Ileal digestibility of amino acids in maize gluten feed for pigs*

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ABSTRACT

Seven batches of maize gluten feed (MGF) were studied for their protein and amino acid standardised ileal digestibility (SID). Each batch was tested on four castrated male pigs Large White (LW) or (LW x Landrace – L) x (LW x Piétrain – P) weighing between 42 and 61 kg, and fitted with an end-to-end ileo-rectal anastomosis. In all batches, lysine, threonine, cystine, tryptophan, aspartic acid and glycine displayed the lowest SID values (on average 62.9, 69.6, 66.7, 64.1, 69.7 and 62.4%, respectively). On the contrary, methionine, arginine, leucine, tyrosine and phenylalanine were among the most digestible amino acids and their average SID reached 84.8, 85.6, 82.3, 83.2 and 82.7%, respectively. Significant differences between batches of MGF appeared for SID of most amino acids. SID of most amino acids decreased as the proportion of N bound to the NDF residue increased.

KEY WORDS: maize gluten feed, ileal digestibility, amino acids, pigs

INTRODUCTION

Maize gluten feed (MGF), by-product of maize starch extraction, is made of the fibrous part of the maize grain and of maize steep liquor, approximately in the proportion of 3 to 1. Although in small proportion, maize germ meal, the chemical

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characterisation of which is close to the one of MGF, is sometimes added to the final product. MGF being the result of a complex industrial process, its composition and its feeding value may be variable. Although this by-product is widely used in pig feeding, only few values relative to its nitrogen (N) and amino acid (AA) ileal digestibility are available in literature (Mariscal-Landin, 1992). The aim of this study, part of a more extensive work on AA ileal digestibility in pig feeds (Jondreville et al., 1995), was to measure the N and AA standardised ileal digestibility in MGF, to evaluate the extent of its variability and to assess the relevance of analytical criteria for its prediction.

MATERIAL AND METHODS

Animals

Castrated male pigs pure bred Large White or (Large White x Landrace) x (Large White x Pietrain) were surgically prepared with an end-to-end ileo-rectal anastomosis at a liveweight of 25 to 30 kg according to the method described by Picard et al. (1984) and Laplace et al. (1989). The experiments lasted from 1991 to 1993 and required the preparation of 16 growing pigs used in trials as four independent groups of four animals. Each batch of MGF was tested on four animals of only one group, at one period, both selected at random (Table 1). Therefore the reliability of our data was based on the initial assumption of the absence of effects, not only of pig ageing or liveweight, but also of other group characteristics on AA ileal digestibility.

| Group | MGF1 | MGF2 | MGF3 | MGF4 | MGF5 | MGF6 | MGF7 |
|-------|------|------|------|------|------|------|------|
| n     | 4    | 4    | 4    | 4    | 3    | 4    | 4    |
| FL    | 79 (3) | 76 (4) | 80 (1) | 79 (4) | 74 (3) | 73 (3) | 79 (3) |
| LW    | 59 (5) | 61 (9) | 59 (8) | 59 (7) | 46 (2) | 42 (2) | 49 (1) |

As described by Green et al. (1987), each test consisted of two phases: an adaptation period of five days followed by a two-day continuous collection period. This combination of the adaptation and collection periods was among those which allowed the most accurate measurement of N excreted by pigs fitted with an end-
to-end ileo-rectal anastomosis (Mariscal-Landin et al., 1995). Twice a day, just after meal, digesta were transferred from cages to a refrigerator and kept at 4°C. At the end of the collection period, the digesta of each pig were homogenised and two samples were taken: one was oven-dried (80°C for 72 h) for dry matter determination and the other one was freeze-dried prior to analysis.

Feedstuffs, experimental diets and feeding

Seven commercial batches of maize gluten feed (MGF1 to MGF7) were selected to be representative for feed used in Europe. MGF6 originated from USA, whereas the other ones originated from European maize starch extraction plants. Each sample of MGF was ground in a hammer mill fitted with a 2 mm screen. Except for MGF4, each of them was mixed with maize starch and sucrose (1:1) and a mineral and vitamin concentrate (55 g/kg diet), so that the N content of the diets was around 31 g/kg DM (Table 2). Because of its low N content, MGF4 was only mixed with the mineral and vitamin concentrate.

