy, kcal/kg	2,960	3,050	3,150	3,200
Dry matter, %	90.25	90.10	91.53	89.70
Crude protein, %	22.11	21.14	19.73	18.31
Calcium, %	0.942	0.899	0.837	0.775
Available phosphorus, %	0.471	0.449	0.418	0.386
Sodium, %	0.224	0.218	0.208	0.198
Potassium, %	0.840	0.812	0.749	0.687
Crude fibre, %	2.985	2.910	2.760	2.614
Ether extract, %	5.039	6.292	6.887	6.973
Mineral matter, %	2.913	2.825	2.637	2.453
Digestible amino acids, %				
Methionine + cystine	0.968	0.844	0.791	0.755
Methionine	0.677	0.559	0.522	0.500
Lysine	1.363	1.189	1.099	1.048
Threonine	0.886	0.773	0.714	0.681
Tryptophan	0.243	0.234	0.213	0.193

¹Vitamin-mineral supplement. (Guaranteed levels per kg of product). Vit. A: 1500,000 UI, Vit. D3: 500,000 UI, Vit. E: 3,333 mg, Vit K3: 250 mg, Riboflavin: 1,000 mg, Thiamine: 300 mg, Vit B6 (Pyrixine): 500 mg, Vit B12: 3,333 mg, Niacin: 6.667 mg, Calcium Pantothenate: 2,000 mg, Folic Acid: 285 mg, Biotin: 8.33 mg, Choline chloride: 70 mg, Methionine: 295 mg, Ferro: Iron mg, Copper: 1,550 mg, Manganese: 14,000 mg, Zinc: 12,130 mg, Iodine: 280 mg, Selenium: 70 mg, Robenidine: 6,000 mg, Licomicin: 734 mg, Antioxidant Additive: 16,667 mg; ²Vitamin supplement. (Guaranteed levels per kg of product). Vit A 1000000 IU, Vit B3 2000000 IU, Vit E 20000 mg, Vit K3 4000 mg, Vit B1 1880 mg, Vit B2 5000 mg, Vit B6 2000 mg, Vit B12 1000 mg, niacin 30000 mg, Pantothenic 13500 mg, Ac. Folic acid 500 mg, Selenium 250 mg, Antioxidant 100000 mg; ³Mineral supplement. (Guaranteed levels per kg of product). Manganese 75000 mg, Zinc 70000 mg, Iron 60000 mg, Copper 85000 mg, Iodine 1500 mg, Cobalt 200 mg

Table 3 - Amino acid composition of the sugarcane yeasts based on the dry matter

Amino acid, %	Yeast			
	1	2	3	4
Methionine	0.26	0.28	0.19	0.28
Cystine	0.11	0.11	0.08	0.14
Methionine + Cystine	0.38	0.39	0.27	0.43
Lysine	1.25	1.27	0.97	1.40
Threonine	0.97	1.01	0.76	1.07
Arginine	0.76	0.79	0.58	0.97
Isoleucine	0.85	0.88	0.66	0.94
Leucine	1.23	1.28	1.00	1.38
Valine	0.99	1.03	0.82	1.12
Histidine	0.35	0.37	0.28	0.40
Phenylalanine	0.74	0.76	0.58	0.82
Glycine	0.82	0.85	0.65	0.91
Serine	0.89	0.90	0.74	0.99
Proline	0.63	0.68	0.52	0.73
Alanine	1.06	1.12	0.95	1.23
Aspartic acid	1.73	1.78	1.58	1.99
Glutamic acid	1.86	1.95	1.54	2.20

Based on the results seen for the total amino acid composition of the yeasts, yeasts 2 and 4 presented higher values when compared to the others; these results were lower than those found in NRC (1994) for lysine (3.23%), leucine (3.19%) and threonine (2.06%). All the yeasts under study proved to be inferior in amino acid composition when compared to the whole yeast studied by Martins (2009), and the yeast extract evaluated by Silva *et al.* (2009). However, only yeast 3 had a lower amino acid composition than the yeast described by Lopes *et al.* (2011).

The values obtained for the apparent ileal digestibility coefficients of the dry matter (ADCDM,%), crude protein (ADCCP,%) and digestible protein (DP,%) of the sugarcane yeasts under study can be seen in Table 4.

