Staple food crops from Brazilian Biofortification Program have high protein quality and hypoglycemic action in Wistar rats

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Abstract

Biofortified staple food crops have become an effective tool by which to address micronutrient deficiencies in many at-risk populations. The effect of these food crops on the micronutrient status has been evaluated, however there is no studies investigating their protein quality, neither by themselves or in a food combination. Thus, this study investigated the effect of combinations of beans and rice, target for biofortification, with high carotenoids content crops (sweet potato and pumpkin) on protein quality, intestinal function and glycemia. Eight groups were evaluated (n=6): 1) rice + Caupi bean (RCB); 2) rice + Pontal bean (RPB); 3) rice + Caupi bean + pumpkin (RCBP); 4) rice + Pontal bean + pumpkin (RPBP); 5) rice + Caupi bean + sweet potato (RCBS); 6) rice + Pontal bean + sweet potato (RPBS); 7) casein (positive control); and 8) nitrogen free (negative control). Tested groups showed high protein quality, and groups containing Caupi bean showed higher true digestibility. Groups containing Pontal bean had a greater reduction of blood glucose compared to casein. Tested groups showed higher fecal excretion of lipids. Combination of food target for biofortification showed high protein quality, and the combinations containing Pontal beans showed hypoglycemic effect.

Keywords: protein; biofortification; blood glucose; fecal lipid; short chain fatty acids.

Practical Application: The Biofortification program not only provide an additional amount of micronutrient but also contribute to the protein intake of the risk population and potentially improve the glycemic status and intestinal function. Thus, the utilization of these foods may have a positive impact on the nutritional status and health of population at risk for micronutrient deficiency.

1 Introduction

Biofortified staple food crops have become an effective tool by which to address micronutrient deficiencies in many at-risk populations (Bouis & Saltzman, 2017; Dias et al., 2018; Glahn et al., 2017). It aims to improve the micronutrient concentration of staple food crops through the best practices of breeding and modern biotechnology (Bouis & Saltzman, 2017; Nutti et al., 2006). These food crops were chosen because their large production and consumption in Brazil, so they can be largely distribute to the population, especially those at nutritional risk (Bouis & Saltzman, 2017; Nutti et al., 2006). The iron bioavailability and the efficiency of these foods in improving the nutritional status of iron have been tested in animals (Dias et al., 2015, 2018; Glahn et al., 2017; Tako et al., 2011, 2015) and in humans (Haas et al., 2016; Vaz-Tostes et al., 2016). However, there is no studies about its protein quality, neither by themselves or in a meal.

Some plant foods are sources of protein, however, often present inadequate amounts of essential aminoacids. Although they are not source of high protein quality, these foods contribute significantly to the overall protein intake of the population, they represent the protein sources of lower cost and therefore higher. The combination of different food sources can improve the balance of essential aminoacids and, consequently, protein quality (Cintra et al., 2007; Rezende et al., 2017).

The protein quality of food can be influenced by phytochemicals in foods such as beans, including phytate, tannins and phenolic compounds, which can bind the minerals, proteins and starches, forming insoluble complexes which impair the absorption of these nutrients (Ramírez-Cárdenasi et al., 2008; Rezende et al., 2017). On the other hand, some compounds present in the beans, such as dietary fiber, tannins, phytates and amylase inhibitors have been inversely correlated to the digestion of carbohydrates, glycemic response and the intestinal function (Dias et al., 2018; Hutchins et al., 2012; Pacifici et al., 2017). Also, dietary fiber can improve the lipid metabolism and modulate the intestinal function, since it can reduce fat digestion and absorption, increasing its excretion in feces (Chen et al., 2014).
There are some studies that evaluated the protein quality in beans (Cruz et al., 2005) and in the combination of rice and bean (Cintrón et al., 2007; Kannan et al., 2001). However, there is no studies evaluating the protein quality in combination of biofortified foods, such as beans, rice, pumpkin and sweet potato and its nutritional benefits.

