Supplemental Tables

Table S1.	Factor	loadings	of	food	groups	in	dietary	patterns	identified	using	principle
component	analysi	is									

	Balanced diet	Prudent diet	Noodle/ meat diet	Rice-based diet
Rice	-2	-7	3	93 *
Grain	8	-5	-2	-93 *
Noodles	3	1	62 *	3
Breads	-5	35	53 *	-4
Cakes	-4	29	31	6
Cookies	36	44 *	3	1
Beans	27	49 *	6	-2
Potatoes	51 *	-3	-1	-3
Kimchi	11	41 *	16	5
Eggs	-3	15	76 *	-3
Fast foods	69 *	39	-2	-2
Green vegetables	72 *	25	1	1
Mushroom	51 *	33	-5	-4
White vegetables	54 *	20	11	0
Fatty fish	66 *	14	12	0
White fish	49 *	2	18	1
Crabs	20	13	6	-2
Processed meats	46 *	-11	40 *	8
Red meats	16	2	66 *	-4
Soups	33	-8	38	4
Chickens	46 *	38	-2	-4
Seaweeds	14	48 *	2	1
Milk	22	29	6	2
Beverages	11	-19	16	13
Coffee	14	-11	23	14
Tea	22	46 *	-5	-5
Fruit	50 *	-4	4	1
Pickle	17	-30	15	5
Nuts	1	50 *	7	-4
Variance explained by each factor	3.7096422	2.3282962	2.2892598	1.796854

		Model 1		Model 2		
-	Low-PRS (n=5,276)	Medium-PRS (n=19,660)	High-PRS (n=12,178)	Medium-PRS (n=21,641)	High-PRS (n=4,201)	
BMI	1	1.004 (0.935-1.077)	1.002 (0.893-1.124)	0.972 (0.881-1.073)	1.003 (0.903-1.113)	
Waist	1	1.014	0.955	0.988	0.903	
circumference		(0.908-1.132)	(0.849-1.073)	(0.847-1.151)	(0.766-1.063)	
Type 2 diabetes	1	1.037	1.019	0.975	1.100	
Hypertension	1	(0.913~1.177) 1.080	(0.890~1.167) 1.103	(0.727~1.308) 1.011	(0.808~1.498) 1.035	
Total cholesterol	1	(0.993-1.174) 1.077	(1.010-1.206) 1.104	(0.902-1.133) 1.099	(0.917-1.168) 1.153	
LDL cholesterol	1	(0.973-1.192) 0.979	(0.991-1.231) 1.005	(0.954-1.267) 0.952	(0.990-1.343) 0.979	
HDL cholesterol	1	(0.899-1.067) 1.064	(0.917-1.100) 1.044	(0.849-1.067) 1.087	(0.867-1.105) 1.063	
TG	1	(0.992-1.140) 1.051 (0.968-1.142)	(0.970-1.124) 1.051 (0.968-1.142)	(0.985-1.198) 1.014 (0.913-1.127)	(0.958-1.180) 1.084 (0.970-1.212)	
Hs CRP	1	0.815	0.875	0.922	1.054	
WBC counts	1	(0.638-1.041) 0.963 (0.906- 1.023)	(0.675-1.135) 1.032 (0.967-1.102)	(0.674-1.261) 0.965 (0.907- 1.025)	(0.759-1.462) 1.030 (0.964-1.100)	

Table S2. Adjusted odds ratios for breast cancer according to the polygenetic risk scores of the best model (PRS) for gene-gene interaction after covariate adjustments

Values represent odd ratios and 95% confidence intervals.

The PRSBM scores of the subjects in the best model were calculated by the summation of the risk alleles of the included genetic variants in the model.

The subjects were divided into three groups by the tertiles of the PRS (0-3, 4-5, and >5).

Low-PRS was the reference for both model 1 and model 2.

The cutoff value of each parameter was as follows: $<25 \text{ kg/m}^2 \text{ BMI}$, 90 cm for men and 85 cm for women waist circumferences; $< 25 \text{ kg/m}^2 \text{ BMI}$, 90 cm for men and 85 cm for women waist circumferences, 230 mg/dL plasma total cholesterol concentrations, 40 mg/dL for men and 50 mg/dL for women plasma HDL cholesterol, 150 mg/dL plasma triglyceride concentrations, <126 ml/dL fasting serum glucose plus diabetic drug intake, 140 mmHg SBP, 90 mmHg DBP plus hypertension medication, 0.5 mg/dL serum high sensitive-C-reactive protein (hs-CRP) concentrations, and 5.6×10^9 /L white blood cell counts

Model 1: adjusted for age, residence area, survey year, body mass index (BMI), education, job, and income.

Model 2: adjusted for age, residence area, survey year, BMI, education, income, smoking, alcohol, energy, physical activity, fat percent intake carbohydrate percent intake, menopause age, initial menstruation age, and pregnancy experience.

*Significantly different from low GRS in logistic regression analysis at * P<0.05, ** P<0.01, *** P<0.001.