Supplementary Material: Identification of Dietary Pattern Networks Associated with Gastric Cancer Using Gaussian Graphical Models: A Case-Control Study

Madhawa Gunathilake, Jeonghee Lee, Il Ju Choi, Young-Il Kim and Jeongseon Kim

Table S1. Association between dietary pattern networks derived from GGMs and intestinal type of GC risk.

Distance wetternes	No. of	No. of	Model I OR	Model II OR	Model III OR	
Dietary patterns	controls	cases	(95% CI)	(95% CI)	(95% CI)	
Vegetables and						
seafood						
T1 (low)	276 (33.3)	69 (43.7)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	57 (36.1)	0.82 (0.56-1.21)	0.84 (0.53-1.33)	0.86(0.53-1.38)	
T3 (high)	277 (33.4)	32 (20.3)	0.46 (0.29-0.73)	0.47 (0.28-0.80)	0.52(0.30-0.91)	
<i>p</i> for trend			< 0.001	0.006	0.021	
Snacks and fats						
T1 (low)	277 (33.4)	70 (44.3)	1.00	1.00	1.00	
T2 (medium)	276 (33.3)	58 (36.7)	0.83 (0.56-1.22)	0.84 (0.54–1.32)	0.91 (0.57-1.45)	
T3 (high)	277 (3.4)	30 (19.0)	0.43 (0.27-0.68)	0.59 (0.33-1.04)	0.60 (0.33-1.08)	
<i>p</i> for trend			< 0.001	0.070	0.084	
Milk and dairy						
T1 (low)	276 (33.3)	78 (49.4)	1.00	1.00	1.00	
T2 (medium)	276 (33.3)	51 (32.3)	0.65 (0.44-0.97)	0.89 (0.57-1.42)	0.85 (0.53-1.38)	
T3 (high)	278 (33.5)	29 (18.4)	0.37 (0.23-0.58)	0.92 (0.50-1.66)	0.94 (0.51-1.76)	
<i>p</i> for trend			< 0.001	0.839	0.982	
Meat						
T1 (low)	276 (33.3)	70 (44.3)	1.00	1.00	1.00	
T2 (medium)	278 (33.5)	49 (31.0)	0.70 (0.47-1.04)	0.83 (0.51-1.35)	0.76 (0.46-1.26)	
T3 (high)	276 (33.3)	39 (24.7)	0.56 (0.36-0.85)	1.17 (0.63–2.15)	1.07 (0.57-2.02)	
<i>p</i> for trend			0.012	0.506	0.679	
Fruit						
T1 (low)	276 (33.3)	85 (53.8)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	48 (30.4)	0.56 (0.38–0.83)	0.79 (0.50–1.25)	0.72 (0.45–1.17)	
T3 (high)	277 (33.4)	25 (15.8)	0.29 (0.18-0.47)	0.50 (0.29–0.87)	0.53 (0.30-0.93)	
<i>p</i> for trend			< 0.001	0.014	0.027	

Model I: crude model; model II: adjusted for age, sex, family history of gastric cancer, smoking status, regular exercise, education, occupation, income and total energy intake; model III: additionally adjusted for *H. pylori* infection status

Diatary patterns	No. of	No. of	Model I OR	Model II OR	Model III OR	
Dietary patterns	controls	cases	(95% CI)	(95% CI)	(95% CI)	
Vegetables and						
seafood						
T1 (low)	276 (33.3)	61 (37.2)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	59 (36.0)	0.96 (0.65-1.43)	1.07 (0.69–1.66)	1.13 (0.72-1.80)	
T3 (high)	277 (33.4)	44 (26.8)	0.72 (0.47-1.09)	0.80 (0.50-1.29)	0.86 (0.53-1.42)	
<i>p</i> for trend	0.115 0.338		0.338	0.533		
Snacks and fats						
T1 (low)	276 (33.3)	64 (39.0)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	56 (34.2)	0.87 (0.58-1.30)	0.95 (0.61-1.47)	1.15 (0.72–1.82)	
T3 (high)	277 (33.4)	44 (26.8)	0.68 (0.45-1.04)	0.98 (0.60-1.64)	1.11 (0.65–1.88)	
<i>p</i> for trend			0.078	0.987	0.743	
Milk and dairy						
T1 (low)	277 (33.4)	72 (43.9)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	51 (31.1)	0.71 (0.48-1.05)	0.78 (0.50-1.21)	0.64 (0.40-1.02)	
T3 (high)	276 (33.3)	41 (25.0)	0.57 (0.38-0.87)	0.94 (0.56-1.57)	0.89 (0.52-1.53)	
<i>p</i> for trend			0.022	0.952	0.898	
Meat						
T1 (low)	276 (33.3)	65 (39.6)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	55 (33.5)	0.84 (0.57-1.25)	0.88 (0.55-1.40)	0.80 (0.49-1.29)	
T3 (high)	277 (33.4)	44 (26.8)	0.68 (0.44-1.02)	0.81 (0.46-1.43)	0.79 (0.44-1.42)	
<i>p</i> for trend			0.071	0.510	0.534	
Fruit						
T1 (low)	276 (33.3)	72 (43.9)	1.00	1.00	1.00	
T2 (medium)	277 (33.4)	55 (33.5)	0.76 (0.52-1.12)	0.80 (0.52-1.24)	0.86 (0.55-1.36)	
T3 (high)	277 (33.4)	37 (22.6)	0.51 (0.33-0.78)	0.55 (0.34-0.89)	0.55 (0.33-0.92)	
<i>p</i> for trend			0.003	0.016	0.019	

