Dose-Dependent Effects of Barley Cooked with White Rice on Postprandial Glucose and Desacyl Ghrelin Levels

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Summary White rice is an indispensable staple food in Japan, although it is a high glycemic index food. The objective of this study was to estimate how barley cooked with white rice might affect postprandial glucose, insulin and desacyl ghrelin concentrations as well as fullness. The study was conducted in randomized crossover design with nine healthy subjects. Blood glucose, insulin, free fatty acid and desacyl ghrelin concentrations and subjective levels of fullness and hunger were measured for 240 min after intake of glucose, white rice, 30% rolled barley (30BAR), 50% rolled barley (50BAR) and 100% rolled barley (100BAR) containing 75 g of available carbohydrate. Postprandial glucose and insulin levels were suppressed by intake of 30BAR, 50BAR and 100BAR comparing with those of white rice. Area under the curves of plasma glucose and insulin concentrations was reduced by barley intake in a dose-dependent manner. Although plasma desacyl ghrelin levels decreased postprandially, the degree of reduction was suppressed by barley intake in a dose-dependent manner. Postprandial desacyl ghrelin levels can be a sensitive biomarker of carbohydrate metabolism. The combination of white rice with barley plays a beneficial role in preventing and treating type 2 diabetes, obesity and other metabolic diseases.

Key Words: white rice, barley, insulin, desacyl ghrelin, fullness

Introduction

Recently, the number of cases of diabetes has rapidly increased worldwide, prompting renewed efforts to implement primary prevention measures. The glycemic index (GI) is a ranking of carbohydrates based on their immediate effects on blood glucose levels [1]. Epidemiologic studies demonstrated that the GI of the diet might be an important factor in preventing type 2 diabetes [2–4]. Japanese and other Asian cuisines, combining white rice (staple food) with another dish, are characterized by low fat intake. A low-fat diet has beneficial effects for the prevention or

treatment of diseases related to insulin resistant syndrome [5]. However, white rice belongs to the group of high GI foods [6]. A previous study indicated that the relative glycemic effects of mixed meals are mainly influenced by GI of the staple food [7]. Furthermore, replacement of carbohydrate in the dish with a low GI diet improved early insulin secretion compared with a high GI diet in metabolic syndrome subjects [8]. These data suggest that staple carbohydrate-rich foods with low GI are needed for the prevention of diabetes. Several studies investigating the effects of mixtures of high-GI and low-GI grains have been reported, but most of these investigations used bread [9, 10], rather than white rice, as the test meal. Barley is a typical low GI food, but it is difficult to ingest barley alone because its texture is very hard. Although white rice is sometimes cooked with 20-30% barley, which can be easily ingested, it is unclear whether this combination affects postprandial

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glucose metabolism.

Ghrelin is one of the hormones involved in controlling body weight, insulin secretion and appetite regulation. The plasma concentration of ghrelin is known to gradually increase before a meal and then decrease immediately after a meal [11-13]. Ghrelin has two major molecular forms: acyl ghrelin and desacyl ghrelin [14]. Plasma desacyl ghrelin accounts for more than 90% of total circulating ghrelin [14, 15], whereas acyl ghrelin is relatively unstable and is rapidly converted into desacyl ghrelin or smaller fragments. This study was undertaken to clarify the effect of barley mix with white rice on postprandial plasma glucose, insulin and desacyl ghrelin levels in healthy subjects.

Experimental Procedures

Subjects

Six healthy men and 3 healthy women were recruited in this study. The study was performed after obtaining written

Table 1. Characteristics of the subjects

Characteristics	Values		
F/M	3/6		
Age (years)	26.7 ± 1.1		
Body weight (kg)	61.1 ± 2.8		
BMI (kg/m ²)	22.0 ± 0.7		
Body fat ratio (%)	18.5 ± 2.6		
Triacylglycerol (mg/dl)	77.0 ± 10.5		
Total cholesterol (mg/dl)	162.9 ± 8.2		
HDL cholesterol (mg/dl)	55.6 ± 2.3		
Fasting plasma glucose (mmol/l)	4.9 ± 0.1		
HbA1c (%)	4.8 ± 0.1		
BUN (mg/dl)	11.3 ± 0.8		
Creatinine (mg/dl)	0.8 ± 0.05		
AST (IU/l)	15.6 ± 0.8		
ALT (IU/l)	15.0 ± 2.1		

Values are means \pm SEM.

