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3 **Association between Dietary Patterns and Serum Adiponectin: A**
4 **Cross-Sectional Study in a Japanese Population**

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19 **Association between Dietary Patterns and Serum Adiponectin: A**
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22 The aim of this study was to evaluate the associations between dietary pattern, adiponectin, and
23 insulin resistance. The study population consisted of 612 men and women aged 35-69 years old
24 who had participated in the baseline survey of Japan Multi-Institutional Collaborative Cohort (J-
25 MICC) Study in Tokushima Prefecture. Diets and lifestyle related variables were assessed by
26 questionnaires. Multiple regression analyses were used to analyse the relations between dietary
27 patterns and high molecular weight adiponectin. For further analysis, path analysis was used to
28 test the hypothesized model of association between dietary pattern, serum adiponectin, and
29 insulin resistance. The result showed that higher score of bread and dairy pattern was directly
30 associated with increased serum level of adiponectin in women, which was inversely related to
31 Homeostasis Model Assessment of Insulin Resistance. In conclusion, higher consumption of
32 bread and dairy products, and low intake of rice may be associated with increased serum
33 adiponectin in women.

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35 Keywords: adiponectin, dairy, dietary pattern, insulin resistance, path analysis

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42 **Introduction**

43 Metabolic syndrome can be defined as a cluster of risk factors, such as
44 abdominal obesity, hypertension, dyslipidaemia, and glucose intolerance (Balkau &
45 Charles 1999; Expert Panel on Detection, Evaluation, and Treatment of High Blood
46 Cholesterol In Adults 2001). The number of people who suffer from this condition is
47 increasing worldwide (Cameron et al. 2004). Most people who suffer from metabolic
48 syndrome have insulin resistance (Grundy et al. 2004) and are at increased risk of type
49 II diabetes and cardiovascular diseases (Alberti et al. 2009). To measure the insulin
50 resistance in large population-based epidemiological investigations, the usage of
51 surrogate assessment of insulin resistance named Homeostasis Model Assessment of
52 Insulin Resistance (HOMA-IR), has been proven to be an effective tool (Antuna-Puente
53 et al. 2011).

54 Adiponectin is a glycosylated adipokine selectively secreted from adipocytes
55 and its function is related to glucose uptake, beta-oxidation, regulation of insulin, and
56 play a role in the regulation of inflammation (Xu et al. 2007; Mittal 2008). Lower
57 concentrations of plasma adiponectin are associated with decreased insulin sensitivity,
58 whereas individuals with high concentrations of serum adiponectin were less frequent to
59 develop type II diabetes (Lindsay et al. 2002) and cardiovascular disease (Pischon et al.
60 2004).

61 In human blood, adiponectin circulates in different isoforms: high molecular
62 weight (HMW), middle molecular weight (MMW), and low molecular weight (LMW)
63 (Pajvani et al. 2003; Waki et al. 2003). Compared with plasma total adiponectin, HMW
64 adiponectin is more useful for predicting insulin resistance and metabolic syndrome
65 (Hara et al. 2006).

66 From the standpoint of preventing metabolic syndrome, type 2 diabetes and
67 cardiovascular diseases, understanding the factors that favourably modify serum
68 adiponectin concentrations would be an important priority. Even though a previous
69 study from Japan suggested that none of the nutrient intakes had significant association
70 with serum adiponectin concentration, (Murakami et al. 2013) another study has shown
71 the significant relation between Japanese traditional dietary pattern and serum
72 adiponectin (Guo et al. 2012). Meanwhile, studies from the U.S. showed significant
73 associations between adherence to Mediterranean dietary pattern (Mantzoros et al.
74 2006) or healthy eating pattern (Fargnoli et al. 2008) and adiponectin.

75 In this study, we tried to investigate the association between major dietary
76 pattern and HMW adiponectin. Furthermore, we analysed the association between
77 dietary pattern, serum adiponectin, and insulin resistance by using path analysis.