Considering this information, the objective of this study was to evaluate the effect of food combinations of beans and rice, target for biofortification, with high carotenoids content crops (sweet potato and pumpkin) on the protein quality, blood glucose and intestinal modulation.

2 Materials and methods

2.1 Sample

Since we aimed to evaluate the protein quality and nutritional benefits of biofortified foods, we used the following staple food crops from Brazilian Biofortification Program: common beans (Phaseolus vulgaris L.) BRS Pontal and BRS Xique-Xique (caipi) (high Fe content); white rice (Oryza sativa) Chorinho; pumpkin (Curcubita moscata) Duchesne and sweet potato (Ipomoea batatas) (high pro-vitamin A carotenoid content). Cultivars were developed and supplied by Empresa Brasileira de Pesquisa Agropecuária (EMBRAPA), Brazil.

2.2 Staple food crop flours preparation

The common beans were cooked in three replicates in a conventional pressure cooker for 40 minutes using a bean/water ratio of 1:2.2 (w/v) and dried in an air oven for 17 hours at 60 °C (Ramírez-Cárdenasi et al., 2008). The rice was cooked in three replicates in a conventional cooker using a rice/water ratio of 1:2.8 (w/v) and dried in an air oven for 17 hours at 60 °C. The pumpkin and sweet potato were peeled and sliced on a multiprocessor and dried in an air oven for 6 hours at 60 °C (Dias et al., 2015). All foods were ground by mill 090 CFT at 3000 rpm, sieved (600 mesh screen) and stored at -12 °C (Ramírez-Cárdenasi et al., 2008).

2.3 Chemical composition of foods

Protein concentration was determined by micro-Kjeldahl method according to the AOAC procedure (Association of Official Analytical Chemists, 2012). The determination of total fiber and soluble and insoluble fractions was performed by the enzymatic-gravimetric method, according to Association of Official Analytical Chemists (2012), using the enzymatic hydrolysis for a heat-resistant amylase, protease, and amyloglucosidase (Total dietary fiber assay Kiyonaga, Sigma®). The lipid concentration was determined by extraction with ethyl ether using Soxhlet apparatus (Association of Official Analytical Chemists, 2012). The ash concentrations were determined by the gravimetric method based on the weight loss of the materials subjected to heating at 550 °C (Association of Official Analytical Chemists, 2012). Moisture was determined by drying 10 grams of the samples for 24 hours at 105 °C in the air oven (Marconi, SP, Brazil) (Association of Official Analytical Chemists, 2012).

Determination of carotenoids

The extraction of pro-vitamin A carotenoids (α and β-carotene) in pumpkin and sweet potato was performed according to Rodriguez et al. (1976). Fifty grams of the pumpkin and sweet potato flours was ground in 60 mL of chilled acetone for approximately 2 minutes in a micro grinder and the material was vacuum filtered on Buchner funnel using filter paper. The filtrate was transferred to a separator funnel, in which 50 mL of cooled petroleum ether were added to transfer the pigment from acetone to petroleum ether. Each fraction was washed three times with distilled water to remove all acetone. The concentration of material was determined by evaporation of the petroleum ether extract using a rotary evaporator at 35 °C. The pigments were re-dissolved in a known amount of petroleum ether and stored in amber glass vials at -18 °C.

The analyses of carotenoids were performed in triplicate by high performance liquid chromatography (HPLC) using the chromatographic conditions developed by Sant'Ana et al. (1998) as follows: HPLC system (Shimadzu, SCL 10AT VP, Japan) comprised of a high-pressure pump (Shimadzu, LC-10AT VP, Japan), autosampler with 50 µL loop (Shimadzu SIL-10A VP, Japan) and diode array detector (DAD) (Shimadzu SPD-M10A, Japan); chromatographic column, Phenomenex Gemini RP-18 (250 mm × 4.6 mm, 5 µm), equipped with a guard column, RP-18 Phenomenex ODS column (4 mm × 3 mm). The mobile phase was methanol: ethyl acetate: acetonitrile (80:10:10, v/v/v) at a flow rate of 2.0 mL/min. Chromatograms were obtained at 450 nm.