Table S2. Association	between	dietary	pattern	networks	derived	from	GGMs	and	diffuse	type of
GC risk.										

Model I: crude model; model II: adjusted for age, sex, family history of gastric cancer, smoking status, regular exercise, education, occupation, income and total energy intake; model III: additionally adjusted for *H. pylori* infection status

Model II OR

1.00 (ref)

0.77 (0.50-1.19)

	Males						
Dietary pattern	T1 (low)	T2 (medium)	T3 (high)	T1 (low)	T2 (medium)	T3 (high)	<i>p</i> -interaction
Vegetable and seafood pattern							
No. controls/cases	180/104	179/115	181/51	97/61	96/47	97/37	
Crude OR	1.00 (ref)	1.11 (0.79–1.56)	0.49 (0.33-0.72)	1.00 (ref)	0.78 (0.49–1.25)	0.61 (0.37–0.99)	0.669
Model I OR	1.00 (ref)	1.22 (0.82-1.80)	0.51 (0.32-0.81)	1.00 (ref)	0.85 (0.50-1.46)	0.76 (0.43-1.34)	0.964
Model II OR	1.00 (ref)	1.25 (0.82–1.91)	0.55 (0.34-0.89)	1.00 (ref)	1.04 (0.58–1.84)	0.82 (0.45–1.51)	0.964
Snacks and fat							
No. controls/cases	179/99	180/100	181/71	96/55	97/60	97/30	
Crude OR	1.00 (ref)	1.00 (0.71-1.42)	0.71 (0.49-1.03)	1.00 (ref)	1.08 (0.68–1.71)	0.54 (0.32-0.91)	0.471
Model I OR	1.00 (ref)	1.07 (0.72-1.60)	0.78 (0.50-1.20)	1.00 (ref)	1.09 (0.65–1.85)	0.62 (0.31-1.22)	0.277
Model II OR	1.00 (ref)	1.03 (0.67–1.58)	0.80 (0.50-1.28)	1.00 (ref)	1.29 (0.73–2.27)	0.65 (0.32–1.34)	0.180
Meat							
No. controls/cases	180/119	180/82	180/69	97/57	96/38	97/50	
Crude OR	1.00 (ref)	0.69 (0.48-0.98)	0.58 (0.40-0.83)	1.00 (ref)	0.67 (0.41–1.11)	0.88 (0.55–1.41)	0.174
Model I OR	1.00 (ref)	0.84 (0.56-1.27)	1.17 (0.72–1.90)	1.00 (ref)	0.72 (0.40-1.30)	0.85 (0.46–1.56)	0.165
Model II OR	1.00 (ref)	0.93 (0.60–1.44)	1.23 (0.74–2.06)	1.00 (ref)	0.67 (0.36–1.27)	0.65 (0.34–1.23)	0.426
Fruits							
No. controls/cases	180/124	180/80	180/66	97/82	96/35	97/28	
Crude OR	1.00 (ref)	0.65 (0.46-0.91)	0.53 (0.37–0.77)	1.00 (ref)	0.43 (0.27–0.70)	0.34 (0.20-0.57)	0.120
Model I OR	1.00 (ref)	0.81 (0.54-1.21)	0.77 (0.50-1.17)	1.00 (ref)	0.54 (0.32-0.93)	0.56 (0.32-1.00)	0.245

Table S3. Interaction between GGM derived dietary patterns and sex in the risk of GC.

Model I: crude model; model II: adjusted for age, family history of gastric cancer, smoking status, regular exercise, education, occupation, income and total energy intake; model III: additionally adjusted for *H. pylori* infection status

1.00 (ref)

0.59 (0.33-1.05)

0.62 (0.34-1.14)

0.403

0.76 (0.48-1.19)



Figure S1. Dietary intake networks for intestinal type GC derived by Gaussian graphical models.



Figure S2. Dietary intake networks for diffuse type GC derived by Gaussian graphical models.