78

79 **Materials and methods**

80 *Study population*

81 The population for the present study consisted of 697 participants aged 35 to 69
82 years old, who were enrolled in the baseline survey of the Japan Multi-Institutional
83 Collaborative Cohort (J-MICC) Study in Tokushima Prefecture, Japan. Details about J-
84 MICC Study have been described in the previous report (Hamajima 2007). In short, the
85 aim of the J-MICC Study was to examine the prospective associations of lifestyle and
86 genetic factors and their interactions with the risk of lifestyle-related diseases. We
87 distributed approximately 98,700 leaflets explaining the objective and method of the J-
88 MICC Study in all over Tokushima city, with a total population of 264,500. Within July

89 25th 2012 to February 27th 2013, there were 697 subjects who read the leaflets and
90 attended the health check-ups performed by our research team. Written inform consent
91 was obtained from participants after we explained about the outline and the objectives
92 of this study. The participation rate was difficult to calculate because of the recruitment
93 method. The study protocol was approved by the review boards of Nagoya University
94 School of Medicine (an affiliate of the former principal investigator, Dr. Nobuyuki
95 Hamajima), Aichi Cancer Center Research Institute (affiliated with the present principal
96 investigator, Dr. Hideo Tanaka), and Tokushima University Hospital.

97 We excluded participants who had history of ischemic heart disease (n = 21),
98 stroke (n = 9), or diabetes mellitus (n = 28), or who received medical treatment with
99 anti-diabetic drugs (n = 22). Participants might be excluded for one or more reasons.
100 Subjects with missing values of serum adiponectin (n = 9) and other variables (n = 24)
101 were also excluded from the study. These exclusions gave final study population of 612
102 participants (437 women and 175 men) for statistical analysis.

103

104 *Questionnaire*

105 In this study, subjects were asked to fill out the self-administered questionnaire
106 about dietary habits, current and previous diseases, medication and supplements
107 consumption, physical activity, and smoking and drinking habits. Questionnaires were
108 sent to participants approximately 2 weeks before the health check-up. Validated short
109 food frequency questionnaire (FFQ) was used for dietary evaluations (Tokudome et al.
110 2004; Tokudome et al. 2005; Goto et al. 2006; Imaeda et al. 2007). Dietary intake
111 information was collected by asking the participants about how often they consumed 46
112 foods and 9 beverages over the past year. Consumption of rice, bread, and noodle at

113 breakfast, lunch and dinner were divided into six categories: rarely, 1-3 times per month,
114 1-2 times per week, 3-4 times per week, 5-6 times per week, and every day. Other foods
115 intake including coffee and green tea, was categorized into 8 categories: rarely, 1-3
116 times per month, 1-2 times per week, 3-4 times per week, 5-6 times per week, once per
117 day, 2 times per day, and ≥ 3 times per day. Beverages consumption was divided into
118 seven categories: rarely, less than 2 cups per week, 3-4 cups per week, 5-6 cups per
119 week, 1-2 cups per day, 3-4 cups per day, ≥ 5 cups per day. Information on the portion
120 size was collected only for staple foods. Average daily consumption of energy and
121 selected nutrients were computed using a program developed by Department of Public
122 Health, School of Medicine, Nagoya City University (Tokudome et al. 2004; Tokudome
123 et al. 2005). A relative validation study was completed by comparing the intake of
124 energy and 26 nutrients evaluated using this FFQ and 3 day-weighed diet records (3d-
125 WDRs) as a reference. Deattenuated, log-transformed, and energy-adjusted Pearson's
126 correlation coefficients for 26 nutrients intake distributed from 0.10-0.86 (Tokudome et
127 al. 2005). This FFQ had substantially high one-year interval reproducibility values for
128 consumption of foods and nutrients assessment (Imaeda et al. 2007).

129 Smoking status was self-reported and classified into current smoker, past smoker,
130 and never, whilst drinking habit was categorized into 3 groups, current drinker, past
131 drinker, and never. Physical activity during leisure time was estimated by multiplying
132 the frequency and the average duration of light exercise (such as walking and golf are
133 3.4 metabolic equivalents [METs]), moderate exercise (such as jogging, swimming and
134 dance are 7.0 METs), and vigorous-intensity exercise (such as marathon is 10.0 METs).
135 The three levels of exercise were summed to obtain the MET-hours/week.

136

137 *Anthropometric and biochemical measurement*

138 Data on anthropometric measurement (height, weight, and waist circumference),
139 fasting plasma glucose, and insulin were obtained at the time when participants came
140 for the health check-up performed by our research team. Participants were requested not
141 to eat breakfast and received medical check-up between 8.00 AM to 11.00 AM. Venous
142 blood samples were collected from all participants and serum was separated within 3
143 hours. HMW adiponectin levels were assessed by the external laboratory using latex
144 turbidimetric immunoassay (SRL, Tokyo, Japan). Serum insulin was measured using a
145 chemiluminescence immunoassay (BML Inc., Tokyo, Japan).