Phytate and phenolic compounds

Phytate content was determined in triplicate by ion exchange and spectrophotometry according to Latta & Eskin (1980), with modifications by Ellis & Morris (1986). The determination of the concentration of phenolic compounds in foods was performed using the Folin Ciocalteu reagent as described by (Singleton et al., 1999).

2.4 Diets

The composition of the experimental diets were based on AIN-93G diet (Reeves et al., 1993). Replacement of casein diet control the flour of biofortified food was based on food protein content and the contribution of these to the daily protein intake of children aged 7 to 10 years of age, according to research conducted by Hinnig & Bergamaschi (2012). In this study, it was observed that the bean contribution to the total protein diet consumption was 8.85%, the rice was 4.01%, the pumpkin and sweet potato was 0.35% each. Thus, based on the amount of protein of these foods it reached a ratio of 50% of rice, beans 50%, 13% sweet potato and 3% pumpkin. The composition of the diet was calculated so that the diets were isocaloric and isonitrogenous (Supplementary Material Table S1).

2.5 Biological assay

Experiment I: evaluation of protein quality

Forty eight male rats (Rattus norvegicus, Wistar) weaning were randomized into eight groups (n= 6): 1) Rice + Caipi Bean (RCB); 2) Rice + Pontal Bean (RPB); 3) Rice + Caipi Bean +
Pumpkin (RCBP); 4) Rice + Pontal Bean + Pumpkin (RPBP); 5) Rice + Caupi Bean + Sweet Potato (RCBS); 6) Rice + Pontal Bean + Sweet Potato (RPBS); 7) Positive Control (Casein); 8) Negative Control (no protein) (Figure 1). The animals were kept in individual cages in a room with 12 hours photoperiod and an average temperature of 22 °C (Figure 1).

Food intake and body weight of the animals were recorded weekly. To determine the digestibility diets were labeled with indigo carmine (200 mg.100 g⁻¹) and feces were collected for a period of 4 days. The amount of protein diets, and fecal nitrogen was determined by semi-micro Kjeldahl method. Evaluation of protein quality was determined by Protein Efficiency Ratio (PER) (Association of Official Analytical Chemists, 1975), Net Protein Ratio (NPR) and true digestibility (Amaya et al., 1991).

**Experiment II: effects on intestinal function, and modulation of glucose**

To evaluate the effect of ingestion of food on the combinations of the modulation of gut function, lipid profile and blood glucose experiment I was extended for further 14 days, keeping the following groups: Rice + Caupi Bean + Pumpkin (RCBP); Rice + Pontal Bean + Pumpkin (RPBP); Rice + Caupi Bean + Sweet Potato (RCBS); Rice + Pontal Bean + Sweet Potato (RPBS); Positive Control (Casein) (Figure 1).

After 28 days and 12 hours fasting, the animals were anesthetized with isoflurane (Osoforine, Cristália®) were euthanized by cardiac puncture. Blood was collected for the analysis of blood glucose and serum lipids and feces from the cecum were collected for lipid analysis and short-chain fatty acids in the feces. Furthermore, colon fragments were removed and fixed in buffered formalin 10% solution for histological analysis.

This study was approved by the Ethics Committee for Animal Experimentation of the Federal University of Viçosa, Viçosa, MG.

**2.6 Blood glucose**

Plasma glucose concentrations were determined by colorimetric methods according to the manufacturer’s instructions (Bioclin®).

