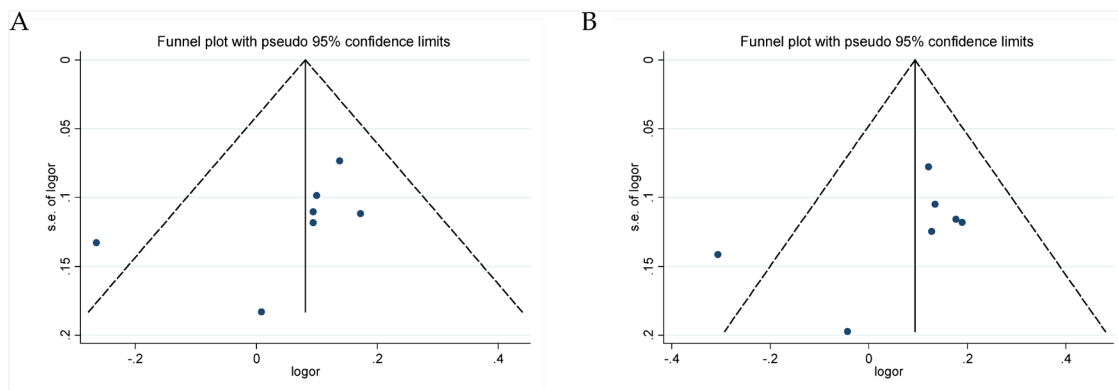
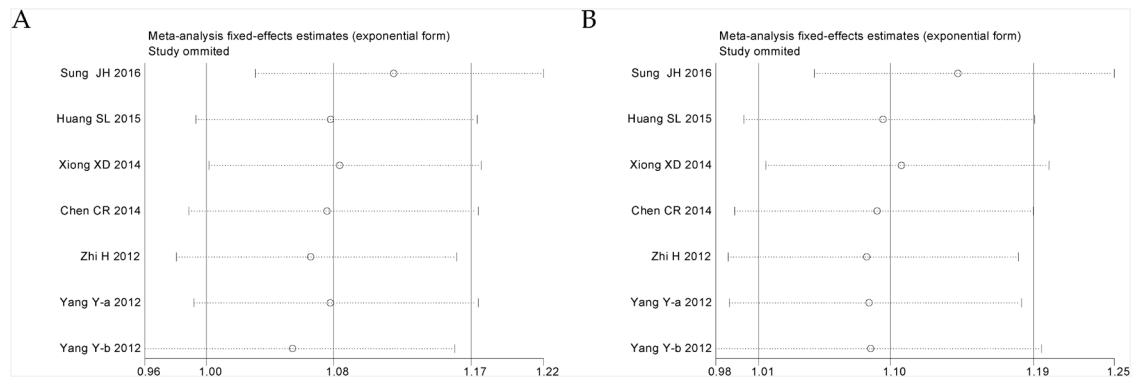


# Significant association between functional microRNA polymorphisms and coronary heart disease susceptibility: a comprehensive meta-analysis involving 16484 subjects