HDL, high density lipoprotein; BUN, blood urea nitrogen; AST, aspartate aminotransferase; ALT, alanine aminotransferase.

 Table 2.
 Energy and macronutrient composition of the test meals

informed consent from all of the subjects, and was approved by the Ethics Committee of the University of Tokushima. The clinical and biological characteristics of the subjects are shown in Table 1. The mean values \pm SEM of age and body mass index (BMI) were 26.7 ± 1.1 years and 22.0 ± 0.7 kg/m², respectively.

Study protocol

The design of this experiment was a randomized and crossover study. The experiment was conducted so that each test day was separated by a washout period of 7 days. All the subjects were asked to avoid heavy exercise and intake of alcohol 24 h before the day of the study. The subjects were instructed to eat and drink the same prescribed foods at 2000 h before a test day. After an overnight fast, venous blood samples were collected at 0840 h and each subject was asked to complete a short questionnaire rating their appetite using a visual analog scale (VAS). The subjects were provided with their test meals (breakfast) at 0900 h and were required to consume each test meal within 20 min. Blood samples were collected at 0 (i.e., immediately before) and 30, 45, 60, 90, 120, 180 and 240 min after the meal. The blood samples were used to analyze glucose, insulin, free fatty acid (FFA) and desacyl ghrelin levels.

Test meals

Five different test meals were used: glucose (GL), white rice (WR), 30% rolled barley (30BAR), 50% rolled barley (50BAR) and 100% rolled barley (100BAR) containing 75 g available carbohydrate. White rice and rolled barley were boiled, and then served. The amounts of available carbohydrate in 30BAR were 52.5 g and 22.5 g, and in 50BAR were 37.5 g and 37.5 g from WR and BAR, respectively (Table 2). Except for GL, all other test meals were made up in water to give a final total volume of 650 ml.

Visual analogue scores (VAS)

VAS was used to measure subjective feelings of hunger and fullness. VAS consisted of 100 mm vertical lines with short questionnaire anchored at each end that expressed the most positive or most negative sensation. Subjects drew a

	GL	WR	30BAR	50BAR	100BAR
Meal volume (ml)	225	375	398	413	450
Water volume (ml)		275	252	237	200
Total volume (ml)	225	650	650	650	650
Energy (kJ)	1255	1435	1460	1477	1515
Protein (g)	0.0	4.6	5.4	5.9	7.2
Fat (g)	0.0	0.9	1.1	1.3	1.7
Available carbohydrate (g)	75.0	75.0	75.0	75.0	75.0
Dietary fiber (g)	0.0	0.5	3.2	4.9	9.2

horizontal line at the point on the vertical line that corresponded to their sensation.

Analysis methods

Plasma and serum samples were separated and stored at -80°C until use. Plasma glucose levels were measured by an enzymatic method (hexokinase method) (Sino-test, Tokyo, Japan). Serum insulin levels were measured by chemiluminescent enzyme immunoassay (EIKEN, Tokyo, Japan). Serum FFA levels were measured enzymatically (EIKEN, Tokyo, Japan). Plasma desacyl ghrelin levels were measured by enzyme linked immunosorbent assay (Mitsubishi Kagaku latoron, Tokyo, Japan). The incremental area under the curve (AUC) for plasma glucose and insulin were calculated for a 240-min period after each test meal.