**2.7 Short Chain Fatty Acids (SCFA)**

The content of propionic acid, acetic and butyric acids in the feces was determined according to Smiricky-Tjardes et al. (2003) with modifications. The determination of SCFA concentrations in the sample was made in a gas chromatograph (Model GC-2010, Shimadzu Scientific Instruments Inc., Japan). The drug was made through the hydrogen gas stream of 1.8 mL/min. An initial oven temperature of 100 °C for 0.5 minutes was used, with subsequent increase of 8 °C/min until the temperature of 180 °C and this was maintained for 1 minute, then increase 20 °C/min to 200 °C, which was maintained for 5 minutes. The identification was made by comparison of the retention time, as external standard using a mixture of free short chain fatty acids (free acid Volatile mix codes. 46975, Sigma Aldrich, USA). Quantitation was done by using the standard curve at concentrations from 2 to 10 mM.

**2.8 Fecal lipids**

The total lipid content in the feces was determined by extraction using a Soxhlet apparatus, according to the analytical method of Association of Official Analytical Chemists (2012).

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**Figure 1.** Scheme showing the experimental design of the study. RCB = Rice and Caupi Beans; RPB = Rice and Pontal Bean; RCBP = Rice, Caupi Bean and Pumpkin; RPBP = Rice, Pontal Bean and Pumpkin; RCBS = Rice, Caupi Bean and Sweet Potato; RPBS = Rice, Pontal Bean and Sweet Potato.
In this procedure, extraction was conducted using ethyl ether as solvent.

2.9 Colon histomorphometric analysis

Semi-serial histological sections of fragments of the colon of 3 mm thickness were obtained using automatic microtome (Reichert-Jung®, Germany) and were stained by the Toluidine Blue technique. The slides were examined under an Olympus CX31 light microscope. To measure crypt depth (CD) and thickness of the circular and longitudinal muscle layers (CML and LML, respectively), twenty random fields per animal were selected (Silva et al., 2016) and were obtained and analyzed using the ImagePro-Plus® software version 4.5 (Media Cybernetics, Rockville USA).

2.10 Statistical analysis

The floors of staple food crops were analyzed in three replicates. The treatments were conducted in a completely randomized design, with six replications. The results were analyzed by analysis of variance. For "F-value" significant, the Newman-Keuls test was used to compare means among the experimental groups. Statistical analyzes was performed using the Statistical Analysis System software, version 9.1. P-value <0.05 was considered statistically significant.

Table 1. Chemical composition of staple food crop flours

<table>
<thead>
<tr>
<th></th>
<th>Caupi Bean</th>
<th>Pontal Bean</th>
<th>Rice</th>
<th>Pumpkin</th>
<th>Sweet Potato</th>
</tr>
</thead>
<tbody>
<tr>
<td>Moisture</td>
<td>8.27± 0.1</td>
<td>10.7± 0.28</td>
<td>7.35± 0.06</td>
<td>9.99± 0.55</td>
<td>9.92± 0.06</td>
</tr>
<tr>
<td>Ash</td>
<td>2.93± 0.02</td>
<td>3.14± 0.03</td>
<td>0.34± 0.02</td>
<td>6.38± 0.07</td>
<td>2.27± 0.06</td>
</tr>
<tr>
<td>Lipids</td>
<td>2.23± 0.22</td>
<td>1.37± 0.3</td>
<td>0.13± 0.13</td>
<td>1.46± 0.14</td>
<td>1.55± 0.34</td>
</tr>
<tr>
<td>Protein</td>
<td>23.15± 0.32</td>
<td>18.86± 0.08</td>
<td>8.83± 0.18</td>
<td>15.86± 0.24</td>
<td>2.63± 0.12</td>
</tr>
<tr>
<td>Total dietary fiber</td>
<td>15.53± 0.92</td>
<td>26.69± 0.45</td>
<td>10.08± 0.1</td>
<td>15.02± 0.03</td>
<td>15.31± 0.31</td>
</tr>
<tr>
<td>Soluble fiber</td>
<td>1.88± 0.11</td>
<td>7.04± 1.27</td>
<td>0.37± 0.08</td>
<td>5.10± 0.25</td>
<td>4.89± 0.38</td>
</tr>
<tr>
<td>Insoluble fiber</td>
<td>13.66± 0.82</td>
<td>19.64± 0.92</td>
<td>0.87± 0.43</td>
<td>9.92± 0.23</td>
<td>10.42± 0.38</td>
</tr>
<tr>
<td>Carbohydrates</td>
<td>48.87w± 0.73</td>
<td>48.87w± 0.73</td>
<td>82.48± 0.05</td>
<td>52.19w± 0.34</td>
<td>69.62w± 0.56</td>
</tr>
<tr>
<td>Total phenolic (mg/100 g)</td>
<td>89.0± 0.09</td>
<td>133± 0.15</td>
<td>6± 0.01</td>
<td>241± 0.12</td>
<td>151± 0.07</td>
</tr>
<tr>
<td>Carotenoids (mg/100 g)</td>
<td>nd</td>
<td>nd</td>
<td>308.84± 1.98</td>
<td>127.11± 0.06</td>
<td></td>
</tr>
<tr>
<td>Phytic acid</td>
<td>0.54± 0.01</td>
<td>0.51± 0.02</td>
<td>0.20± 0.03</td>
<td>0.03± 0.32</td>
<td>0.10± 0.1</td>
</tr>
</tbody>
</table>

