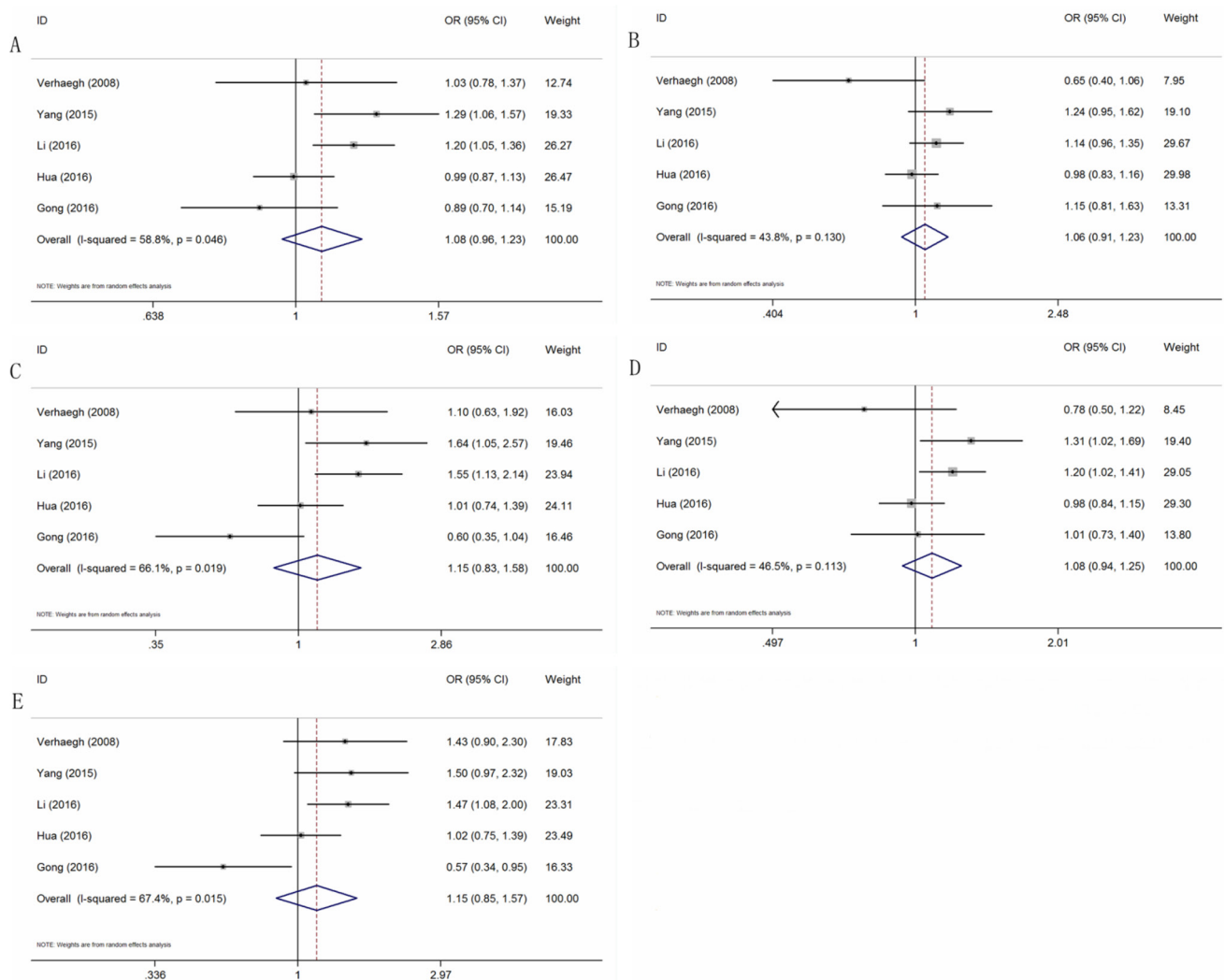
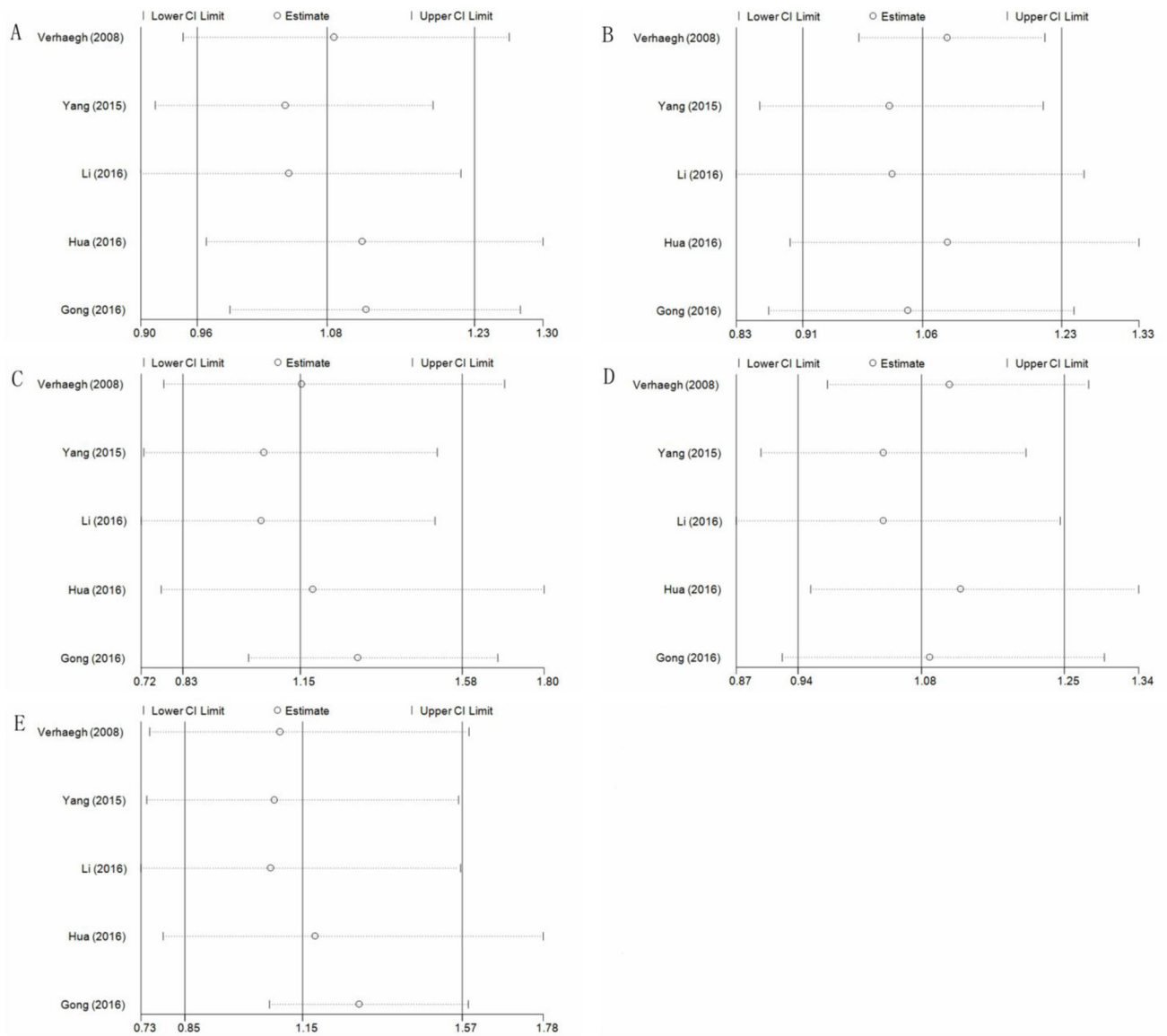