Animals were fed twice daily (8 a.m. and 4 p.m.) a diet in meal form mixed with water (1:2, v/v) at the time of feeding. Drinking water was continuously...

Table 2

| Chemical composition of maize gluten and formulation of diets | MGF1 | MGF2 | MGF3 | MGF4 | MGF5 | MGF6 | MGF7 |
|--------------------------------------------------------------|------|------|------|------|------|------|------|
| Chemical composition of MGF, g/kg DM                          |      |      |      |      |      |      |      |
| N                                                            | 35.0 | 36.8 | 39.2 | 31.8 | 37.9 | 34.9 | 38.4 |
| Fat                                                          | 33.1 | 22.5 | 28.2 | 40.9 | 21.2 | 33.5 | 49.5 |
| CF                                                           | 83.2 | 77.8 | 92.7 | 75.0 | 79.3 | 93.9 | 80.6 |
| Ash                                                          | 65.0 | 55.2 | 63.3 | 61.4 | 54.7 | 109.5 | 83.1 |
| NDF                                                          | 388.0 | 378.0 | 413.7 | 389.2 | 304.4 | 384.7 | 366.1 |
| ADF                                                          | 98.4 | 90.4 | 105.6 | 95.0 | 92.8 | 133.9 | 94.0 |
| ADL                                                          | 12.4 | 7.4 | 10.9 | 12.2 | 8.2 | 24.4 | 9.3 |
| N-NDF                                                        | 8.0 | 3.5 | 6.2 | 11.0 | 3.9 | 7.3 | 4.1 |
| Starch                                                        | 256.6 | 235.6 | 188.7 | 355.7 | 232.4 | 154.2 | 202.1 |

Formulation of diets, g/kg DM

|          | MGF | Starch/sucr. | Min./vit. conc. |
|----------|-----|--------------|-----------------|
| MGF      | 886 | 59           | 55              |
| Starch/sucr. | 855 | 90          | 55              |
| Min./vit. conc. | 813 | 132      | 55              |

1 maize starch and sucrose mixture (1:1)

2 the mineral and vitamin concentrate provided per kg DM feed: 303 mg MgCl₂6H₂O, 303 mg KCl, 8437 mg NaCl, 22000 mg NaHCO₃, 9548 mg CaHPO₄, 10849 mg CaCO₃, 512 mg FeSO₄7H₂O, 303 mg ZnSO₄·H₂O, 80 mg CuSO₄·5H₂O, 121 mg MnSO₄·H₂O, 5 mg CoCO₃, 2 mg Ca (IO₃)₂, 27500 IU vit. A, 2200 IU vit. D3, 4070 IU vit. E, 21 mg vit. K3, 6 mg vit. B1, 14 mg vit. B2, 13 mg vit. B6, 51 mg vit. PP, 80 mg vit. C, 1 mg folic acid, 70 mg vit. B12, 0.03 mg biotin, 30 mg vit. B3, 804 mg choline, 50 mg carbadox
available. The feeding scale was set at 90 g meal per kg metabolic weight (live-weight\(^{0.75}\)), i.e., taking into account the dry matter concentration in our maize gluten feed based diets, around 76 g dry matter ingested per kg metabolic weight. Every week, the feed supply was individually adjusted, based on weighings at four-week intervals and expected increase in liveweight between weighings.

**Chemical analysis**

All the chemical determinations were performed in duplicate. Each batch of MGF, each diet and each sample of digesta was analysed for N and AA. The N content was measured by the Kjeldahl method (AFNOR, 1977a), with a Kjel-foss automatic apparatus (Foss-Electric, Denmark). AA were determined on a Beckman 6300 AA analyser after 23 h hydrolysis in HCl 6 N at 110°C (AFNOR, 1993a). Methionine and cystine were determined after oxidation with performic acid. Tryptophan was determined by HPLC followed by fluorometric detection (Kontron Instrument, France). Hydrolysis was carried out in alkaline solution of Ba(OH)\(_2\) at 125°C for 16 h (Landry and Delhaye, 1992).