Data presented as mean and standard deviation. nd = not determined. Means with different letters in the same line present significant difference (p<0.05) by Tukey test.

Table 2. Effect of the staple food crops intake on the protein quality

<table>
<thead>
<tr>
<th></th>
<th>PER-R</th>
<th>NPR-R</th>
<th>TD</th>
<th>TD-R</th>
</tr>
</thead>
<tbody>
<tr>
<td>Casein</td>
<td>3.43± 0.74</td>
<td>100± 0</td>
<td>100± 0</td>
<td>94.24± 1.32</td>
</tr>
<tr>
<td>RCB</td>
<td>2.73± 0.36</td>
<td>80.3± 14.37</td>
<td>3.39± 0.5</td>
<td>83.11± 12.47</td>
</tr>
<tr>
<td>RPB</td>
<td>3.33± 0.3</td>
<td>97.02± 8.84</td>
<td>4.22± 0.27</td>
<td>103.38± 6.61</td>
</tr>
<tr>
<td>RCBP</td>
<td>3.11± 0.39</td>
<td>90.63± 11.4</td>
<td>3.84± 0.45</td>
<td>94.09± 11.17</td>
</tr>
<tr>
<td>RPBP</td>
<td>3.30± 0.36</td>
<td>96.28± 10.66</td>
<td>4.14± 0.38</td>
<td>101.34± 9.3</td>
</tr>
<tr>
<td>RCBS</td>
<td>3.14± 0.42</td>
<td>91.45± 12.25</td>
<td>3.98± 0.51</td>
<td>97.43± 12.48</td>
</tr>
<tr>
<td>RPBS</td>
<td>2.69± 0.43</td>
<td>78.48± 12.79</td>
<td>3.62± 0.47</td>
<td>88.76± 11.62</td>
</tr>
</tbody>
</table>

Data presented as mean and standard deviation. RCB = Rice and Caupi Beans; RPB = Rice and Pontal Bean; RCBP = Rice, Caupi Bean and Pumpkin; RPBP = Rice, Pontal Bean and Pumpkin; RCBS = Rice, Caupi Bean and Sweet Potato; RPBS = Rice, Pontal Bean and Sweet Potato; PER = protein efficiency quotient; PER-R = Relative PER; NPR = net protein ratio; NPR-R = Relative NPR; TD = true digestibility; TD-R = relative true digestibility. Means with lower case in the same columns present significant difference by the Newman-Keuls test (p<0.05). n = 6, period of 14 day.
3.3 Effect of the staple food crops intake on the blood glucose

Animals fed diets with food combinations containing pontal beans, RPBP and RPBS, had lower blood glucose concentrations (p<0.05) than animals fed the control diet (Figure 2). That effect could be related to the higher soluble fiber content in the pontal bean (Table 1).