146 Homeostatic Model Assessment (HOMA) is a method for assessing β -cell
147 function and insulin resistance from basal (fasting) glucose and insulin or C-peptide
148 concentrations. The equation used to calculate HOMA-IR was $\text{insulin } (\mu\text{U/mL}) \times$
149 $\text{plasma glucose (mg/dL)}/405$ (Matthews et al. 1985).

150 Body mass index (BMI) was calculated as weight (kg) divided by the square of
151 height (m^2).

152

153 *Statistical Analysis*

154 To extract the dietary pattern, we used principal component analysis (PCA) by
155 PRINCOMP procedure from SAS software package (version 9.4). The components of
156 dietary patterns obtained from PCA reflect the combinations of foods consumed by each
157 participant. In determining the number of dietary patterns to be retained, we considered
158 the eigenvalues (≥ 1.0), scree test, and interpretability. Retained dietary patterns were
159 named based on the highest factor loadings on each pattern. We analysed the relation

160 between retained dietary pattern scores and serum adiponectin in men and women
161 separately, by using multiple regression analysis adjusted for potential confounders.
162 Model 1 adjusted for age. Model 2 included all variables from model 1 and additionally
163 adjusted for energy intake, physical activity, drinking habit, smoking habit, and
164 menopausal status (in women only). Model 3 included all variables from model 2 and
165 further adjusted for BMI. Hypothetical path diagram was constructed based on the result
166 of multiple regression analysis and the results of prior studies.

167 To examine the baseline characteristics, continuous variables are shown as
168 median (25th and 75th percentiles) and categorical variables are presented as the count
169 and proportion. The differences across the quartiles of dietary pattern scores were
170 examined using the Chi-square test, Kruskal-Wallis test, or Fisher's exact test.

171 The hypothetical path diagrams were analysed using path analysis, performed by
172 IBM[®] SPSS[®] AMOS[™] (Version 22). Path analysis is an analysis involving
173 assumptions about the direction of causal relationships between linked sequences and
174 configuration of variables (Porta 2008). Listwise deletion method was used to handle
175 the missing data. The covariates included in the path analysis were the available
176 predictors of adiponectin, dietary pattern, and BMI: age, energy intake, physical activity,
177 drinking habit, smoking habit, and menopausal status. The assessment of model fit was
178 performed using the following parameters: chi-square test (χ^2), RMSEA (Root Mean
179 Square Error of Approximation), CFI (Comparative Fit Index), and NFI (Normed-Fit
180 Index).

181 Chi-square is a traditional measure for estimating the overall model fit and
182 evaluate the discrepancy between the sample covariance matrix and the fitted
183 covariance matrix (Hu & Bentler 1999). The insignificant result at 0.05 threshold is

184 necessary for good model fit (Barrett 2007). Nevertheless, χ^2 is sensitive to sample size.
185 As a result, χ^2 nearly always rejects the model when the sample size is large. Therefore,
186 it is necessary to use the incremental fit indices as well (Choi et al. 2014). Incremental
187 fit indices show the relative improvement of hypothesized model compared to the null
188 model – typically no correlation among observed variables was assumed (Kline 2011).

189

190 **Results**

191 Five dietary patterns were extracted by PCA. These patterns accounted for
192 34.09% of the total variance of food intakes. There were 9 other factors showing
193 eigenvalue ≥ 1.0 , but the interpretation of those components was difficult. The scree plot
194 (see supplementary figure 1) suggested five factors to be retained. Table 1 shows the
195 factor loadings of each food item in the extracted dietary patterns. The first dietary
196 pattern (eigenvalue 7.24) was named vegetable pattern because it had high factor
197 loadings for vegetables. The second principal component (eigenvalue 2.59) was named
198 high-fat pattern because of the high loadings for ham, mayonnaise, and deep fried foods.
199 The third pattern (eigenvalue 2.21) had positive factor loadings for squid, shellfish, and
200 cod roe – thus called seafood pattern. The fourth pattern (eigenvalue 1.99) was named
201 bread and dairy pattern because of the high loadings for bread, butter, milk, and yogurt,
202 and negative loading for rice. The fifth pattern (eigenvalue 1.65) had high loadings for
203 milk, egg, tofu, and natto (fermented soybeans); we identified this as protein pattern.