# Significant association between lncRNA H19 polymorphisms and cancer susceptibility: a meta-analysis

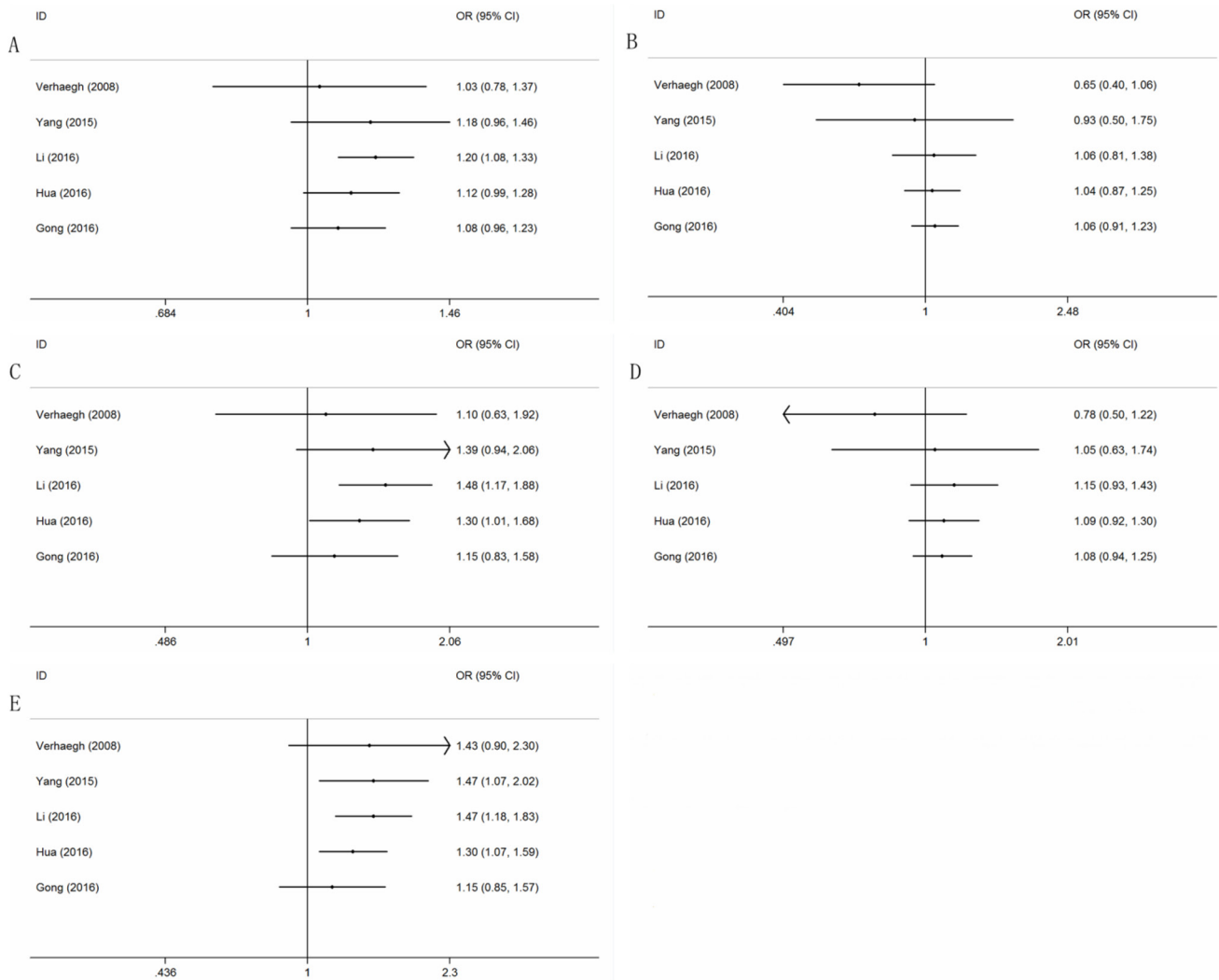
## Supplementary Materials



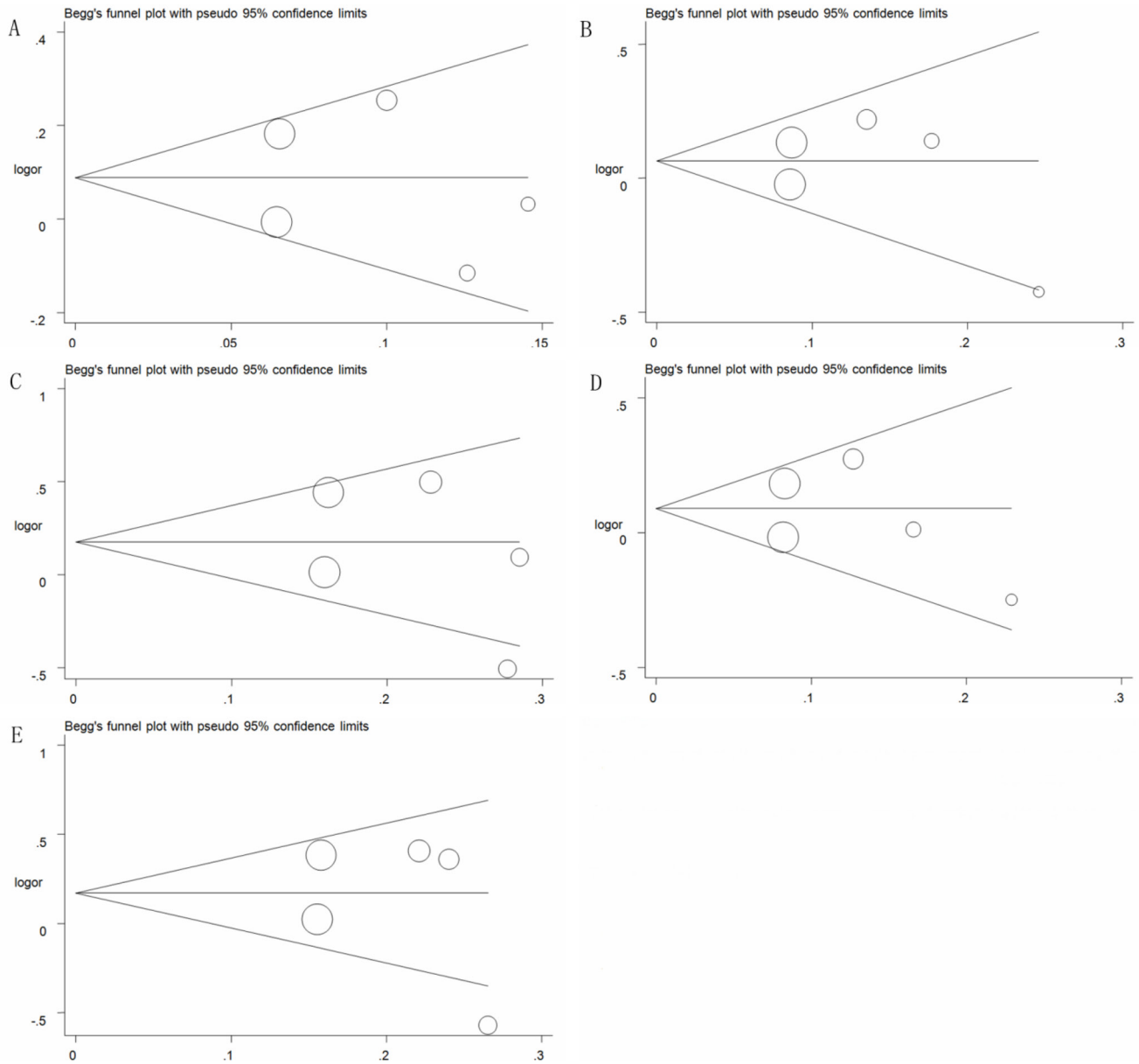
**Supplementary Figure 1: OR and 95% CIs of the associations between rs2839698 G>A polymorphism and cancer risk. (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model).**



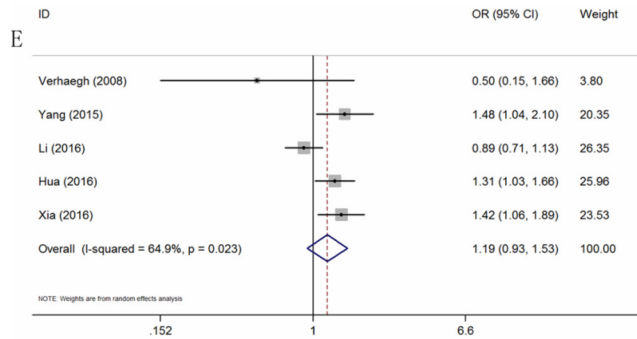