The dry matter (DM) content of each batch of MGF was determined by drying to constant weight at 103°C (AFNOR, 1976). For the determination of ash content, samples were placed at 550°C for 8 h (AFNOR, 1977b). Neutral detergent fibre (NDF), acid detergent fibre (ADF) and acid detergent lignin (ADL) determinations were sequentially carried out following the van Soest et al. (1991) method, and crude fibre (CF) according to the Weende method (AFNOR, 1993b), by means of a Fibertec system (Tecator, Sweden). The concentration of N in the NDF residue (N-NDF) was evaluated by the successive analysis of NDF and N, as described above. Fat content was measured without hydrolysis prior to analysis (OJEC, 1984), by means of a Soxtherm apparatus (Gerhardt, Germany). According to the Ewers method (OJEC, 1980), starch was hydrolysed by hot hydrochloric acid before measurement of specific rotation (POL.400, Isabiolgie, France).

**Digestibility calculation**

Apparent ileal digestibility (AID) was calculated from N and AA concentrations in the diets and in the ileal digesta. Standardised ileal digestibility (SID) was recalculated, according to the following formula, as suggested by Low (1982):

\[
SID \text{ AA} (%) = AID \text{ AA} (%) + \left( \frac{\text{basal endogenous AA (g/kg DMI)} \times 100}{\text{dietary AA (g/kg DM)}} \right)
\]

For this correction, assumption was made that the basal endogenous losses were in constant proportion to the amount of dry matter ingested (DMI). The basal en-
Endogenous output was previously determined in our laboratory (unpublished results), in 20 replicates, on pigs weighing on average 70 kg, prepared and managed according to the above described procedure, and fed a protein-free diet made of maize starch (580 g per kg), sugar (300 g per kg), wood cellulose (35 g per kg), soya oil (30 g per kg) and a mineral and vitamin concentrate (55 g per kg). The values of basal endogenous losses are given in Table 4.

Data analyses

The effect of the batch of MGF (B) on N and AA SID was analysed using the pig liveweight (LW) as covariate; the LWxB interaction also was tested. The introduction of LW in the model aimed at checking the absence of its effect on SID. However, the effect of group of pigs could not be tested independently of the batch of MGF.

Attempts for prediction of N and AA SID were made through regressions of the SID values on the chemical characteristics of the diets (g/kg DM) and on the ratios of the analytical data to the AA concentrations, as suggested by Sève et al. (1994). The analyses were performed using the MINITAB statistical software (MINITAB, 1996), by means of the GLM procedure, for the analyses of variance, and by means of the STEPWISE procedure, for the regression analyses.

RESULTS

For MGF5, digestibility was calculated on three replicates only because of the death of one pig during experiments (Table 1). On an average the level of feeding of pigs ranged from 73 to 80 g DM/kg LW^0.75. The average liveweight of pigs fed the different batches of MGF ranged from 42 to 61 kg.

Composition of MGF

The composition of each batch of MGF is given in Tables 2 and 3. The starch, ash and fat concentrations roughly doubled from MGF6 to MGF4, from MGF5 to MGF6 and from MGF5 to MGF7 and MGF4, respectively. The N and NDF concentrations were less variable as they increased by 23% from MGF4 to MGF3 and by 36% from MGF5 to MGF3, respectively. Except for CF and ADF (r=0.83, P<0.05), ash and ADF (r=0.85, P<0.05), and N and N-NDF (r=0.83, P<0.05), the analytical criteria were not correlated together. Particularly, the proportion of ADF in NDF was variable as it was around 24 to 26% in all batches except in MGF5 and MGF6, the ADF content of which reached 30 and 35% of the NDF, respectively.
Glutamic acid displayed the highest concentration, followed by leucine, alanine, arginine and aspartic acid, whereas tryptophan content was the lowest, followed by methionine and cystine. There was a relationship between N-NDF and the AA profile of protein: the proportion of lysine, tryptophan, aspartic acid and glycine in protein tended to increase with N-NDF (r = 0.93, 0.95, 0.95 and 0.93; P < 0.01, respectively), whereas the proportion of cystine and glutamic acid tended to decrease (r = -0.91, P < 0.01 and r = -0.76; P < 0.05, respectively).

