Short communication: Applicability of feeding behaviour traits as high-throughput phenotyping methods for identifying protein-efficient pigs 4 Author: Esther Ewaoluwagbemiga¹, Giuseppe Bee¹ and Claudia Kasper^{1,2,}* 6 ¹ Swine Research Unit, Agroscope Posieux 1725, Switzerland

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- Short title: Feeding behaviour to select protein-efficient pigs

Abstract

 The objective of this study was to explore the potential of using automatically recorded feeding behaviour as a proxy for protein efficiency (PE) by investigating the relationship between feeding behaviour and PE. A total of 402 Swiss Large White pigs were used in this experiment (204 females and 198 castrated males). Pigs were fed ad libitum on a reduced protein diet (80% of standard) from 20kg to 100kg BW. Individual daily feed intake was monitored and carcass composition at slaughter was determined by dual-energy X- ray absorptiometry (DXA). The PE was calculated as the ratio of protein in the carcass (estimated by DXA) to the total protein consumed. Feeding behaviour traits monitored were daily feed intake (DFI; g/day), feed intake per visit (FIV; g/visit), number of daily visits (NDV; visits/day), duration of visits (DUV; min/visit), feeding rate (FR; g/min), and feeder occupation (FO; min/day). Regression analysis was used to estimate the relationship between PE and feeding behaviour, while correcting for the effects of sex, experimental series and age. Weak Pearson's correlations (-0.25 to 0.12) were found between PE and 28 feeding behaviour traits. Beta (β) estimates from this analysis for feeding behaviours were also very low (0.0093% to 0.087%). An increase in FR (g/min) will increase PE by 0.087%

 and an increase in DFI (g/day) will decrease PE by 0.0093%. In conclusion, feeding behaviours are not suitable for the identification of protein-efficient pigs, as estimates are negligible.

Keyword

Swine, protein retention, nitrogen, environmental pollution, carcass

Implications

 This study suggests that feeding behaviour traits recorded via automatic feeder are not reliable predictors of protein efficiency in Swiss Large White pigs receiving a protein- reduced diet. Despite the large differences in protein efficiency, only negligible changes in a range of feeding behaviours were observed. Hence, feeding behaviours are not suitable proxies for the high-throughput phenotyping of protein efficiency and the selection of live animals for use in nutrition experiments or for breeding.

Introduction

 Protein is an important nutrient for livestock, necessary for growth, building and repair of tissues, production of milk, and immune functions. However, less than 50% of ingested dietary protein in pigs is converted into carcass muscle (Kasper et al., 2020a; and Millet et al., 2018). The excess dietary protein that is not converted to meat or used for maintenance cannot be stored in the body for future use but it is rather excreted as waste, thereby contributing to environmental pollution (Wang et al., 2018). Given the growing human demand for animal protein worldwide, limited arable land for livestock production, competition between humans and livestock for plant protein, and the need to preserve the

 environment, there is a pressing need to maximize protein efficiency (PE) of livestock. Both genetic (Shirali et al., 2013; Kasper et al., 2020a) and non-genetic (e.g. nutrition and weight at slaughter; Ruiz-Ascacibar et al., 2017) means are currently being considered in improving protein efficiency in pigs. However, PE is a difficult trait to measure due to costs and time involved, and the necessity to anesthetize or slaughter the animals. PE can be measured chemically, which is laborious and costly. An alternative is the use of modern imaging technology, such as DXA (dual-energy X-ray absorptiometry) (Kasper et al., 2020b), which has, after the initial investment, lower operating costs, but still requires a considerable amount of human resources for image acquisition and processing. Preferably, a proxy can be used as a high-throughput phenotyping method for the selection of protein-efficient pigs, which is ideally recorded automatically, in a non- destructive fashion and/or in the course of standard rearing conditions. Such a proxy could be feeding behavior (Ding et al., 2018) that allows the evaluation and selection of animals during their lifetimes without the need for slaughter. Previous studies have reported phenotypic correlations of -0.01 to 0.35 between feed conversion ratio (FCR) and feeding behaviour traits (Carcò et al., 2018; Kavlak and Uimari 2019), and slightly higher correlations of 0.147 to 0.408 were reported between feeding behaviour traits and protein retention (Carcò et al., 2018). In addition, an increasing number of farms are using computerized feeding systems that automatically record individual feeding patterns without disturbing the animals. These feeding systems allow for easy evaluation of feeding behaviour traits and facilitates the collection of large amounts of data required for genetic experiments. Thus, if adequate and consistent correlations between feeding behaviour and protein efficiency exist, feeding behaviour can be used as a proxy trait for PE to select protein-efficient pigs.