Among the foods tested in the pre-starter phase, yeasts 2, 3 and 4 had a higher percentage of ADCDM (83.64, 80.95 and 82.87 respectively). Yeast 2 had the highest value for ADCCP (81.22%) and yeast 4 had the highest percentage of DP (16.57%).

In the starter period, the highest percentage of ADCDM and ADCCP were for Yeast 2 (78.09 and 78.28% respectively); for DP value, Yeasts 2 and 4 had the highest percentages (14.13 and 14.07%). The same behaviour was seen during the growth and final periods, where Yeast

2 obtained the greatest values for ADCDM (76.74 and 72.98%) and for ADCCP (74.75 and 71.21%). For DP, Yeasts 2 and 4 had better results (13.49 and 13.22%) and (12.85 and 13.07%) respectively in both experimental phases.

According to Lopes *et al.* (2015), the use of whole yeast in chicken diets may reduce nutrient digestibility due to the cell wall, which is thick and resistant to enzymatic digestion, making the intracellular content unavailable.

The highest values seen for DP in Yeasts 2 and 4 in all the experiments can be explained by these yeasts having the highest levels of crude protein (18.05 and 21.12% respectively), and high protein digestibility coefficients, which increases the percentage of DP.

The digestibility coefficients of the dry matter and crude protein, as well as the digestible protein of the sugarcane yeasts under study, decreased with advancing age; this is considered normal, since according to Nery *et al.* (2007), in the early phases of a bird's life, pre-starter and starter, greater nitrogen retention occurs for the deposition of protein tissue.

The multiple regression equations generated to estimate the apparent digestibility coefficients of the dry matter (ADCDM) and the apparent digestibility coefficients of the crude protein (ADCCP) of the yeasts

Table 4 - Mean value for the apparent ileal digestibility coefficients of the dry matter and crude protein, and values for digestible protein of the sugarcane yeasts based on the dry matter

Parameter	Yeast				Mean	CV	P
	1	2	3	4			
ADCDM (%)	74.21 b	83.64 a	80.95 a	82.87 a	80.42	1.32	0.0029
ADCCP (%)	72.59 b	81.22 a	76.74 ab	78.46 ab	77.25	2.61	0.0524
DP (%)	12.42 c	14.66 b	11.46 c	16.57 a	13.77	2.65	0.001
Starter (18th day)							
ADCDM (%)	65.08 b	78.09 a	65.73 b	67.98 b	69.22	2.97	0.01
ADCCP (%)	66.80 b	78.28 a	68.11 b	66.66 b	69.96	3.73	0.01
DP (%)	11.43 b	14.13 a	10.17 c	14.07 a	12.45	3.75	0.01
Growth (28th day)							
ADCDM (%)	62.24 c	76.74 a	61.42 c	68.83 b	67.31	3.10	0.01
ADCCP (%)	63.09 b	74.75 a	63.28 b	62.61 b	65.93	4.80	0.01
DP (%)	10.79 b	13.49 a	9.45 c	13.22 a	11.74	4.60	0.01
Final (38th day)							
ADCDM (%)	66.60 c	72.98 a	59.82 d	68.04 b	66.89	0.81	0.01
ADCCP (%)	61.40 b	71.21 a	62.64 b	61.89 b	64.28	5.36	0.01
DP (%)	10.50 b	12.85 a	9.35 c	13.07 a	11.44	4.89	0.01

* ADCDM - apparent digestibility coefficient of dry matter; * ADCCP - apparent digestibility coefficient of crude protein; * DP - digestible protein; *CV - coefficient of variation; * P - probability; Mean values followed by the same letter in a row do not differ statistically by Tukey's test at 5% probability

are shown in Tables 5 and 6 respectively. To predict the values for ADCDM and CP, a total of 20 significant models were generated.

To verify the fit of the data, the coefficient of determination (R^2) and the Akaike information criterion (AIC) were used. Some significant models that were generated use the intercept as part of the equation, however, most of the equations did not use the intercept in the model. Janssen (1989), when elaborating the European Table of Energy Values for Poultry, did not take the intercept into consideration, the same as Rostagno *et al.* (2011), when elaborating the Brazilian Tables for Poultry and Swine.

The coefficients of determination of the regression equations (R^2) ranged from 0.73 to 0.99 for the ADCDM, with an AIC of from 270.85 to 272.85; for the ADCCP, the R^2 ranged from 0.789 to 0.998, with an AIC of 234.84 to 236.79, using equations with three to five variables.