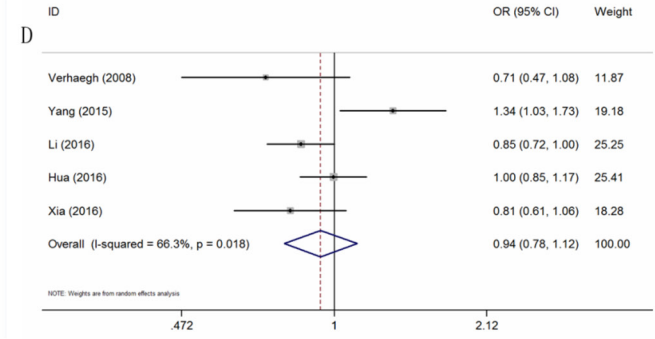
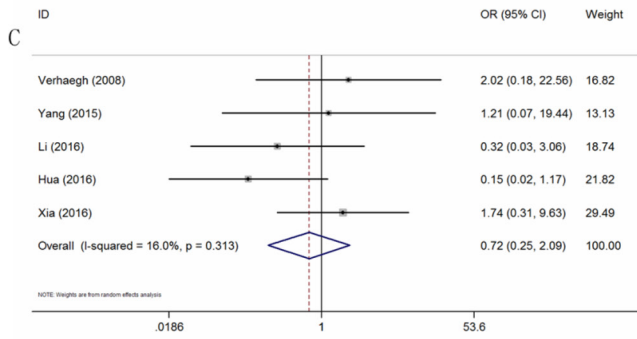
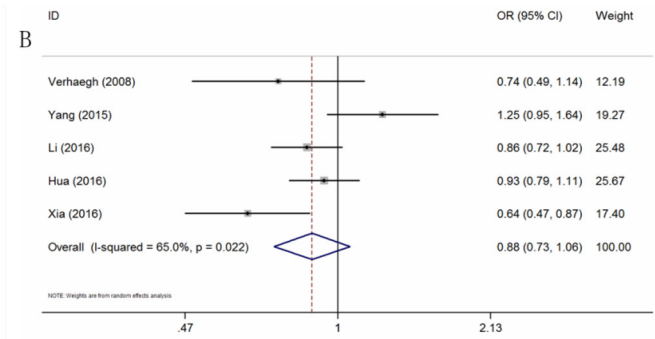
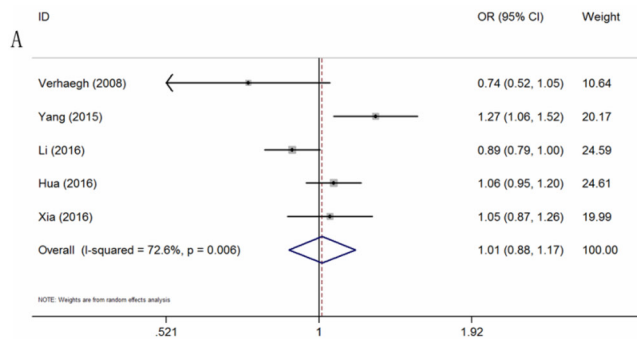
**Supplementary Figure 2: Sensitivity analyses according to publication year in rs2839698 G>A polymorphism. (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model).**