204 Table 2 shows the associations between extracted dietary patterns and serum
205 adiponectin in men and women by using multiple regression analyses. The result
206 showed that bread and dairy pattern scores had positive significant association with
207 serum adiponectin only in women ($p = 0.004, 0.009, \text{ and } 0.015$ for model 1, 2, and 3,

208 respectively). Addition of a product term between bread and dairy pattern score and
209 menopausal status to model 2 and model 3 showed no evidence of significant
210 interaction ($p = 0.41$ for model 2 and $p = 0.08$ for model 3, data not shown).

211 Table 3 shows the baseline characteristics of the study participants according to
212 quartile of bread and dairy pattern score. Participants who had higher scores of bread
213 and dairy pattern were more likely to be older, female, had lower energy intake, took
214 part in more leisure time physical activity, and had higher concentrations of HDL
215 cholesterol and serum adiponectin.

216 Figure 1 presents the path analysis model for women. This model showed good
217 fit model to the data, with $\chi^2 = 18.634$, d.f. = 12, $p = 0.098$, standardized root mean
218 square residual (SRMR) = 0.019, comparative fit index (CFI) = 0.992, and root mean
219 square error of approximation (RMSEA) = 0.036. Significant association could be
220 found between bread and dairy pattern and adiponectin ($p = 0.017$). Hereinafter, a
221 negative association was seen between adiponectin and HOMA-IR ($p = 0.004$). Further,
222 BMI was significantly inversely related to adiponectin and significantly positively
223 influenced the HOMA-IR. The pathways of the path analysis model for Figure 1 is
224 shown in Table 4.

225 The analyses of calcium, vitamin D, and potassium and their associations with
226 adiponectin separately are shown in Supplementary Table 1. Significant association
227 could only be found between potassium and adiponectin in women ($p = 0.019$, 0.004,
228 and 0.039 for model 1, 2, and 3, respectively).

229 We found positive significant associations of bread and dairy pattern with
230 adiponectin when waist circumference was used instead of BMI in model 3
231 (Supplementary Table 2) ($p = 0.030$).

232 In the supplementary figure 2, waist circumference was used instead of BMI for
233 the path analysis. The model showed a good fit, with $\chi^2 = 20.511$, d.f. = 12, $p = 0.058$,
234 SRMR = 0.020, CFI = 0.989, and RMSEA = 0.040. Similar to the result in figure 1,
235 bread and dairy pattern was positively significantly associated with adiponectin ($p =$
236 0.028), followed with inverse significant association between adiponectin and HOMA-
237 IR ($p < 0.001$). Menopausal status was found to be positively significantly related to
238 waist circumference. Furthermore, waist circumference was inversely significantly
239 associated with adiponectin and positively related to HOMA-IR. Supplementary table 3
240 shows the pathways of path analysis model for supplementary figure 2.

241

242 **Discussion**

243 In our study, derived dietary patterns were comparable to those of other studies
244 in Japanese populations. For instance, the Japan Public Health Center-based Prospective
245 Study (Nanri et al. 2013) extracted prudent, westernized, and traditional dietary patterns
246 from its population. In another Japanese population, Nanri et al. (Nanri et al. 2008)
247 obtained the healthy, high-fat, seafood, and westernized breakfast dietary pattern – with
248 bread, margarine, and yogurt as part of its components.

249 Our study showed that dietary pattern characterized by high intake of bread and
250 dairy products such as milk, butter, and yogurt, and low intake of rice was significantly
251 related to a higher concentration of serum adiponectin, which is associated with the
252 lower value of HOMA-IR in women. From our analysis, we infer that bread and dairy
253 intake is one of the determinants of serum adiponectin, which has beneficial effects on
254 the improvement of insulin resistance. Our result is in line with earlier systematic
255 review study which revealed that higher intake of dairy products may have a positive

256 effect on insulin sensitivity (Turner et al. 2015). However, in our study population, the
257 indirect effect of the bread and dairy pattern score on HOMA-IR was considered to be
258 rather small, as suggested by the product of the two standardized estimates ($0.11 \times [-$
259 $0.13]$).

