

S8 Table. Five significant miRNA-related SNPs from Ye et al. (40) and risk of EA (A) or BE (B).

A.

	SNP	Gene	Alleles [†]	Controls		EA cases		OR*	95% CI	P
				N	MAF [‡]	N	MAF [‡]			
1	rs213210	pri-miR-219-1	G/A	3204	0.069	2491	0.080	1.12	(0.96-1.30)	0.147
2	rs6505162	miR-423, -3184	C/A	3207	0.455	2495	0.455	1.04	(0.96-1.12)	0.348
3	rs14035	RAN	A/G	3206	0.310	2494	0.316	1.04	(0.95-1.13)	0.398
4	rs11614913	miR-196a-2	T/C	3204	0.402	2495	0.416	1.02	(0.94-1.10)	0.629
5	rs11077	XPO5	C/A	3206	0.415	2495	0.423	1.01	(0.93-1.10)	0.749

B.

	SNP	Gene	Alleles [†]	Controls		BE cases		OR*	95% CI	P
				N	MAF [‡]	N	MAF [‡]			
1	rs213210	pri-miR-219-1	G/A	3201	0.069	3284	0.074	1.03	(0.90-1.19)	0.623
2	rs6505162	miR-423, -3184	C/A	3204	0.455	3285	0.449	1.01	(0.94-1.08)	0.831
3	rs14035	RAN	A/G	3203	0.310	3288	0.304	0.99	(0.92-1.07)	0.785
4	rs11614913	miR-196a-2	T/C	3201	0.402	3285	0.410	1.00	(0.93-1.07)	0.993
5	rs11077	XPO5	C/A	3203	0.415	3287	0.427	1.04	(0.97-1.12)	0.305

[†]Minor/major alleles, [‡]Minor allele frequency, *OR adjusted for age, sex, ev1-ev4, using additive model (per-allele)