3.4 Effect of the staple food crops intake on the intestinal modulation

The control group and the RPBP group showed higher crypt depth (p<0.05) and longitudinal muscle layer than the other test groups (RPBS, RCBP, RCBS) (Table 3 and Figure 3).

The excretion of fecal fat of animals fed diets containing combinations of test foods was higher (p<0.05) than the animals

**Table 3.** Effect of staple food crops intake on gut morphometry of Wistar rats.

<table>
<thead>
<tr>
<th></th>
<th>Crypt Depth</th>
<th>Middle Muscle Layer</th>
<th>Longitudinal Muscle Layer</th>
</tr>
</thead>
<tbody>
<tr>
<td>Control</td>
<td>167.67 ± 23.07</td>
<td>44.44 ± 7.46</td>
<td>140.84 ± 14.16</td>
</tr>
<tr>
<td>RPBP</td>
<td>151.82 ± 4</td>
<td>33.94 ± 4.59</td>
<td>126.25 ± 6.19</td>
</tr>
<tr>
<td>RPBS</td>
<td>130.45 ± 6.71</td>
<td>36.77 ± 7.37</td>
<td>110.45 ± 12.08</td>
</tr>
<tr>
<td>RCBP</td>
<td>133.57 ± 11.04</td>
<td>39.04 ± 3.76</td>
<td>109.21 ± 7.13</td>
</tr>
<tr>
<td>RCBS</td>
<td>132.28 ± 15.94</td>
<td>41.61 ± 6.76</td>
<td>121.83 ± 10.11</td>
</tr>
</tbody>
</table>

Data presented as mean and standard deviation. RPBP = Rice, Pontal Bean and Pumpkin; RPBS = Rice, Pontal Bean and Sweet Potato; RCBP = Rice, Caupi Bean and Pumpkin; RCBS = Rice, Caupi Bean and Sweet Potato; Results in μm. Means with lower case in the same columns present significant difference by the Newman-Keuls test (p<0.05). n = 6, period of 14 days.
Biofortified foods present high protein quality and hypoglycemic action in rats

Among the test groups, the RCBP group showed the lowest (p<0.05) fecal fat excretion (Figure 4E). Furthermore, the RPBP group showed high (p<0.05) acetate concentration in the feces (Figure 4C).

4 Discussion

This study evaluated the effect of food combinations of beans and rice, target for biofortification, with high carotenoids content crops (sweet potato and pumpkin) on the protein quality, blood glucose and intestinal modulation in rats. The results showed that the all experimental groups presented a similar feed efficiency ratio (FER) (Supplementary Material Table S2), indicating that the diets were isocaloric. The PER and NPR indices of test groups did not differ of the casein group, indicating high protein quality. Among test groups it was observed that the RCB and RPBS groups had the lowest average of PER and NPR and RPB groups and RPBP showed the highest averages for these indices (Table 2). Thus, the combination of different plants studied, such as a cereal, a legume and a vegetable resulted in PER and NPR values similar to those of casein, indicating high biological value of the protein combinations.

RCB group, RCBP and RCBS showed higher values of TD and TD-R than the other test groups containing pontal beans in their combinations. Among the test groups the RPBP group showed lowest true digestibility (Table 2). The higher digestibility found in the groups containing caupi bean may be associated with lower contents of soluble dietary fiber and phenolic compounds compared to pontal beans. The soluble dietary fiber can form complexes in the intestinal content, prevent access of digestive enzymes, reducing protein digestibility (Kendall et al., 2010).

Animals fed with diets containing pontal beans, RPBP and RPBS, had lower blood glucose concentrations than animals fed with the control diet (Figure 2). This result may be associated with the content of dietary fiber, especially the soluble fraction, present in pontal beans which was higher than the caupi bean. The soluble fiber slows the glucose contact with the intestinal absorptive area, thereby reducing its absorption. Moreover, the fibers may increase peripheral insulin sensitivity (Aleixandre & Miguel, 2016; Stewart & Zimmer, 2018).