Statistical analysis

Data are shown as mean \pm SEM. Firstly, the statistical difference was determined by 1-way ANOVA, with Fisher's protected least significant difference as a posthoc test to compare 3 test meals (GL, WR and 100BAR). Secondly, the statistical difference was determined to compare white rice with combining white rice and barley in varying proportions. Data of 30BAR and 50BAR were compared with WR respectively. Groups were compared for differences using Student's *t* test for paired comparisons. Correlation was calculated using stepwise linear regression by combining data from the 4 groups. Note that GL was not included in this analysis because this test meal had a different volume. All of the statistical analyses were performed with the Stat View 5.0 (SAS Institute, Inc., Cary, NC). *p* value less than 0.05 was considered significant in all analyses.

Results

Effects of GL, WR and 100BAR intake on postprandial glucose, insulin and FFA levels

Plasma glucose levels at 30, 45 and 60 min in 100BAR were significantly lower than in GL (p < 0.01; 30 and 45 min, p<0.05; 60 min) and WR (p<0.01; 45 min, p<0.05; 30 and 60 min). No differences were observed between GL and WR (Fig. 1a). Serum insulin levels increased after each meal reaching a peak value at 45 min for GL and WR, and at 30 min for 100BAR. Serum insulin levels for 100BAR at 30–120 min were significantly lower than for GL (p < 0.01; 45, 60, 90 and 120 min, p<0.05; 30 min). At 60 and 90 min serum insulin levels for 100BAR were significantly lower than those in WR (p < 0.05, respectively). Serum insulin level at 60 min for WR was markedly lower than for GL (p < 0.05) (Fig. 1b). Serum FFA levels for 100BAR fell more slowly than for GL and WR. At 45 and 120 min serum FFA levels for 100BAR were significantly higher than for GL (p < 0.05; respectively), and significantly higher than GL and WR at 180 min (p<0.01; respectively). Serum FFA level for GL quickly increased and returned to fasting level at 240 min, and were significantly higher than those for WR and BAR (p<0.01; respectively) (Fig. 1c).

In the postprandial early stage (0–30 and 0–60 min), plasma glucose AUCs for 100BAR were significantly lower than those for GL (p<0.01) and WR (p<0.05). At 0–120, 0–180 and 0–240 min, plasma glucose AUCs tended to be lower for 100BAR compared to those for GL and WR, although this observation was not statistically significant (Fig. 1d). For 100BAR, serum insulin AUCs were significantly lower than those for GL at all time periods (0–30, 0–60, 0–120, 0–180, 0–240 min) (p<0.01). At 0–30, 0–60, 0–120 and 0–180 min, serum insulin AUCs for WR were significantly lower than those for GL (p<0.05; 0–30, 0–60 and 0–180 min, p<0.01; 0–120 min). Also, serum insulin AUCs at 0–120 and 0–180 min for 100BAR were significantly lower than those for WR (p<0.01; respectively) (Fig. 1e).

Dose-dependent effects of barley intake on postprandial glucose, insulin and FFA levels

Plasma glucose levels for 30BAR and 50BAR increased more slowly than for WR. Indeed, the plasma glucose level at 45 min for 50BAR and 30BAR was significantly lower (p < 0.05) or lower (p = 0.07) than that for WR, respectively (Fig. 2a). Serum insulin levels increased after each meal, reaching a peak value at either 45 min for WR or at 30 min for 30BAR and 50BAR. Serum insulin levels for 30BAR and 50BAR immediately dropped from peak values, and were significantly lower than that for WR at 60 min (p < 0.05; respectively) (Fig. 2b). Serum FFA levels for 50BAR fell more slowly than those for WR; see values at 120 and 180 min in Fig. 2c (p < 0.05; respectively). The plasma glucose AUCs and insulin AUCs were slightly lower for the 30BAR and 50BAR groups compared to the WR group, although these differences were not statistically significant (Fig. 2d, e).

Effects of GL, WR, 30BAR, 50BAR and 100BAR intake on postprandial desacyl ghrelin levels

Postprandial plasma desacyl ghrelin levels after ingestion of 100BAR remained unchanged from the fasting level. By contrast, postprandial plasma desacyl ghrelin levels after intake of GL and WR were markedly suppressed between 30 and 180 min. Specifically, at 60 min the plasma desacyl ghrelin level for WR was lower than for 100BAR (p<0.05); at 120 min, plasma desacyl ghrelin for GL was lower than for WR (p<0.05) (Fig. 3a). The degree of reduction in the level of plasma desacyl ghrelin was mirrored by the postprandial insulin concentration but not by the glucose concentration.