260 The positive association between dairy products and serum adiponectin may be
261 because of its contents, namely milk fat, vitamin D, calcium, potassium, whey protein,
262 magnesium, or a combination of these components. Previous study showed that serum
263 25(OH)D was positively related to serum adiponectin (Vaidya et al. 2012). Meanwhile,
264 dietary calcium may take part in the regulation of oxidative and inflammatory stress
265 (Zemel & Sun 2008)– which negatively correlated with adiponectin (Furukawa et al.
266 2004). Increased magnesium intake may also have a favourable effect on adiponectin
267 (Cassidy et al. 2009). Nonetheless, when we analysed the intake of calcium, vitamin D,
268 and potassium and their associations with adiponectin separately, we could find
269 significant association between only potassium and adiponectin (Supplementary Table
270 1). We consider that it is not calcium, vitamin D, or potassium per se that may increase
271 the concentration of serum adiponectin, but the combination of those nutrients with
272 other components within dairy products. In addition, when we analysed the correlation
273 between main food items in bread and dairy pattern and adiponectin – only bread and
274 butter were positively significantly correlated with adiponectin. As for the
275 macronutrients intake, we did not find any significant correlation (data not shown).
276 Possibly, the effect of single food items or nutrients were too small to be detected –
277 compared to when they were measured together as a dietary pattern. In addition, the
278 intercorrelation among individual food items and nutrients makes it hard to analyse their
279 effect separately (Hu 2002).

280 Different from our results, one study stated that only low-fat dairy products were
281 associated with increased serum adiponectin in a Japanese population (Niu et al. 2013).
282 In our FFQ, we did not separate low- and high-fat dairy products. Nevertheless, the
283 validation study of our FFQ using diet record showed that the proportion of low-fat
284 dairy products in this population was low (data not shown). Further studies may be
285 needed to clarify whether only low-fat dairy products are associated with high serum
286 levels of adiponectin.

287 In the bread and dairy pattern, the factor loading of rice was negative. This may
288 be because people consume bread as the staple food instead of rice. Separate analysis
289 between rice intake and adiponectin showed that there was an inverse significant
290 association between rice intake and adiponectin. Therefore, low intake of rice may in
291 part contribute to the positive association between bread and dairy pattern and
292 adiponectin. The possible reason for this is that rice, in general, has high glycaemic
293 index/load and raise the blood glucose level (Foster-Powell et al. 2002). It has been
294 suggested that glycaemic index and glycaemic load are inversely associated with
295 adiponectin (Qi et al. 2006).

296 The result of multiple regression analysis for the association between bread and
297 dairy pattern score and serum adiponectin in women (Table 2) showed that there was no
298 substantial difference in the results between model 2 (not adjusted for BMI) and model
299 3 (additionally adjusted for BMI). Therefore, it is considered that BMI might not be the
300 main underlying pathway in the positive association between bread and dairy pattern
301 and serum adiponectin. This finding was concordant with the result of path analysis
302 (Figure 1), which showed no significant association between bread and dairy pattern
303 and BMI in this study population.

304 We speculated that insignificant association between bread and dairy pattern and
305 adiponectin in men was possibly due to a small number of men in our study. Another
306 possible reason is that consumption of bread and dairy products was below the amount
307 required to have impact on adiponectin. Further, testosterone may suppress the
308 production of adiponectin in men (Swarbrick & Havel 2008).

309 Several studies have reported the positive significant association between dietary
310 patterns and serum adiponectin. Higher adherence to Mediterranean diet was
311 significantly associated with increasing concentrations of serum adiponectin in the U.S.
312 (Mantzoros et al. 2006). Another study in the U.S. (Fargnoli et al. 2008) indicated that
313 higher Alternate Healthy Eating Index (AHEI) score (which reflects a healthier dietary
314 pattern) was associated with higher HMW adiponectin concentrations. Meanwhile,
315 investigation in a healthy Mediterranean women population revealed the positive effects
316 of high non-refined cereals and low-fat dairy consumption on adiponectin (Yannakoulia
317 et al. 2008). Compared to those studies, there was no common food content between
318 those dietary patterns and our dietary pattern, except for low-fat dairy products. One
319 reason for the difference may be that average food and nutritional intake greatly differ
320 among the countries or population studied.

