**S4 Table.** Association between genetic risk score and the risk of colorectal cancer, stratified by age<sup>a)</sup>

	< 50 Years	≥ 50 Years				
Genetic risk score	No. of controls/cases	Crude OR (95% CI)	Multivariate OR (95% CI) <sup>b)</sup>	No. controls/cases	Crude OR (95% CI)	Multivariate OR (95% CI) <sup>b)</sup>
Unweighted		,	,		,	,
≤ 11	57/16	1.0 (reference)	1.0 (reference)	216/64	1.0 (reference)	1.0 (reference)
12-16	164/70	1.52 (0.82-2.83)	1.60 (0.83-3.09)	578/271	1.58 (1.16-2.17)	1.74 (1.23-2.46)
≥ 17	70/55	2.80 (1.45-5.40)	3.21 (1.59-6.47)	210/156	2.51 (1.77-3.55)	2.82 (1.91-4.16)
Weighted						
<b>T</b> 1	70/18	1.0 (reference)	1.0 (reference)	253/80	1.0 (reference)	1.0 (reference)
T2	133/61	1.78 (0.98-3.25)	1.94 (1.02-3.67)	514/231	1.42 (1.06-1.91)	1.43 (1.03-1.98)
T3	88/62	2.74 (1.49-5.05)	3.04 (1.58-5.82)	237/180	2.40 (1.75-3.30)	2.47 (1.73-1.98)

OR, adds ratio; CI, confidence interval; T, tertile; SNP, single nucleotide polymorphism. <sup>a)</sup>Genetic risk score was calculated using 13 SNPs, <sup>b)</sup>Adjusted for sex, family history of colorectal cancer, education, obesity, physical activity, smoking, alcohol consumption, and dietary inflammatory index.