Fullness and hunger

Peak fullness scores occurred 30 min after each test meal. The peak score for 100BAR was significantly higher than for GL (p<0.05). Fullness score at 180 and 240 min after ingestion of 100BAR was significantly higher than after ingestion of GL and WR at the corresponding time points (p<0.01; respectively). Fullness scores at 240 min after intake of GL and WR returned to fasting level values, but remained above fasting level for 100BAR (Fig. 4a). Fullness score at 45 min for 50BAR was significantly higher than for WR and 30BAR (p<0.05; respectively). For 30BAR and 50BAR, high fullness scores at 180 and 240 min were significantly higher than for WR (p<0.05; 180 min, p<0.01; 240 min) (Fig. 4b). Hunger scores showed reverse patterns of fullness scores (Fig. 4c, d).

Correlation

Significant variables were entered into a stepwise linear



regression model in which the difference from baseline in plasma glucose levels (Δ plasma glucose), serum insulin levels (Δ serum insulin) and fullness scores (Δ fullness) were used as independent variables and difference from baseline in desacyl ghrelin levels (Δ plasma desacyl ghrelin) were used as a dependent variable. This analysis revealed that Δ serum insulin was an independent predictor of Δ plasma desacyl ghrelin (Δ plasma desacyl ghrelin = 65.877– 3.886* Δ serum insulin, $r^2 = 0.074$, p < 0.0001).

Discussion

In this study, plasma glucose level at 45 min for 50BAR and 30BAR was significantly lower or lower, and serum insulin levels for 50BAR and 30BAR were significantly lower than that for WR at 60 min. Because viscous fibers have been shown to reduce postprandial glucose and insulin levels [16-18], these effects might be due to the viscous fiber component of barley. Among the various grains, barley



Fig. 3. Postprandial plasma desacyl ghrelin level after ingestion of glucose, white rice, 100% barley (a), and white rice, 30% barley, 50% barley (b). X, glucose (GL); open circle, white rice (WR); closed circle, 100% barley (100BAR). *p<0.05, GL vs 100BAR; †p<0.05, GL vs WR; ‡p<0.05, 100BAR vs WR. Open circle, white rice (WR); closed triangle, 30% barley (30BAR); closed square, 50% barley (50BAR). *p<0.05, WR vs 30BAR; †p<0.05, WR vs 50BAR. Mean values with their standard errors.

is highly viscous in character. Indeed, the viscosity of barley is mainly derived from the β -glucan content [19]. β -glucan has been reported to suppress postprandial glucose levels in a dose-dependent manner [20]. A possible mechanism to explain these results might involve delayed or reduced carbohydrate absorption from the gut [21]. The plasma glucose AUCs and insulin AUCs were not statistically significant for the 30BAR and 50BAR compared to the WR. Because it is calculated for the total amount of increase, AUC is difficult to detect the difference with each time point.

In most previous studies using a mixture of high-GI and low-GI grains, the glucose and insulin levels were suppressed compared with those of high-GI when more than 50% of low-GI grain was used [9, 10]. Furthermore, the addition of toppings to a carbohydrate-rich food had the effect of reducing the glycemic response [22-24]. The effect was particularly notable for high GI carbohydrate-rich food. The GI of white rice is higher than that of wheat flour bread, barley flour bread or barley porridge [6, 25], which were used as high-GI grains in previous studies [9, 10]. In this study, the combination of white rice and barley and even 30BAR, showed a definite effect on the suppression of postprandial insulin peak value. And it tended to suppress of postprandial glucose peak value. By contrast, the effect of the combination of white rice and pre-germinated brown rice was not observed [26]. This finding might be explained by the higher GI of brown rice compared with that of barley [27]. Therefore, it is conceivable that GI values and combination rates of each food could be important factors in terms of this phenomenon.