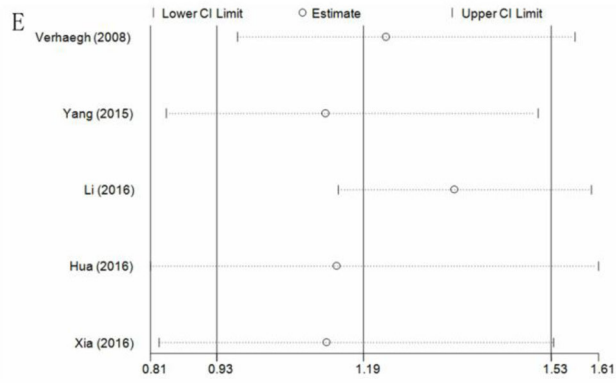
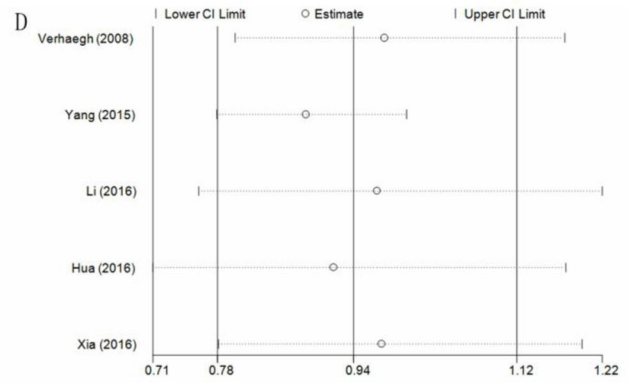
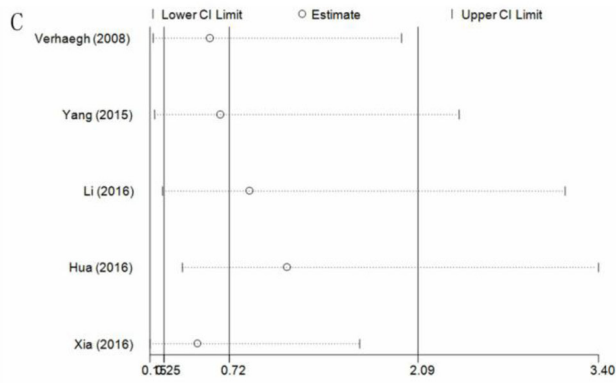
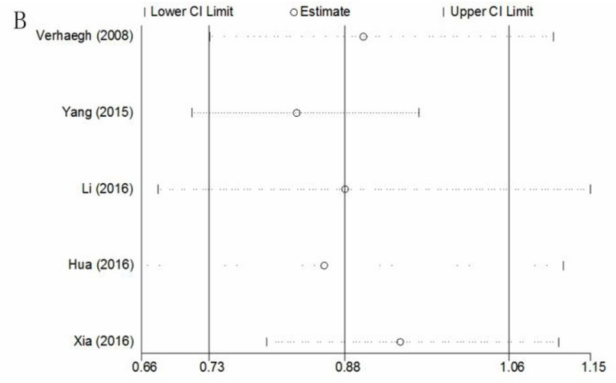
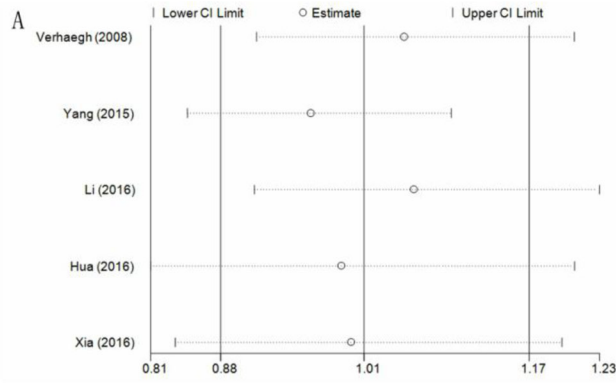
**Supplementary Figure 3: Cumulative meta-analyses according to publication year in rs2839698 G>A polymorphism. (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model).**