321 This study has several limitations. First, the relationship between bread and
322 dairy pattern and serum adiponectin should be interpreted with caution in term of time
323 sequence, because cross-sectional study design was used. Second, we had no data on the
324 validity of dietary pattern. However, other studies showed that validity and
325 reproducibility of the dietary pattern assessed from FFQ were acceptable (Hu et al.
326 1999; Nanri et al. 2012). Third, information on dietary intake and lifestyle factors were
327 collected using self-reported questionnaire. Thus, random measurement error might be

328 inevitable. Fourth, in the FFQ, we did not further classify the type of bread (refined or
329 whole grain) and dairy products (low-fat or high-fat). Nevertheless, our dietary records
330 showed that majority of our study participants consumed refined bread and normal dairy
331 products. Fifth, dietary patterns might depend on the statistical method used and
332 interpretation might be subjective, and it is uncertain if the results are applicable to
333 other populations. Nonetheless, our dietary patterns were similar to those reported for
334 another study in a Japanese population (Nanri et al. 2008). Lastly, the subjects in our
335 study were Japanese, thus, our result may not be generalizable to other ethnic
336 populations.

337 In conclusion, the result of the present study show that dietary pattern
338 characterized by high consumption of bread and dairy products, and low intake of rice
339 was positively but weakly related to higher concentration of serum adiponectin in
340 women. Further studies are needed to clarify the biological mechanisms for the
341 beneficial effects of dairy products toward serum adiponectin.

342

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351

352 **Disclosure statement**

353 The authors declare that there is no conflict of interest.

354

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360

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- 536 Table 1. Factor loading matrix for the extracted dietary patterns
- 537 Table 2. Associations between extracted dietary patterns and serum adiponectin in men
538 and women
- 539 Table 3. Baseline characteristics of the study participants based on the quartiles of the
540 bread and dairy pattern intake
- 541 Table 4. Pathways of the path model
- 542 Figure 1. Path diagram of the association between bread and dairy pattern, serum
543 adiponectin, and insulin resistance in women. * $p < 0.05$ ** $p < 0.01$ *** $p < 0.001$; solid
544 lines represent significant relationships; dashed lines represent non-significant
545 relationships; drinking habit: does not have a drinking habit = 0, has a drinking habit =
546 1; smoking habit: does not have a smoking habit = 0, has a smoking habit = 1;
547 menopausal status: woman with pre-menopausal status = 0, woman with post-
548 menopausal status = 1; BMI = Body Mass Index, HOMA-IR = Homeostasis Model
549 Assessment Insulin Resistance.

Table 1. Factor loading matrix for the extracted dietary patterns

Food items	Vegetable pattern	High-fat pattern	Seafood pattern	Bread and dairy pattern	Protein pattern
Rice	-	-0.20	0.26	-0.51	-
Bread	-	0.25	-0.25	0.55	-
Noodle	-	0.37	-	-	-
Bread & margarine	-	0.31	-0.17	0.25	-
Bread & butter	-	-	-	0.35	-
Milk	-	-	-	0.36	0.38
Yogurt	0.30	-	-	0.44	-
Miso	0.41	-	0.16	-	-
Chilled tofu & toppings, boiled tofu	0.41	-	0.27	-	0.28
Natto (fermented soybeans), soya beans (other cooked beans)	0.45	-0.20	0.23	0.16	0.33
Egg	0.30	0.21	-	-	0.36
Chicken	0.21	0.32	-	-0.31	0.24
Beef, pork	0.22	0.34	-0.15	-0.25	-
Liver	-	-	0.24	0.17	-
Ham, sausage, salami, bacon	0.20	0.44	-	-	0.16
Fish (such as sashimi, boiled fish, grilled fish)	0.40	-0.16	0.34	-	-
Small fish eaten with its bone	0.46	-0.21	0.23	0.20	-
Canned tuna	0.16	0.18	0.14	0.22	-
Squid, shrimp, crab, octopus	0.34	0.34	0.34	-	-0.30
Shellfish (such as clam, oyster)	0.33	0.27	0.46	0.15	-0.26
Cod roe, salmon roe	0.20	0.20	0.51	-	-
Fish paste cake, steamed seasoned fish paste	0.42	0.20	0.22	-	-
Deep-fried tofu with thinly sliced vegetables, deep-fried bean curd, deep fried tofu	0.50	-	0.18	-	-
Potato, taro, sweet potato	0.61	-	-	-	-
Pumpkin	0.56	-	-	-	0.19
Carrot	0.57	-	-0.19	-0.18	0.28
Broccoli	0.48	-	-	-	0.19
Green leafy vegetables (such as spinach, Japanese mustard spinach, edible chrysanthemum)	0.57	-0.25	-	-0.20	-
Green & yellow vegetables (such as green pepper, green beans)	0.57	-0.25	-	-0.17	-
Cabbage	0.56	-	-	-	-
Radish (boiled & grated radish)	0.60	-	-	-	-
Dried radish	0.46	-	0.29	0.16	-
Burdock, bamboo shoot	0.49	-	-	-	-
Other light-colored vegetables (such as cucumber, onion, bean sprouts, Chinese cabbage, lettuce)	0.61	-	-0.26	-0.26	-
Mushroom (shiitake, enoki, shimeji)	0.61	-	-0.25	-0.15	-
Seaweed (such as hijiki, kelp)	0.59	-0.19	-	-	-0.24
Mayonnaise (including potato salad)	0.17	0.58	-	-	-
Deep-fried food	0.20	0.60	-	-0.28	-0.15
Stir fry food (dish made with small amount of oil)	0.37	0.35	-0.23	-0.28	-
Mandarin orange, orange, grapefruit	0.48	-0.16	-0.19	0.20	-0.29
Other fruits (strawberry, kiwi, apple, watermelon)	0.49	-0.26	-0.28	0.23	-0.21
Peanuts, almond	0.36	-	-0.38	-	-0.23
Western confectionary (such as cake, cream puff)	0.29	0.26	-0.39	-	-0.41
Japanese confectionary (such as steamed bun)	0.46	0.20	-0.20	-	-0.30
Green tea	0.24	-	0.21	-	-0.32
Coffee	-	0.26	-0.27	0.17	-
Eigenvalue	7.24	2.59	2.21	1.99	1.65
Cumulative variance explained (%)	15.7	21.4	26.2	30.5	34.1