The insulin level peaked earlier and fell more rapidly for 30BAR, 50BAR and 100BAR than for the GL and WR group of subjects. Because the rise in plasma glucose for 30BAR, 50BAR and 100BAR was smaller than for GL and WR, an excessive amount of insulin secretion might be suppressed. Also, it was reported that intake of cereal fibers accelerated the response of the acute glucose-dependent insulinotropic polypeptide (GIP) and insulin response, result in the reduction of postprandial glucose and insulin values the following day [28]. Repeated lower postprandial insulinemic responses may prevent the onset of type 2 diabetes by preventing damage to the β cell function. Recovery of β cell function or a decrease in insulin resistance will improve early insulin secretion over the long term in patients with type 2 diabetes [29].

We also studied the effect of the ingestion of a carbohydrate-enriched breakfast on the plasma ghrelin level. Percentage change in ghrelin concentration between 30 and 180 min were negatively correlated with the percentage change in insulin but not with the observed changes in glucose level [30]. Insulin levels after ingestion of a high fiber meal were lower than those after ingestion of a low fiber meal. Furthermore, postprandial reductions of desacyl ghrelin levels were suppressed after intake of a high fiber meal. A stepwise linear regression model revealed that plasma desacyl ghrelin levels were predictors of serum insulin levels but not of plasma glucose levels. In contrast, an arabinoxylan fiber-enriched meal affects serum ghrelin levels but not postprandial insulin responses in healthy subjects [31]. These apparently conflicting results might be due to differences in the amount of carbohydrate used in



Fig. 4. Postprandial fullness after ingestion of glucose, white rice, 100% barley (a), white rice, 30% barley, 50% barley (c), and postprandial hunger after ingestion of glucose, white rice, 100% barley (b), white rice, 30% barley, 50% barley (d). X, glucose (GL); open circle, white rice (WR); closed circle, 100% barley (BAR). * p<0.05, GL vs 100BAR; †p<0.05, GL vs WR; ‡p<0.05, 100BAR vs WR. Open circle, white rice (WR); closed triangle, 30% barley (30BAR); closed circle, 50% barley (50BAR). *p<0.05, WR vs 30BAR; †p<0.05, WR vs 50BAR. Mean values with their standard errors.

each study (i.e., 50 g in previous study compared with 75 g in our study). Therefore, postprandial desacyl ghrelin levels can act as a sensitive biomarker for the degree of postprandial carbohydrate metabolism.

Intake of 30BAR, 50BAR and 100BAR resulted in prolonged postprandial fullness and in a less pronounced reduction of desacyl ghrelin level compared with the WR group. These findings are consistent with previous work, which concluded postprandial satiety is maintained for longer after intake of a high fiber meal compared to a low fiber meal [32]. There are several inconsistent reports between postprandial ghrelin levels and fullness and satiety. Ghrelin responds rapidly and dose-dependently to carbohydrate intake and is positively correlated with subjective measures of hunger and negatively correlated with fullness [30]. In contrast, postprandial alterations of ghrelin level seem not to play an important role in the acute regulation of satiety [33, 34]. Furthermore, conflicting reports suggest intracerebroventricular administration of desacyl ghrelin to rats can result in either increased [35] or decreased [36, 37] food intake. Thus, the relationship between postprandial ghrelin level and fullness is still obscure and needs to be clarified in further studies. However, it is important to keep in mind that prolonged fullness after intake of 30BAR, 50BAR and 100BAR may induce a reduction of energy intake and improve glucose metabolism.

In conclusion, results from the present study suggest that the combination of white rice with barley plays a beneficial role in the prevention and treatment of type 2 diabetes, obesity and other metabolic diseases.

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Abbreviations

GI, glycemic index; BMI, body mass index; VAS, visual analog scale; FFA, free fatty acid; GL, glucose; WR, white rice; 30BAR, 30% rolled barley; 50BAR, 50% rolled barley; 100BAR, 100% rolled barley; AUC, area under the curve; HDL, high density lipoprotein; BUN, blood urea nitrogen; AST, aspartate aminotransferase; ALT, alanine aminotransferase.

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