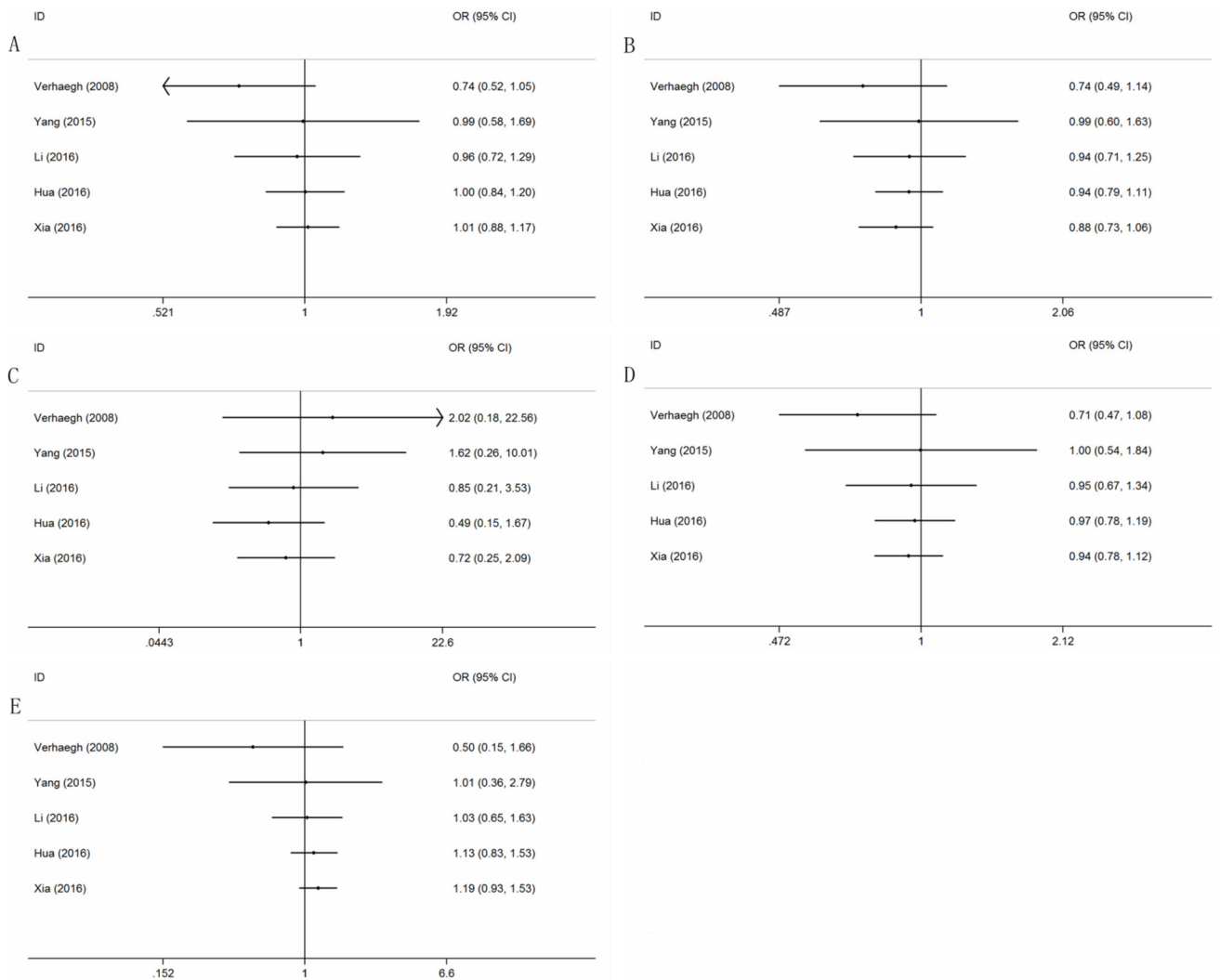
**Supplementary Figure 4: Funnel plot analysis to detect publication bias in rs2839698 G>A polymorphism.** (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model). Circles represent the weight of the studies.



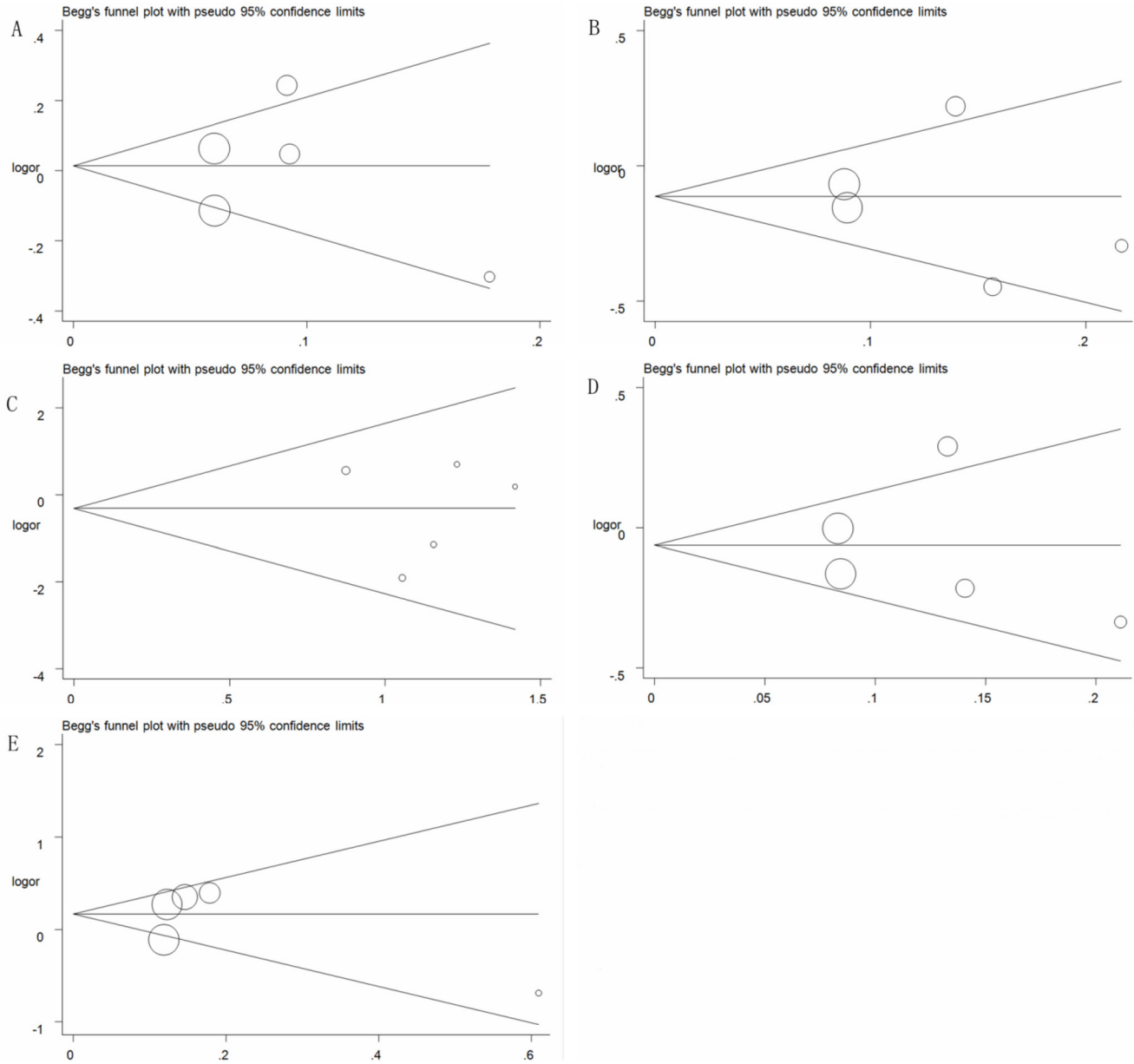
**Supplementary Figure 5: OR and 95% CIs of the associations between rs217727 G>A polymorphism and cancer risk. (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model).**