Factor loadings are equivalent to Pearson correlation between the food items and the dietary patterns.

For simplicity, factor loadings less than ± 0.15 were indicated by dash.

Table 2. Associations between extracted dietary patterns and serum adiponectin in men and women participants

Dietary pattern			Parameter estimate	Standard error	R ^{2†}	p
Vegetable pattern	Men	Model 1	-0.011	0.008	0.012	0.191
		Model 2	-0.009	0.009	0.108	0.308
		Model 3	-0.011	0.009	0.167	0.219
	Women	Model 1	0.003	0.005	0.016	0.526
		Model 2	0.004	0.005	0.032	0.484
		Model 3	0.0003	0.005	0.155	0.948
High-fat pattern	Men	Model 1	-0.007	0.013	0.003	0.589
		Model 2	-0.0001	0.013	0.102	0.991
		Model 3	-0.003	0.013	0.160	0.842
	Women	Model 1	-0.002	0.007	0.015	0.774
		Model 2	-0.004	0.008	0.032	0.626
		Model 3	0.004	0.007	0.156	0.611
Seafood pattern	Men	Model 1	-0.017	0.016	0.008	0.293
		Model 2	-0.014	0.016	0.106	0.382
		Model 3	-0.011	0.016	0.162	0.504
	Women	Model 1	-0.009	0.007	0.018	0.251
		Model 2	-0.011	0.008	0.036	0.155
		Model 3	-0.008	0.007	0.158	0.243
Bread and dairy pattern	Men	Model 1	0.007	0.015	0.003	0.627
		Model 2	0.008	0.015	0.104	0.575
		Model 3	0.006	0.014	0.161	0.667
	Women	Model 1	0.024	0.008	0.034	0.004
		Model 2	0.022	0.008	0.047	0.009
		Model 3	0.019	0.008	0.167	0.015
Protein pattern	Men	Model 1	0.002	0.019	0.002	0.922
		Model 2	0.002	0.018	0.102	0.932
		Model 3	-0.001	0.018	0.160	0.949
	Women	Model 1	0.003	0.009	0.015	0.734
		Model 2	0.005	0.009	0.032	0.541
		Model 3	-0.002	0.008	0.155	0.786

Model 1: Adjusted for age

Model 2: Adjusted for age, energy intake, physical activity, drinking habit, smoking habit, and menopausal status (women only)

Model 3: Adjusted for age, energy intake, physical activity, drinking habit, smoking habit, menopausal status (women only), and BMI

†R² of the model including dietary pattern and all covariates

Table 3. Baseline characteristics of the study participants based on the quartiles of the bread and dairy pattern intake

