Supplementary Information for Association study between *XPG* Asp1104His polymorphism and colorectal cancer risk in a Chinese population

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Table S1. The baseline characteristics of all qualified studies in this meta-analysis.

Study	year	country	Ethnicity	Design	Case/control	Genotype distribution										HWE for P	MAF
						cases					controls						
						GG	GC	CC	G	C	GG	GC	CC	G	C		
Pardini	2008	Czech	Caucasian	HB	532/532	334	177	21	845	219	356	153	23	865	199	0.21	0.187[a]/0.267[b]
Emel	2011	Turkish	Caucasian	PB	79/247	43	34	2	120	38	148	83	16	379	115	0.35	0.232[a]/0.267[b]
Gil	2012	Polish	Caucasian	HB	132/100	86	35	11	207	57	64	31	5	159	41	0.62	0.205[a]/0.267[b]
Liu	2012	China	Asian	HB	1028/1085	233	603	192	1069	987	329	537	219	1195	975	0.99	0.449[a]/0.444[b]
zhu	2013	China	Asian	HB	878/884	286	459	133	1031	725	355	405	124	1115	653	0.62	0.369[a]/0.444[b]

[a] the MAF in our study;
[b] the MAF in the HapMap database
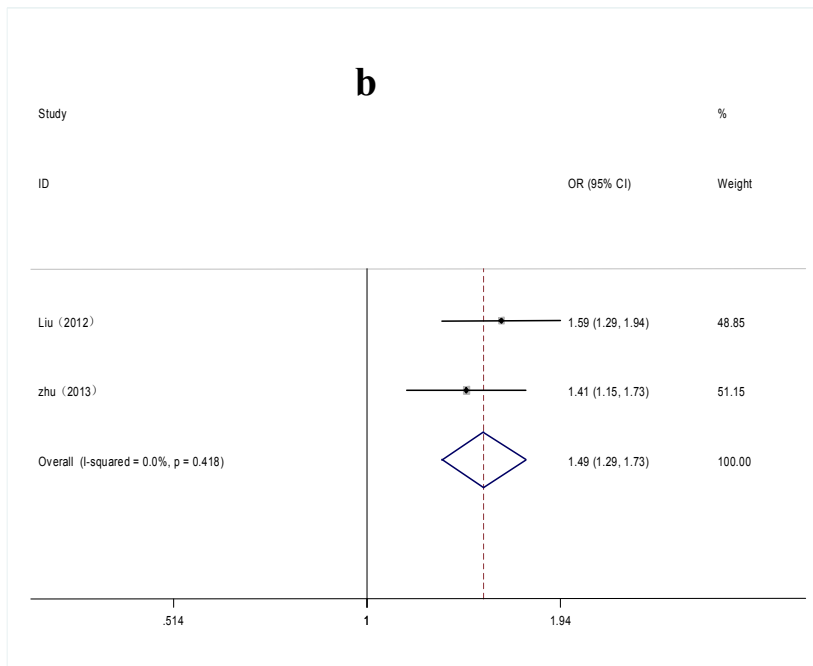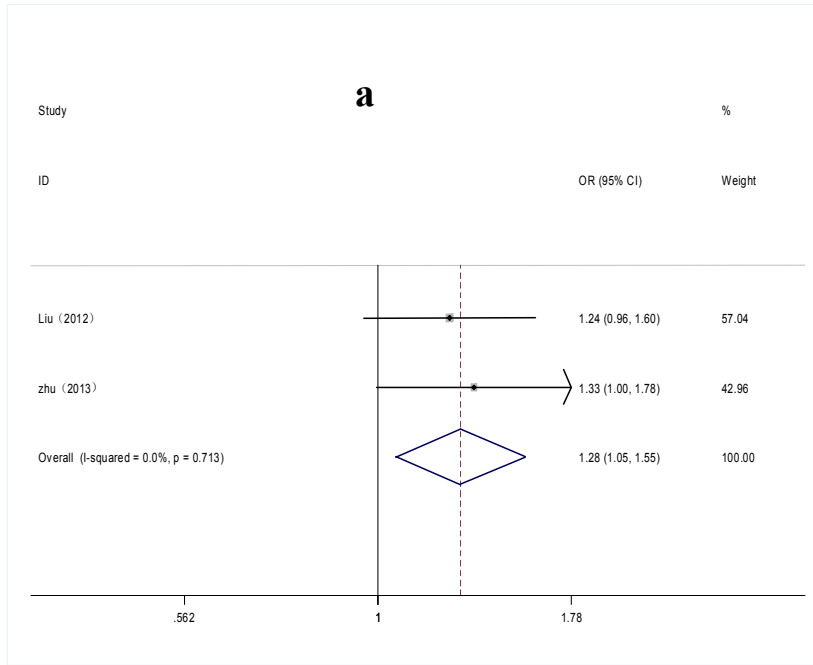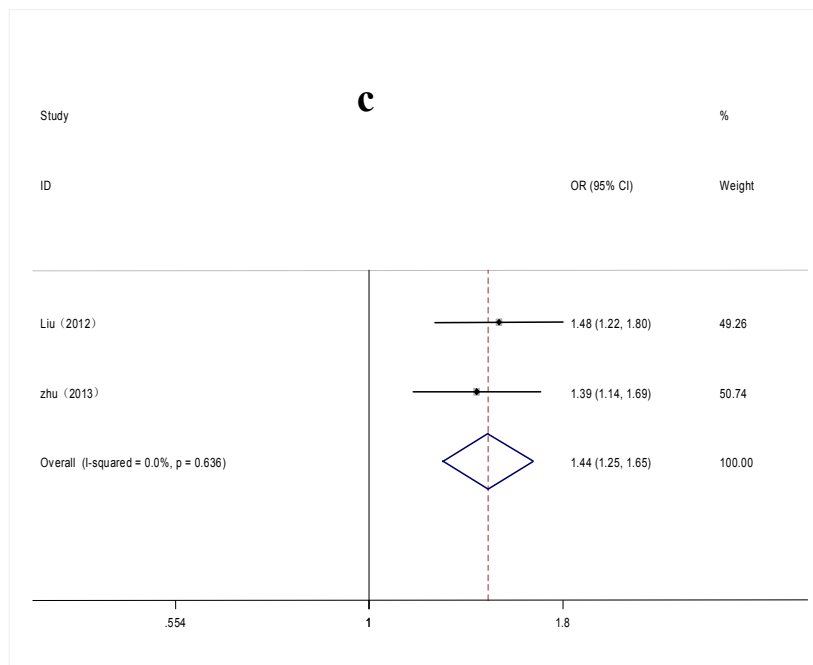
Figure S1. Odds ratios (ORs) for associations between *XPG* Asp1104His polymorphism and colorectal cancer in Asian populations. (based on all models: (a) His/His vs. Asp/Asp, (b) Asp/His vs. Asp/Asp, (c) His/His + Asp/His vs. Asp/Asp