For the predicted values of the ADCDM, among the equations generated using the intercept, Model 1 ($ADCDM = 774.4 - 0.424*age - 7.51*DM - 45.01*EE$) gave a better fit to the data, considering the value of the coefficient of determination ($R^2 = 0.73$) and the lowest value for the AIC (270.85). For the equations that did not use the intercept, Model 4 had the best fit ($ADCDM =$

$-0.424*age + 3.02*CP + 126.93*E - 0.03*GMD$) with $R^2 = 0.99$ and an AIC of 270.94.

For the values predicted for the ADCCP comparing the equations that made use of the intercept, Model 1 ($ADCCP = 606.9 - 0.428*age - 5.8*DM - 0.01*GMD$) with an R^2 of 0.789 and an AIC of 234.84 gave the best fit to the data. For the equations that did not use the intercept as part of the model, Equation 6 ($ADCCP = -0.426*age + 2.6*CP + 139.7*EE - 0.03*GMD$) best fit the data, with a value for R^2 of 0.998 and the smallest value for AIC (235.47).

High values for the coefficient of determination (R^2) were obtained with the remaining equations that used or did not use the intercept in the models, however, as the AIC was also used, whose criterion of choice is the lowest value, the models that presented both combinations were those that best fit the data, these being chosen to predict a more accurate value.

In the models generated for the ADCDM and ADCCP, with or without the use of the intercept, the age parameter was included in all equations. It can be seen that for each day that the birds age, the values go down on average by 0.424% for the ADCDM and 0.428% for the ADCCP.

Table 5 - Prediction equations for the apparent digestibility coefficient of dry matter in broiler chickens as a function of the physicochemical composition of the yeasts¹

RE	Coefficient of regression									R ²	AIC
	Inter.	Age	DM	CP	MM	EE	NDF	MGD	CE		
1	774.41	-0.424	-7.51	-	-	-45.01	-	-	-	0.73	270.859
2	581.70	-0.424	-5.63	-	-	-	0.83	-	-	0.73	271.616
3	20.63	-0.424	-	-	-2.14	133.8	3.31	-	-	0.73	272.859
4	-	-0.424	-	3.02	-	126.93	-	-0.03	-	0.99	270.941
5	-	-0.424	-	-11.93	-77.31	-	39.13	0.90	-	0.99	272.859
6	-	-0.424	-	-	-14.43	105.95	7.6	0.14	-	0.99	272.859
7	-	-0.424	1.40	-13.05	-	212.89	13.8	-	-	0.99	272.859
8	-	-0.424	-	-3.0	-	129.18	5.1	-	0.010	0.99	272.859
9	-	-0.424	-	-	-1.47	127.32	3.1	-	0.004	0.99	272.859
10	-	-0.424	0.20	-	-2.20	138.7	3.4	-	-	0.99	272.859

1 - values based on dry matter; Inter. - intercept; RE- regression equation; CP - crude protein; EE - ether extract; MM - Mineral matter; NDF - neutral detergent fibre; MGD - mean geometric diameter; CE - crude energy; R²- coefficient of determination; AIC- Akaike information criterion

Table 6 - Prediction equations for the apparent digestibility coefficient of crude protein in broiler chicken as a function of the physicochemical composition of yeasts¹

RE	Coefficient of regression									R ²	AIC
	Inter.	Age	DM	CP	MM	EE	NDF	MGD	CE		
1	606.9	-0.428	-5.8	-	-	-	-	-0.01	-	0.789	234.85
2	-410.6	-0.428	-	-1.2	12.3	-	-	-	0.1	0.789	236.76
3	-408.2	-0.428	-	-	11.6	-	-0.78	-	0.1	0.789	236.76
4	-312.9	-0.428	-	-	8.52	29.9	-	-	0.1	0.789	236.76
5	-397.5	-0.428	-	-	9.2	-	-	0.02	0.1	0.789	236.79
6	-	-0.426	-	2.6	-	139.7	-	-0.03	-	0.998	235.48
7	-	-0.429	-	3.4	-3.3	115	-	-	-	0.968	235.67
8	-	-0.428	-	-11.5	-78.1	-	38.4	0.90	-	0.998	236.76
9	-	-0.428	1.6	-15.2	-	228.3	14.9	-	-	0.998	236.76
10	-	-0.428	-	-	-17.4	102.3	7.94	0.2	-	0.998	236.76

1 - values based on dry matter; RE - regression equation; CP - crude protein; EE - ether extract; MM - Mineral matter; NDF - neutral detergent fibre; MGD - mean geometric diameter; CE - crude energy; R²- coefficient of determination; AIC- Akaike information criterion

The estimated values for the ADCDM and ADCCP are shown in Tables 7 and 8 respectively, generated using the models that best fit the data for chemical composition. According to Siqueira *et al.* (2011), prediction models can be evaluated by means of bioassays, by simulation, or by the comparison of predicted values with those seen in real situations, constituting a determining step for verifying model applicability.