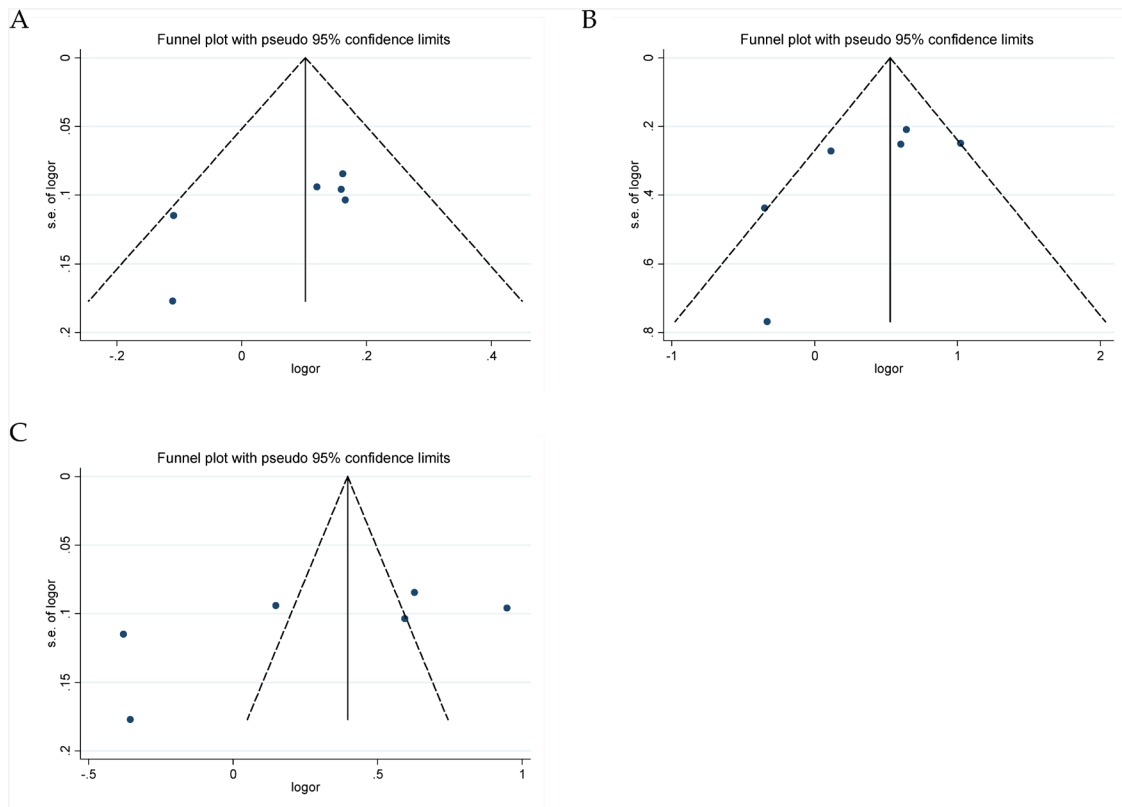
## SUPPLEMENTARY FIGURES



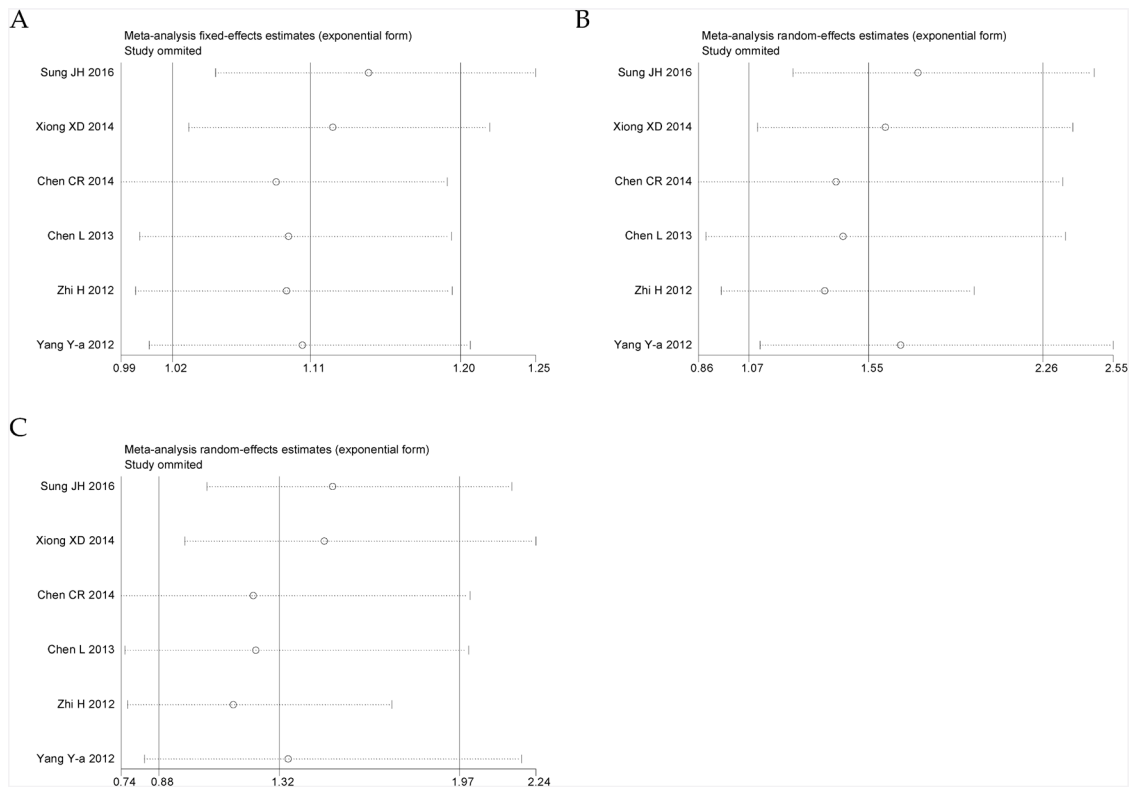
**Supplementary Figure 1: Funnel plots for microRNA-196a2 rs11614913 T>C polymorphism and CHD. A. dominant model (TC + CC vs. TT); B. heterozygote model (TC vs. TT).**



**Supplementary Figure 2: Sensitivity analyses for microRNA-196a2 rs11614913 T>C polymorphism and CHD. A. dominant model (TC + CC vs. TT); B. heterozygote model (TC vs. TT).**



**Supplementary Figure 3: Funnel plots for microRNA-499 rs3746444 A>G polymorphism and CHD. A.** allele model (G vs. A); **B.** recessive model (GG vs. AA + AG); **C.** homozygote model (GG vs. AA).



**Supplementary Figure 4: Sensitivity analyses for microRNA-499 rs3746444 A>G polymorphism and CHD. A.** allele model (G vs. A); **B.** recessive model (GG vs. AA + AG); **C.** homozygote model (GG vs. AA).