Furthermore, the RPBP group showed high acetate concentration in feces. The acetate can reduce the blood glucose by activating hepatic AMPK pathway which reduce expression of gluconeogenesis enzymes (Sakakibara et al., 2006). Another factor that may have contributed to the difference in blood glucose between the groups is the fact that these two groups showed lower concentrations of butyric acid in the feces (Figure 4B). The butyrate may increase the gene expression of GLUT2 in intestinal cells (Andrade et al., 2013; Chen et al., 2014), increased glucose uptake and thus leading to increased blood glucose levels.

The control group and the RPBP group showed higher crypt depth and longitudinal muscle layer than the other test groups (RPBS, RCBP, RCBS) (Table 3 and Figure 3). Conversely, Andrade et al. (2013) observed an increased thickness of the

Figure 4. Concentration of short chain fatty acids (SCFA) and fecal fat excretion in feces of animals fed casein and diets containing biofortified food combinations (n = 6), period of 28 days. (A) Propionic acid; (B) Butyric acid; (C) Acetic acid; (D) Total SCFA; (E) Total fecal lipid. RPBP = Rice + Pongal Bean + Pumpkin; RPBS = Rice + Pontal Bean + Sweet Potato; RCBP = Rice + Caupi Bean + Pumpkin; RCBS = Rice + Caupi Bean + Sweet Potato. Different letters indicate significant difference by the Newman-Keuls test (p <0.05).
muscle layers of the intestine of animals fed soy compared to animals fed casein. The result of this study may be related to higher total concentration of SCFA (Figure 4D) present in animal feces of control and RCBP group compared to other groups. It is known that SCFAs may modulate the proliferation of intestinal cells, increasing the crypt depth and intestinal muscle layer (Nilsson et al., 2013). Moreover, the control group received cellulose as dietary fiber source, and this carbohydrate has the ability to stimulate proliferation of intestinal cells (Rezaei et al., 2014).

The excretion of fecal fat of animals fed diets containing combinations of test foods was higher than the animals receiving the control diet (casein). Among the test groups, the RCBP group showed the lowest fecal fat excretion (Figure 4E). The increased excretion of fecal fat may be related to the high content of soluble dietary fiber present in the food testing. Soluble fiber slows digestion and absorption of fat, thereby increasing its excretion in the feces (Cherem & Bramorski, 2008). Moreover, Marques et al. (2015) in an in vitro study showed that caupi bean peptides were able to inhibit the HMG-CoA reductase and reduce the micellar solubilization of cholesterol, thereby increasing the fecal excretion of lipids. The rice peptides can bind to bile acids and reduce its availability for micelle formation, thus reducing the micellar lipid transport and increasing their fecal excretion (Cam & De Mejia, 2012). Thus, these results show that the combination of food target for biofortification presents high protein quality, and the combinations containing Pontal beans can promote a reduction in the blood glucose, hence it can present a hypoglycemic effect.

5 Conclusion

The combination of beans and rice, and these with biofortified pumpkin and sweet potato presented PER and NPR indices similar to control group, indicating high protein quality. The combination of pontal bean and pumpkin flours increased the total concentration of SCFA in the feces and the intestinal crypt depth compared to the other groups, which suggests an improvement in the intestinal function. Moreover, all food combinations increased fecal excretion of lipids, and combinations containing the pontal beans showed hypoglycemic action.

Overall, our results indicate that the foods from the Biofortification program not only provide an additional amount of micronutrient but it can also contribute to the protein intake of the risk population and potentially improve the glycemic status and intestinal function. Thus, the utilization of these foods may have a positive impact on the nutritional status and health of population at risk for micronutrient deficiency.

Acknowledgements

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References


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Supplementary Material

Supplementary material accompanies this paper.

Table S1. Food Composition and caloric density of experimental diets (g.100-1).

Table S2. Total food consumption, weight gain and food efficiency ratio (FER).

This material is available as part of the online article from http://www.scielo.br/IDSCIELO