Comparing the estimated values for the generated equations with the values determined in the biological

trials for the ADCDM and the ADCCP of the sugarcane yeasts under study, it can be seen that the models, with or without use of the intercept, predict values close to those determined. The mean sum of squares is similar between the models.

According to Nascimento *et al.* (2009), the importance of determining prediction equations is a result of the difficulty in carrying out bioassays and the fact that most laboratories do not have a calorimeter, making it possible to work with equations generated from simple chemical analysis.

Table 7 - Estimation of the apparent digestibility coefficient of dry matter by means of the prediction equations, based on the dry matter

Equation with intercept			
$ADCDM_1 = 774.4 - 0.424*(age) - 7.51*(DM) - 45.01*(EE)$			$R^2 = 0.73$
Equation without intercept			
$ADCDM_4 = -0.424*(age) + 3.02*(CP) + 126.93*(EE) - 0.03*(GMD)$			$R^2 = 0.99$
Yeast	$ADCDM^1$	$ADCDM^2$	$ADCDM^2$
1	67.04	66.76	73.04
2	78.08	77.65	76.13
3	66.98	66.76	69.57
4	71.93	71.71	73.62
Mean	71.00	70.72	73.09
SS ³	-	119.8507	119.8507
Mean SS ⁴	-	29.96267	29.96267

¹ADCDM - apparent digestibility coefficient of dry matter, seen *in vivo* in broiler trials; ²- estimates of ADCDM by the prediction equations; ³- sum of squares; ⁴- mean sum of squares

Table 8 - Estimation of the apparent digestibility coefficient of crude protein by means of the prediction equations, based on the dry matter

Equation with intercept			
$ADCCP_1 = 606.9 - 0.428*(age) - 5.8*(DM) - 0.01*(GMD)$			$R^2 = 0.789$
Equation without intercept			
$ADCCP_6 = -0.426*(age) + 2.6*(CP) + 139.7*(EE) - 0.03*(GMD)$			$R^2 = 0.998$
Yeast	$ADCCP^1$	$ADCCP_1^2$	$ADCCP_6^2$
1	65.97	64.32	69.97
2	76.46	74.83	74.63
3	67.69	65.78	69.50
4	67.40	66.17	68.54
Mean	69.38	67.77	70.66
SS3	-	122.1227	120.984
Mean SS4	-	30.53067	30.246

¹ADCCP - apparent digestibility coefficient of crude protein, seen *in vivo* in broiler trials; ²- estimates of ADCCP by the prediction equations; ³- sum of squares; ⁴- mean sum of squares

Under such conditions, it is up to the nutritionist to choose the equation to be used in formulating a feed, since the choice will depend on the laboratory analyses available, considering that some of these analyses require more sophisticated equipment that can make the procedure more costly.

This study pioneers prediction of the apparent digestibility coefficients of dry matter and crude protein

by means of regression equations generated from the chemical composition of sugarcane yeast. The literature is scarce concerning the ileal digestibility of the dry matter and crude protein of sugarcane yeast determined with broilers. This work alone will not be sufficient to determine whether the generated equations can accurately predict the values of the digestibility coefficients of all the yeasts on the market, it is therefore necessary that further studies be carried out to clarify their peculiarities.

CONCLUSIONS

1. The yeasts under study showed variability in their amino acid composition; however, Yeasts 2 and 4 presented a higher percentage of amino acids;
2. The apparent digestibility coefficients of dry matter and crude protein differed between yeasts. For all experimental phases, Yeast 2 presented the highest values for the digestibility coefficient, as well as for the values of digestible protein;
3. It was seen that the ADCDM and ADCCP of the yeasts reduced on average 0.424 and 0.428% respectively for each day the birds aged;
4. Four models, with or without use of the intercept, showed a better fit to the data for physicochemical composition in the equations generated to predict the values of the digestibility coefficients of dry matter and crude protein.

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