**Supplementary Figure 6: Sensitivity analyses according to publication year in rs217727 G>A polymorphism. (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model).**

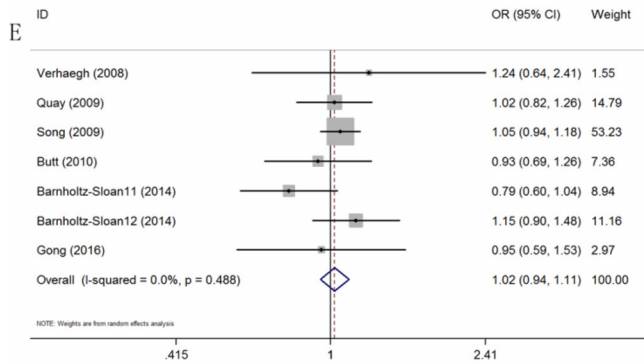
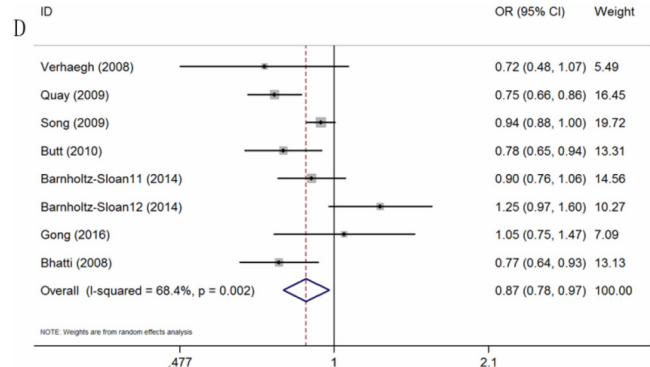
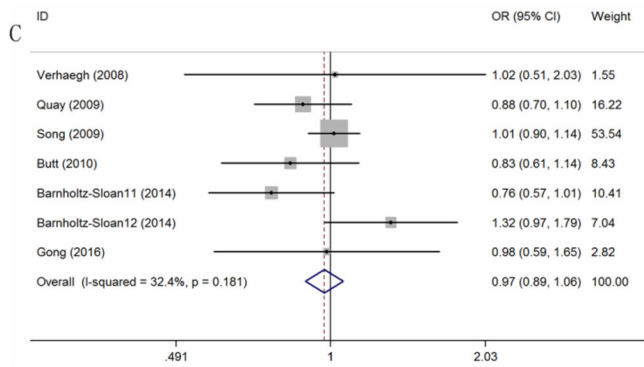
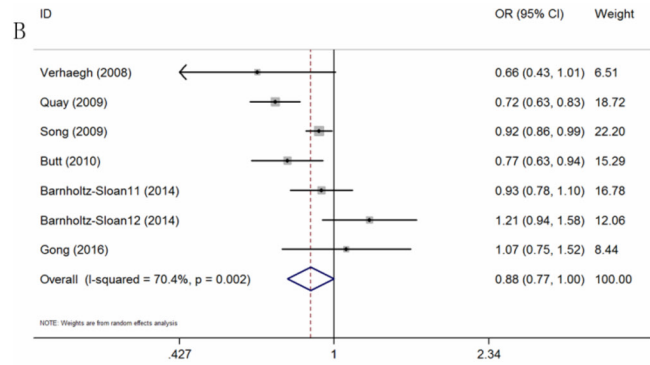
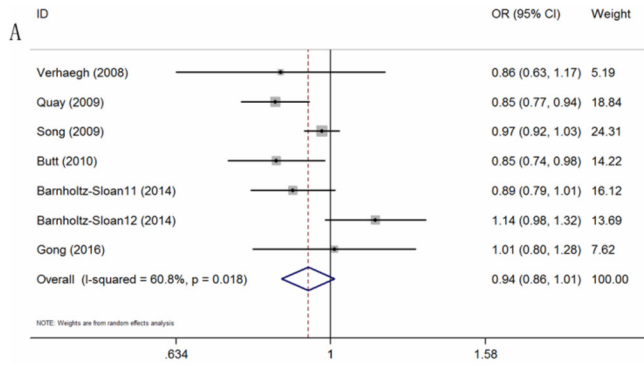


**Supplementary Figure 7: Cumulative meta-analyses according to publication year in rs217727 G>A polymorphism. (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model).**