Standardised ileal digestibility

The results of the analysis of variance and the SID values are presented in Table 4. The LWxB interaction was significant (P < 0.05) for phenylalanine and tyrosine: the digestibility of these two AA increased with the pig liveweight in MGF2, whereas it was independent of the pig liveweight in the other batches. As the interaction LWxB was significant for these two AA only, the results presented are relative to the model without interaction. No effect of the pig liveweight was detected, whereas the batch of MGF had a significant effect on the digestibility of N and most AA except for threonine, arginine, glutamic acid and glycine.
Table 4

|       | MGF1  | MGF2  | MGF3  | MGF4  | MGF5  | MGF6  | MGF7  | P²   | P²   | Batch  | r.s.d. |
|-------|-------|-------|-------|-------|-------|-------|-------|------|------|--------|--------|
| N     | 61.8  | 68.3  | 63.4  | 62.2  | 70.3  | 69.4  | 74.8  | NS   | **  | 3.0    |        |
| (3.1) |       |       |       |       |       |       |       |      |     |        |        |
| Lys   | 62.4  | 65.4  | 59.9  | 57.7  | 65.9  | 64.7  | 64.6  | NS   | *   | 3.4    |        |
| (2.6) |       |       |       |       |       |       |       |      |     |        |        |
| Thr   | 70.1  | 71.9  | 68.8  | 68.8  | 68.1  | 68.3  | 71.1  | NS   | NS  | 3.4    |        |
| (4.6) |       |       |       |       |       |       |       |      |     |        |        |
| Met   | 80.1  | 85.0  | 86.1  | 82.5  | 87.8  | 83.3  | 89.0  | NS   | *   | 3.0    |        |
| (3.0) |       |       |       |       |       |       |       |      |     |        |        |
| Cys   | 63.0  | 69.4  | 62.0  | 60.4  | 71.1  | 69.3  | 71.6  | NS   | NS  | 5.1    |        |
| (6.2) |       |       |       |       |       |       |       |      |     |        |        |
| Trp   | 59.2  | 65.0  | 61.3  | 59.9  | 64.8  | 65.9  | 72.8  | NS   | **  | 3.7    |        |
| (3.9) |       |       |       |       |       |       |       |      |     |        |        |
| Ala   | 75.9  | 82.8  | 78.3  | 75.0  | 81.5  | 77.4  | 83.4  | NS   | *** | 1.9    |        |
| (1.7) |       |       |       |       |       |       |       |      |     |        |        |
| Arg   | 86.7  | 86.7  | 83.4  | 86.5  | 87.0  | 82.9  | 86.2  | NS   | NS  | 2.5    |        |
| (1.6) |       |       |       |       |       |       |       |      |     |        |        |
| Asp   | 67.5  | 73.0  | 69.5  | 65.7  | 72.9  | 70.1  | 69.4  | NS   | *   | 3.0    |        |
| (3.5) |       |       |       |       |       |       |       |      |     |        |        |
| Glu   | 76.8  | 79.6  | 75.8  | 77.0  | 77.0  | 80.8  | 77.1  | NS   | NS  | 2.8    |        |
| (2.0) |       |       |       |       |       |       |       |      |     |        |        |
| Gly   | 60.3  | 63.2  | 59.6  | 57.4  | 62.6  | 70.6  | 63.5  | NS   | NS  | 4.9    |        |
| (6.6) |       |       |       |       |       |       |       |      |     |        |        |
| His   | 70.2  | 73.4  | 70.9  | 69.6  | 72.4  | 74.1  | 80.3  | NS   | *** | 2.2    |        |
| (2.5) |       |       |       |       |       |       |       |      |     |        |        |
| Ile   | 73.6  | 77.7  | 73.6  | 73.2  | 76.0  | 76.2  | 79.3  | NS   | **  | 2.1    |        |
| (1.6) |       |       |       |       |       |       |       |      |     |        |        |
| Leu   | 79.2  | 85.1  | 79.9  | 78.6  | 83.0  | 83.6  | 86.6  | NS   | *** | 1.7    |        |
| (1.5) |       |       |       |       |       |       |       |      |     |        |        |
| Phe   | 79.6  | 85.6  | 82.2  | 80.9  | 83.1  | 82.1  | 85.2  | -    | -   | 2.7    |        |
| (1.3) |       |       |       |       |       |       |       |      |     |        |        |
| Ser   | 73.9  | 78.6  | 75.1  | 75.3  | 77.6  | 74.0  | 80.5  | NS   | *   | 2.9    |        |
| (3.8) |       |       |       |       |       |       |       |      |     |        |        |
| Tyr   | 83.3  | 84.4  | 82.7  | 81.7  | 85.9  | 81.1  | 83.4  | -    | -   | 2.9    |        |
| (4.5) |       |       |       |       |       |       |       |      |     |        |        |
| Val   | 69.5  | 74.3  | 72.2  | 70.7  | 74.4  | 71.0  | 75.7  | NS   | *   | 2.4    |        |
| (2.4) |       |       |       |       |       |       |       |      |     |        |        |