	Quartiles of bread and dairy pattern intake				<i>P</i>
	Q1	Q2	Q3	Q4	
Age (years) [†]	50 (41, 60)	50 (41, 61)	53 (44, 61)	59 (50, 64)	<0.0001
Sex [‡]					
Male	52 (34.0)	51 (33.3)	34 (22.2)	38 (24.8)	0.047
Female	101 (66.0)	102 (66.7)	119 (77.8)	115 (75.2)	
BMI (kg/m ²) [†]	23.0 (20.6, 26.0)	22.3 (20.1, 24.4)	22.5 (20.0, 24.4)	22.5 (20.5, 24.4)	0.123
Waist circumference (cm) [†]	81.0 (74.0, 90.0)	79.5 (73.0, 87.0)	80.0 (73.5, 85.0)	79.0 (73.5, 86.0)	0.259
Smoking [‡]					
Current	18 (11.8)	20 (13.1)	14 (9.2)	16 (10.5)	0.497
Past	36 (23.5)	27 (17.7)	25 (16.3)	25 (16.3)	
Never	99 (64.7)	106 (69.3)	114 (74.5)	112 (73.2)	
Drinking [‡]					
Current	65 (42.5)	77 (50.3)	70 (45.8)	70 (45.8)	0.863
Past	2 (1.3)	3 (2.0)	2 (1.3)	2 (1.3)	
Never	86 (56.2)	73 (47.7)	81 (52.9)	81 (52.9)	
Total energy intake (kcal/day) [†]	1659 (1492, 1885)	1542 (1375, 1720)	1533 (1335, 1690)	1562 (1396, 1733)	<0.0001
Physical activity (MET-hours/week) [†]	5.1 (0.4, 16.8)	3.8 (0.4, 15.3)	5.1 (0.4, 17.9)	9.2 (1.3, 28.5)	0.002
Triglycerides (mg/dl) [†]	87 (60, 128)	80 (56, 116)	77 (58, 103)	72 (56, 109)	0.159
HDL cholesterol (mg/dl) [†]	57 (48, 68)	59 (51, 71)	61 (55, 70)	63 (50, 71)	0.043
Fasting plasma glucose (mg/dl) [†]	89 (84, 93)	89 (85, 95)	89 (85, 94)	90 (85, 96)	0.529
Insulin (μU/ml) [†]	4.7 (3.4, 6.8)	4.5 (2.9, 6.3)	4.3 (3.2, 6.2)	4.3 (3.0, 5.7)	0.323
HOMA-IR [†]	1.04 (0.76, 1.54)	1.01 (0.63, 1.43)	0.95 (0.69, 1.41)	0.94 (0.64, 1.34)	0.446
Serum Adiponectin (μg/mL) [†]	3.68 (2.42, 6.12)	4.28 (2.45, 6.70)	4.88 (3.29, 6.76)	4.78 (3.24, 7.38)	0.004

[†]Median (25%; 75%), [‡]Number (%)

BMI = Body Mass Index; MET = metabolic equivalent; HOMA-IR = Homeostasis Model Assessment Insulin Resistance

Table 4. Pathways of the path model

	Standardized estimate	Unstandardized estimate	Standard error	<i>p</i>
BMI ← bread and dairy pattern score	-0.045	-0.002	0.002	0.357
BMI ← energy intake	0.054	0.048	0.043	0.267
BMI ← physical activity	-0.049	-0.003	0.003	0.322
BMI ← smoking	0.012	0.003	0.011	0.798
BMI ← menopausal status	0.085	0.011	0.006	0.085
Adiponectin ← bread and dairy pattern score	0.107	0.018	0.008	0.017
Adiponectin ← drink	0.074	0.037	0.022	0.093
Adiponectin ← age	0.105	0.003	0.002	0.177
Adiponectin ← smoking	-0.026	-0.024	0.041	0.563
Adiponectin ← BMI	-0.36	-1.405	0.171	<0.001
Adiponectin ← menopause	0.062	0.031	0.038	0.418
HOMA-IR ← adiponectin	-0.126	-0.131	0.045	0.004
HOMA-IR ← BMI	0.478	1.93	0.174	<0.001
HOMA-IR ← physical activity	-0.047	-0.013	0.011	0.251
HOMA-IR ← bread and dairy pattern score	-0.004	-0.001	0.007	0.926

BMI = Body Mass Index; HOMA-IR = Homeostasis Model Assessment Insulin Resistance

Figure 1.