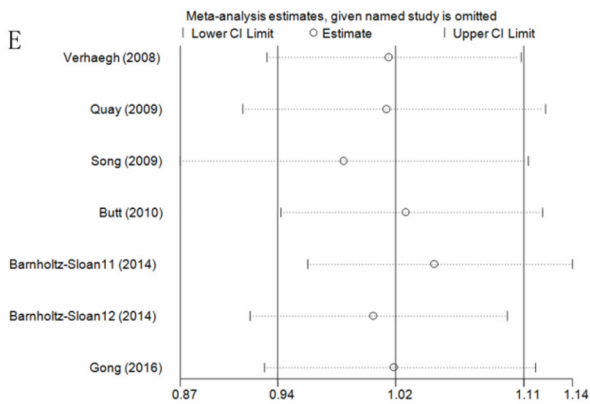
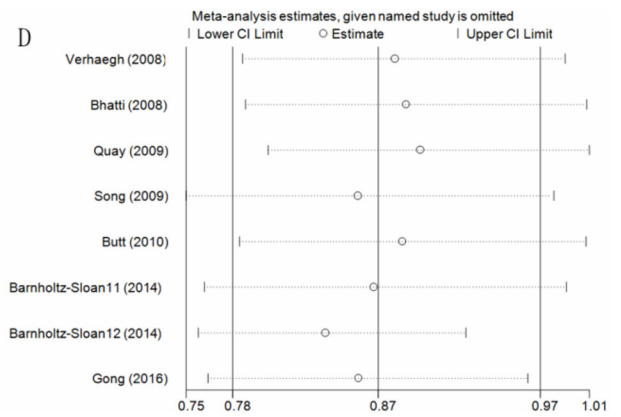
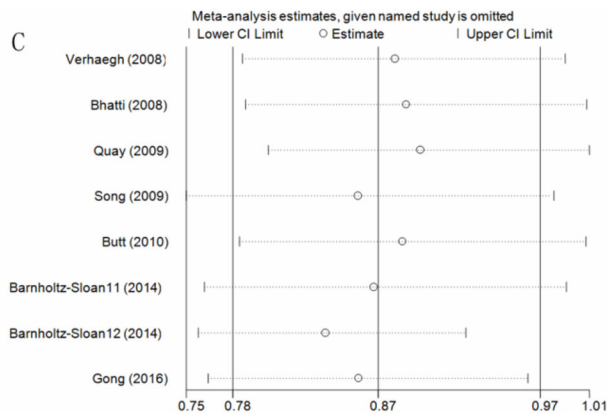
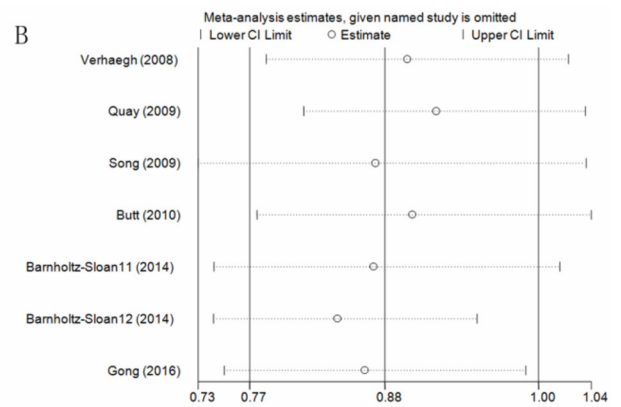
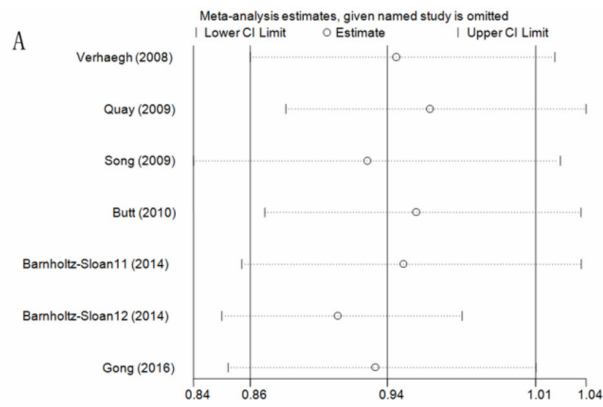


**Supplementary Figure 8: Funnel plot analysis to detect publication bias in rs217727 G>A polymorphism.** (A) for A vs. G model; (B) for GA vs. GG model; (C) for AA vs. GG model; (D) for GA+AA vs. GG model; (E) for AA vs. GG+GA model). Circles represent the weight of the studies.

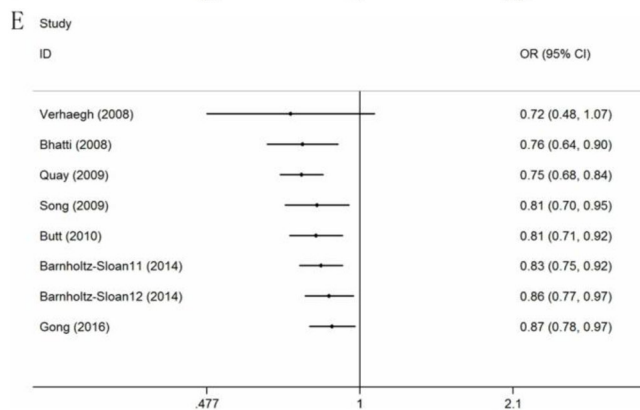
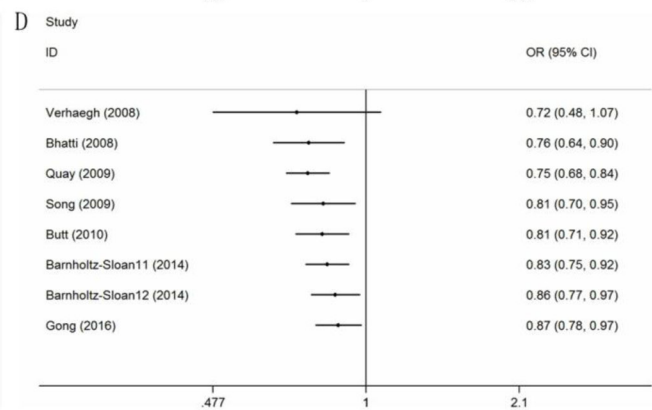
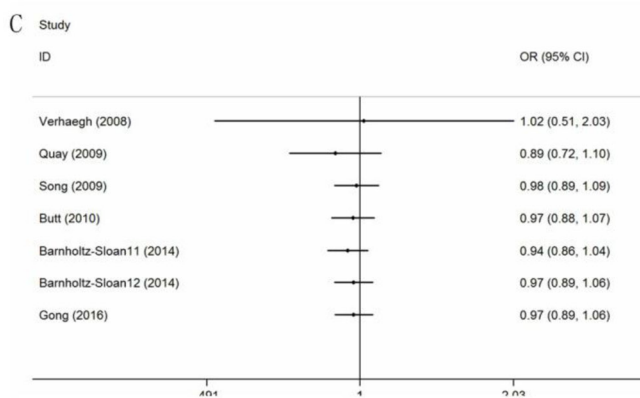
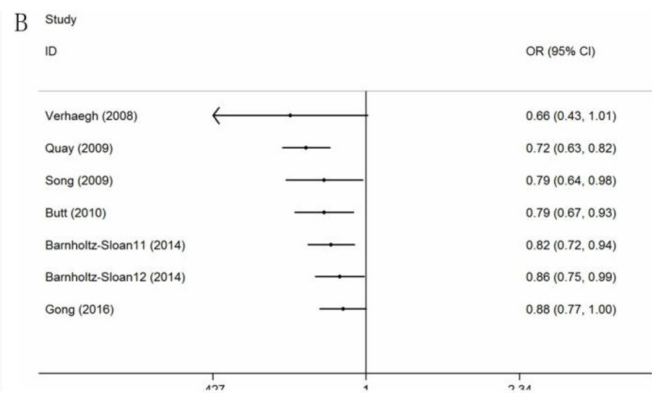
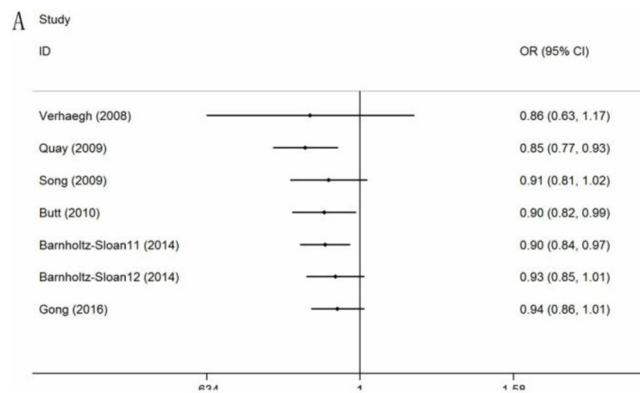