1 expressed as percentage units – the following values of basal endogenous losses were used for calculation of SID, g/kg DMI: N 1.145, Lys 0.261, Thr 0.253, Met 0.086, Cys 0.104, Trp 0.081, Ala 0.293, Arg 0.201, Asp 0.413, Glu 0.511, Gly 0.405, His 0.088, Ile 0.193, Leu 0.297, Phe 0.165, Ser 0.247, Tyr 0.110, Val 0.261

2 probability, NS – P>0.10, *P<0.05, **P<0.01, ***P<0.001

3 residual standard deviation

4 standard deviation
Overall, MGF2, MGF5 and MGF7 displayed the highest digestibility values, and MGF1 and MGF4 the lowest. The largest significant differences between batches were observed for tryptophan, N and cystine (13.6, 13.0 and 11.2 percentage units, respectively), whereas the smallest significant differences were observed for isoleucine and valine (6.1 and 6.2 percentage units, respectively). In all batches, lysine, threonine, cystine, tryptophan, aspartic acid and glycine were among the least digestible amino acids whereas methionine, arginine, leucine, tyrosine and phenylalanine were among the most digestible ones.

**Prediction of standardised ileal digestibility**

None of the analytical criteria were relevant predictors of the SID of threonine, tryptophan, arginine, glutamic acid, glycine and histidine. For the other amino acids, the ratio of N-NDF to total N concentrations provided the equations with the lowest residual standard deviation and the highest coefficient of determination. The SID of lysine, cystine, alanine, aspartic acid, leucine, phenylalanine and valine decreased significantly (P<0.05) as the proportion of N bound to the NDF residue increased. There was a trend (P<0.10) toward a decrease of SID of N, isoleucine and serine, with the increase in this ratio (Table 5).