Materials and Methods

Protein efficiency and feeding behaviours

 A total of 402 Swiss large white pigs (Nfemales = 204, Ncastrated-males = 198) from eight farrowing series were used. The pigs were fed ad libitum a starter diet from weaning until 20 kg BW, and grower and finisher diet from 20 to 60 and 60 to 100 kg BW, respectively. The starter diet was formulated according to the current Swiss feeding recommendations for swine, whereas the levels of digestible protein and essential amino acids were 20% lower than the recommended levels (Ruiz-Ascacibar et al., 2017). Pigs were slaughtered at a BW of approximately 100 kg and the left carcass side including the whole head were scanned with dual-energy X-ray absorptiometry (GE Lunar i-DXA, GE Medical Systems, Glattbrugg, Switzerland), which estimates, among other things, lean tissue content. The lean tissue content obtained from DXA was used in the following calibration equation to estimate the protein content retained in animal's carcass (Kasper et al., 2020). .

93 *g protein carcass* = $-482.745 + 0.23 \times (g \text{lean tissue DXA x 2})$

 Protein efficiency of the carcass was thereafter calculated as the ratio of protein retained in the carcass (corrected for protein content at 20 kg BW) to the total protein intake.

98 **betta protein efficiency** = $\frac{g\,protein\,carcass-g\,protein\,starat}{g\,protein\,intake}$

 Protein content of pigs at 20 kg BW was estimated by multiplying the actual live weight of pigs when the reduced-protein diet experiment started with a chemically determined

102 protein content per kg carcass (protein/kg carcass) of piglets slaughtered at 21.26 ± 1.59 kg in a previous experiment (*g protein start*) (Kasper et al., 2020).

 The time of feeding, duration of feeding and quantity of feed consumed per visit was recorded for each animal with an automatic feeder as described earlier (Ruiz-Ascacibar et al., 2017). Record of visits with no feed intake (i.e. 0 g) were removed from the dataset. In total, 702,162 visits were recorded for the 402 pigs. Feeding behaviour traits evaluated were daily feed intake (DFI; g/day), number of daily visits to the feeder (NDV; visits/day), feed intake per visit (FIV; g/visit), duration per visit (DUV; min/visit), feeding occupation (FO; min/day), and feeding rate (FR; g/min/day). Feeding behaviour traits were computed individually, and the mean of each individual was used for further analysis.

Statistical analysis

 Data were analyzed with R software V 3.6.3. The relationships between protein efficiency and feeding behaviour traits were estimated with linear models, and corrected for the effects of sex, experimental series and age at slaughter. However, due to the high correlations between some of the feeding behaviour traits (Pearson correlation coefficients r = -0.84 to 0.92, *p* < 0.01), residuals of FIV (rFIV), residuals of FO (rFO) and the residuals of DUV (rDUV) were used in the final regression model. rFIV was the linear regression of NDV, DUV and DFI on FIV, rFO was the linear regression of DUV, DFI and FR on FO, and rDUV was the linear regression of NDV, DFI, FR, FO, and FIV on DUV.

126 **Results**

- 127 Means and standard deviation of PE and feeding behaviour traits are presented in Table
- 128 1. Pearson's correlation coefficients between PE and feeding behaviour traits were weak,
- 129 and varied between -0.25 to 0.12 (Table 1). Negative Pearson's correlations with PE were
- 130 observed for DFI, FIV, DUV and FO (i.e. higher daily feed intake, feed intake per visits,
- 131 durations at visits, and feeding occupation is associated with lower PE), and positive
- 132 correlations with PE were observed for NDV and FR (i.e. more daily visits and higher
- 133 feeding rate is associated with higher PE).
- 134 **Table 1:** Descriptive statistics (mean ± SD) and Pearson's correlation coefficients of protein 135 efficiency and feeding behaviour traits in pigs

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138 Multiple linear regression analysis revealed significant relationships of PE with FR, rFIV,

139 DFI, and NDV (Table 2). A one-unit increase in FR was associated with an increase in PE

140 by 0.087%, a one-unit increase in FIV decreased PE by 0.0168%, and a unit increase in

141 DFI decreased PE by 0.0093%. Despite their significant difference from zero, these slopes

142 were very flat (Figure 1).