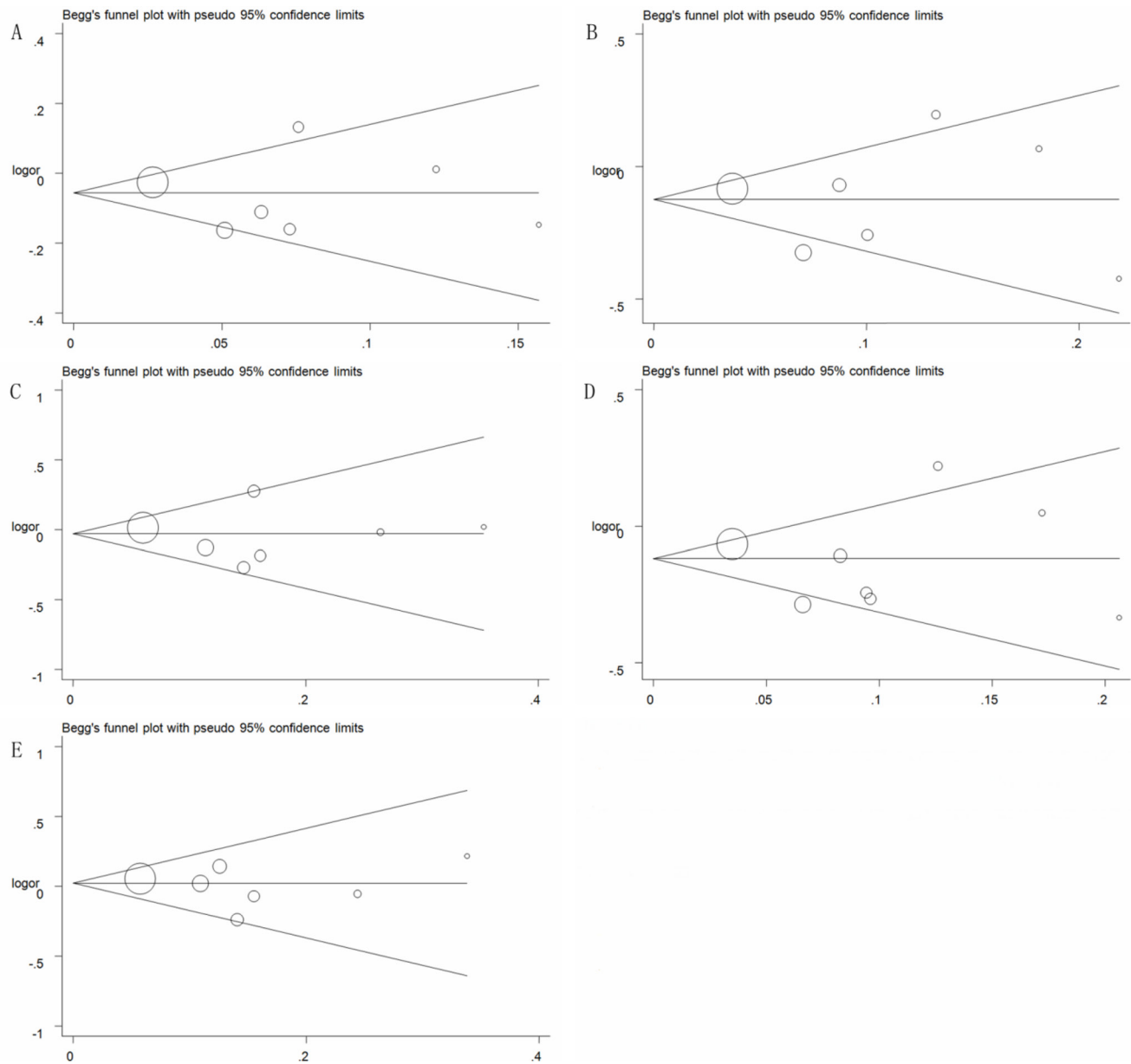
**Supplementary Figure 9: OR and 95% CIs of the associations between rs2107425 C>T polymorphism and cancer risk. (A) for T vs. C model; (B) for CT vs. CC model; (C) for TT vs. CC model; (D) for CT+TT vs. CC model; (E) for TT vs. CC+CT model.**



**Supplementary Figure 10: Sensitivity analyses according to publication year in rs2107425 C>T polymorphism. (A) for T vs. C model; (B) for CT vs. CC model; (C) for TT vs. CC model; (D) for CT+TT vs. CC model; (E) for TT vs. CC+CT model).**



**Supplementary Figure 11: Cumulative meta-analyses according to publication year in rs2107425 C>T polymorphism. (A) for T vs. C model; (B) for CT vs. CC model; (C) for TT vs. CC model; (D) for CT+TT vs. CC model; (E) for TT vs. CC+CT model.**



**Supplementary Figure 12: Funnel plot analysis to detect publication bias in rs2107425 C>T polymorphism.** (A) for T vs. C model; (B) for CT vs. CC model; (C) for TT vs. CC model; (D) for CT+TT vs. CC model; (E) for TT vs. CC+CT model). Circles represent the weight of the studies.