| Table 5 | Prediction of nitrogen and amino acid standardised ileal digestibility (%) in maize gluten feed according to the ratio of the N concentration in the NDF residue (N-NDF) to the total N concentration (g/g) |
|---------|------------------------------------------------------------------------------------------|
| Intercept | s.d. | N-NDF/N | s.d. | P² | R² | r.s.d. |
| N | 73.7 | (3.4) | 37.1 | (17.4) | 0.09 | 47.7 | 3.8 |
| Lys | 67.8 | (1.8) | 27.2 | (9.4) | * | 62.9 | 2.1 |
| Met | 89.2 | (2.1) | 24.9 | (10.5) | 0.06 | 52.9 | 2.3 |
| Cys | 73.8 | (2.9) | 39.8 | (14.9) | * | 58.7 | 3.3 |
| Ala | 85.2 | (1.3) | 33.9 | (6.9) | ** | 83.0 | 1.5 |
| Asp | 74.3 | (1.3) | 25.4 | (6.5) | * | 75.4 | 1.4 |
| Ile | 79.0 | (1.6) | 18.7 | (8.0) | 0.07 | 52.5 | 1.8 |
| Leu | 86.9 | (2.0) | 25.8 | (10.0) | * | 57.4 | 2.2 |
| Phe | 86.0 | (1.3) | 18.4 | (6.8) | * | 59.8 | 1.5 |
| Ser | 79.8 | (1.8) | 18.7 | (9.1) | 0.10 | 45.8 | 2.0 |
| Tyr | 85.4 | (1.1) | 12.5 | (5.8) | 0.08 | 48.5 | 1.3 |
| Val | 76.2 | (1.3) | 20.7 | (6.7) | * | 65.9 | 1.5 |

1 standard deviation
2 probability, *P<0.05, **P<0.01
3 coefficient of determination
4 residual standard deviation
DISCUSSION

Composition of MGF

The composition of the batches is in the range reported in the electronic database (AFZ, 1998) for more than 700 batches of MGF collected in France (N=27 to 42, CF=47 to 111, ash=39 to 103 and starch=91 to 362 g/kg DM). The overall absence of correlations between the analytical criteria is probably the result of the variability in the processes implemented during starch extraction, as well as the variability in the level of incorporation of the ingredients (maize steep liquor, envelopes and sometimes maize germ meal), a maize gluten feed is made from. The negative correlation between N-NDF and N may originate from an increasing level of incorporation of maize steep liquor, rich in N, and, consequently, a decreasing level of incorporation of fibrous residues in the final product. Accordingly, N-NDF is positively correlated with the concentrations of lysine, tryptophan, aspartic acid and glycine, mainly located in the fibrous part of the maize grain, and negatively correlated with glutamic acid, mainly located in the starchy endosperm of maize (Landry and Moureaux, 1980; Chung and Pomeranz, 1985).

Standardised ileal digestibility

The analysis of the effect of liveweight confirmed that this parameter is of little influence on ileal digestibility of amino acids in growing pigs. This is in accordance with previous demonstration that the end-to-end ileo-rectal anastomosis is appropriate to study ileal digestibility of amino acids in growing pigs, even in the long term (Laplace et al., 1994; Bodin et al., 1998). No obvious explanation appeared for the significant LWxB interaction observed for phenylalanine and tyrosine. However, it should be considered with care as the effect of the animal, confounded with the effect of liveweight in our experimental design, may have influenced the results.

On average, the SID of N in MGF was by 19.5 percentage units lower than in maize (Jondreville et al., 2000a). Our SID values were quite low compared with other literature data: with a single batch of MGF (CF=112 and N=39 g/kg DM), Mariscal-Landin (1992) calculated that the SID of N reached 75.9% and the value reported in the Rhodimet Nutrition Guide (RPAN, 1993) reaches 74%. These authors also reported high SID values for methionine, alanine, arginine, leucine, phenylalanine and tyrosine and low SID values for lysine, threonine, cystine, aspartic acid and glycine. Overall, AA were similarly ranked, regarding their SID, in maize and in MGF (Jondreville et al., 2000a). However, the extent of the difference in SID between maize and MGF differed according to the AA. Overall, the lower the digestibility, the greater the difference: large differences
were observed for lysine, aspartic acid and glycine (20.5, 18.9 and 20.7 percentage units, respectively), whereas small differences were observed for methionine, arginine, leucine, phenylalanine and tyrosine (8.7, 5.8, 10.5, 8.8, 6.2 percentage units, respectively).