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148 FIV is the model residual of FIV; rFO is the model residual of FO, and rDUV is the model residual of DUV. Due to high correlations

149 between some feeding behaviour traits, model residuals of FIV, FO and DUV were used in the final regression model to avoid 148 FFIV is the model residual of FIV; rFO is the model residual of FO, and rDUV
149 between some feeding behaviour traits, model residuals of FIV, FO and D
150 collinearity. Original values of FR, DFI and NDV were used in

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Discussion

 In this study, we investigated the potential of using automatically generated feeding behaviour traits as proxies for protein efficiency for high-throughput phenotyping methods in pigs. The advantage of this approach is that protein-efficient pigs may be chosen based on their feeding behaviour data collected during the rearing period, and large phenotypic data sets become available for genetic experiments. However, the result of this study showed weak relationships between feeding behaviour traits and protein efficiency. From linear regression analysis, it is clear that there was no association between PE and feeding behaviour traits (Figure 1).

 The mean carcass PE in this study was 0.38, which is in line with the carcass PE of Kasper et al. (2020a), who investigated the potential to genetically improve protein efficiency in pigs. The mean FO, DFI and FR observed in this study were slightly lower to those reported by Kavlak and Uimari (2019) (FO: 61 min/d; DFI: 2100 g/d; FR: and 40 g/min) and those reported by Do et al. (2013) (FO: 78min/d; DFI: 2340 g/d; FR: 31 g/min). However, the mean feeding behaviour traits reported by Kavlak and Uimari (2019) were recorded over five periods (i.e. growth phases) and the study of Do et al. (2013) was on entire males. In our study, the focus was on the whole growth cycle (from weaning to slaughter weight), castrated males and females were used and dietary protein was reduced, which may influence feeding behaviour patterns.

 The relationships between PE and feeding behaviour traits in this study were weak (i.e. very low estimates), although regression analysis showed statistical significance of some feeding behaviour traits, probably due to large sample size. For instance, a one-unit

 increase in FR (g/min) will increase PE by 0.087% and a one-unit increase in DFI (g/day) will decrease PE by 0.0093%, which is certainly of no biological relevance, and will therefore have little to no impact on PE. This result is similar to the findings of Carco et al. (2018), where FR was the only moderately correlated trait with protein retention among other feeding behaviour traits (r = 0.41), but they concluded that FR has very little influence on feed efficiency. Colpoys et al. (2016) did not find any significant relationship between feeding behaviour traits and estimated protein retention. The large variations that exist among individuals with respect to feeding behaviour traits (Carco et al., 2018) may account for the little to no relationships found between feeding behaviour traits and protein efficiency. Carco et al. (2018) described feeding behaviour as a flexible strategy each pig follows to cover its nutrient requirement, which consequently may lead to variations in feeding behaviour between individuals. The nutrient requirement of each individual may be influenced by feed composition, among other factors such as genotype, health and welfare. For example, Schiavon et al. (2018) found that a reduction in the essential amino acid content of the diet increased feeding rate and feed intake. Since reduced dietary protein was fed in the present study, pigs might have responded in different ways to the diet in terms of feeding behaviour, which could have led to larger variations in feeding behaviours. This weak to no relationship observed between PE and feeding behaviours shows that differences in PE between pigs is not affected by feeding pattern. However, other factors such as metabolic processes that regulate PE and genotype may influence PE.

 Although little to no relationship was found between feeding behaviour and protein efficiency in this present study, further studies could estimate heritability of feeding

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- Conceptualization: E.E., G.B. and C.K. Data Curation: E.E. Formal Analysis: E.E. Funding
- Acquisition: G.B. and C.K. Methodology: E.E. and G.B. Project Administration: C.K.
- Software: E.E. Supervision: G.B. and C.K. Visualization: E.E. Writing Original Draft
- Preparation: E.E. Writing Review & Editing: E.E., G.B. and C.K.

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