By calculating SID, no attempt for evaluating the possible influence of the nature of the feedstuffs on total endogenous losses was made. The correction of AID for basal endogenous losses only aimed at avoiding the digestibility results to be biased by the dietary level of N and AA (Fan et al., 1994; Sève and Henry, 1996) and at getting additive values of digestible AA supply (Furuya and Kaji, 1991). However, there is evidence that total endogenous losses largely exceed basal endogenous losses. Even corrected for basal endogenous N losses, ileal N thus contains variable amounts of specific endogenous N, depending on the characteristics of the feedstuffs (e.g. De Lange et al., 1990; Grala et al., 1998). Thus, factors affecting either the feed specific endogenous losses, or the undigested dietary fraction, or both, may influence the ranking of AA within a single batch, as well as the differences between the batches.

Dietary fibre may impair SID by encapsulating dietary proteins and limiting their access to digestive enzymes (Green et al., 1987; Leterme et al., 1989; Sève et al., 1994). This phenomenon probably explains the low digestibility of lysine, aspartic acid and glycine, located in the most fibrous part of the maize grain and the high digestibility of leucine, tyrosine and phenylalanine, mainly located in the endosperm fraction of the maize grain. As reported by Sève et al. (1994) and Jondreville et al. (2000b) in wheat milling by-products, the proportion of N bound to the NDF residue is a relevant predictor for SID of N and most AA in MGF. However, in most cases, this criteria, which may account for the variable level of recovery of dietary AA, explains less than 65% of the total variability.

Relatively high concentrations of lysine, threonine and particularly tryptophan in the endogenous protein, compared with dietary protein, undoubtedly contributed to their overall low SID in MGF. Many authors reported increasing amounts of N ileal endogenous losses with increasing amounts of dietary fibre (Sauer et al., 1977; Green et al., 1987; Mariscal-Landin et al., 1995, Schulze et al., 1995). SID was then expected to decrease with the increase in dietary fibre concentration. Such a decrease was observed by Sève et al. (1994) and Jondreville et al. (2000b) in wheat milling by-products. In these studies, the ratio of the dietary ADF or ADL to AA concentrations were relevant predictors for SID of most AA. The absence of relevance of any of the fibre criteria in the present study may originate from their low variability in MGF: the proportion of N bound to the NDF residue varied by 250%, whereas the ratios of dietary ADF to AA concentration varied roughly by 60%. In wheat milling by-products, the two criteria varied by about 300% from wheat middlings to wheat bran. It may also be due to either the great variability in the fibre composition of these by-products, or to the lack of rele-
vance of the classical analytical characterisations we performed for a reliable description of fibre in MGF.

At the end, despite the large variability of the batches of MGF we used, both in terms of industrial origin and of composition, a quite simple criteria (N-NDF) allows to explain part of the variability in SID of most AA. However, further investigations, both for a better characterisation of these feedstuffs and for a better understanding of the factors influencing endogenous losses in pigs fed such by-products, are required for a more accurate prediction of AA SID.

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STRESZCZENIE

Strawność jelitowa aminokwasów glutenu kukurydzianego u świń

Standardową strawność jelitową (SID) białka i aminokwasów siedmiu partii glutenu kukurydzianego oznaczano na wieprzach rasy Wielka Biała (LW) lub mieszancach (LW x Landrace) x (LW x Pietrain) o masie ciała od 42 do 61 kg z przetokami mostkowymi na końcu jelita biodrowego. Każda partia glutenu była testowana na 4 zwierzętach.

Wartości SID: lizyny – 62,9; treoniny – 69,6; cystyny – 66,7; tryptofanu – 64,1; kwasu asparaginowego – 69,7 i glicyny – 62,4 wszystkich badanych partii glutenu były najmniejsze. Najlepiej były trawione: metionina – 84,8; arginina – 85,6; leucyna – 82,3; trypsyna – 83,2 i fenyloalanina – 82,7. Istotne różnice SID pomiędzy badanymi partiami glutenu stwierdzono dla większości aminokwasów. SID większości aminokwasów była mniejsza, gdy zwiększał się udział azotu związanego z NDF.