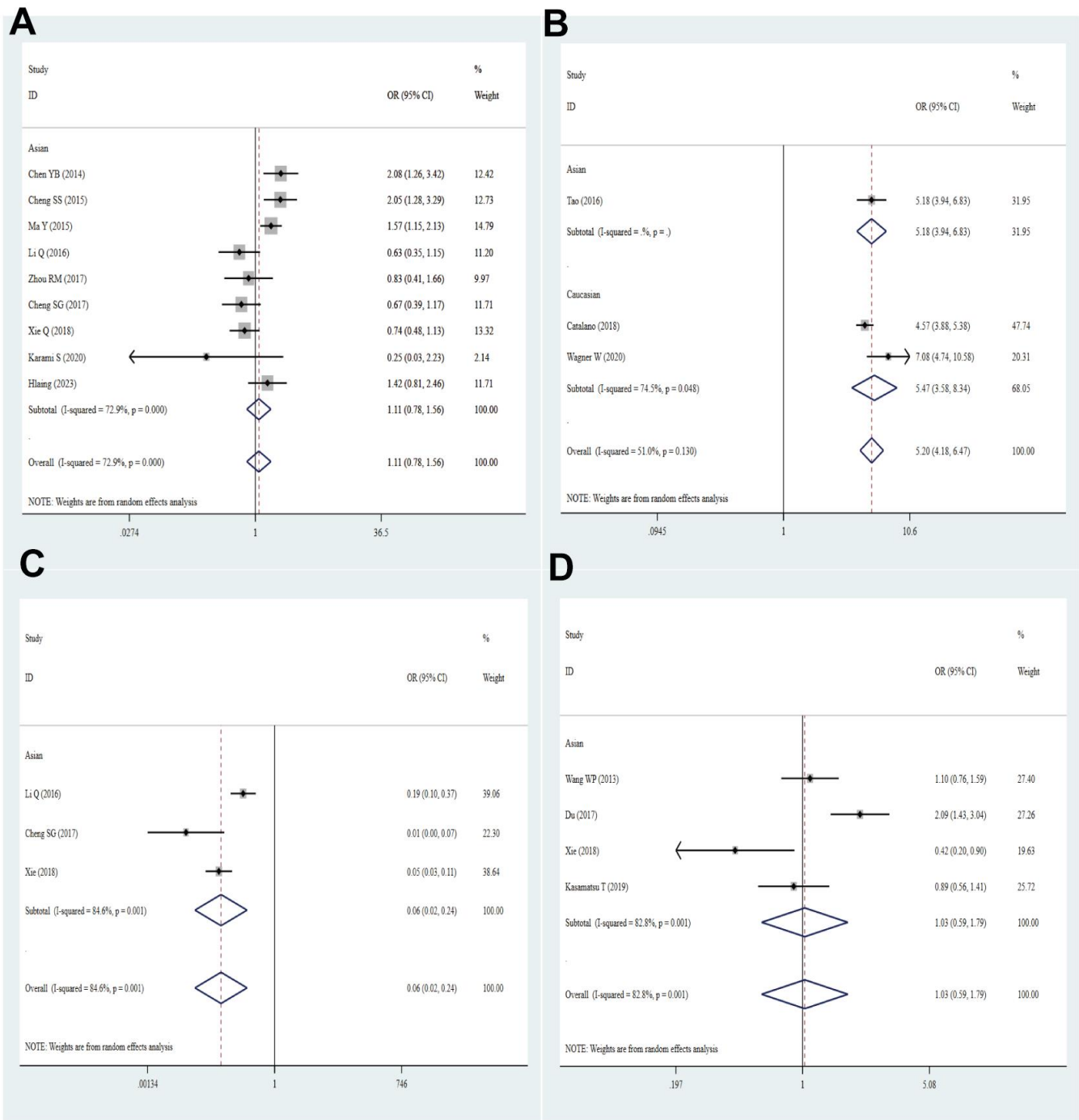
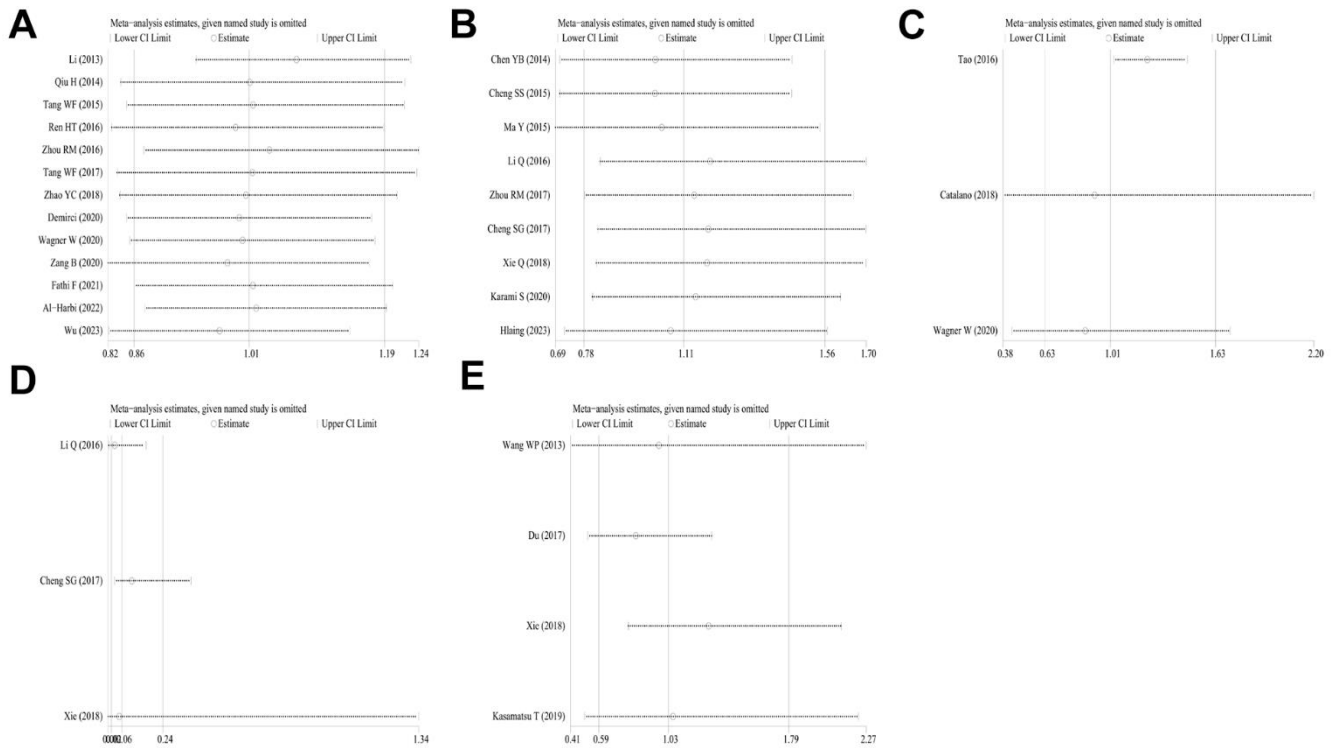


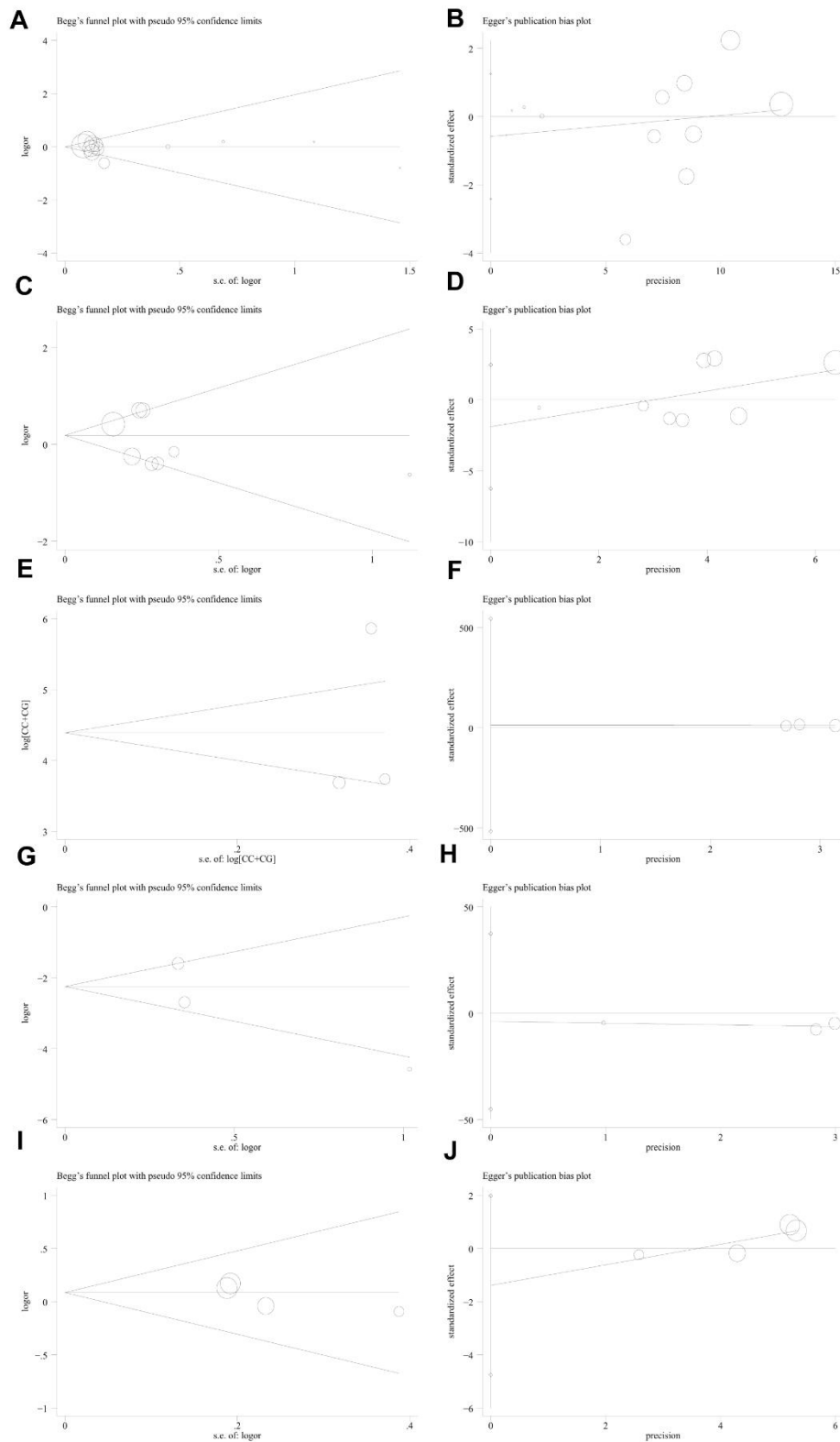
SUPPLEMENTARY FIGURES



Supplementary Figure 1. Forest plot for the association between PD-L1 gene polymorphisms and cancer susceptibility in the homozygote model. For each publication, the estimation of OR and its 95% CI was plotted with a box and a horizontal line. The diamonds represented the pooled ORs and 95% CIs. (A) rs2890658 polymorphism; (B) rs10815225 polymorphism; (C) rs17718883 polymorphism; (D) rs2297136 polymorphism.



Supplementary Figure 2. Analyses of the influence of a single study on the total combined OR in the dominant model. (A) PD-1.6 polymorphism, (B) rs2890658 polymorphism; (C) rs10815225 polymorphism; (D) rs17718883 polymorphism; (E) rs2297136 polymorphism.



Supplementary Figure 3. Begg's funnel plot and Egger's linear regression test for the publication biases under the dominant model. (A) Begg's test for PD-1.6 polymorphism; (B) Egger's test for PD-1.6 polymorphism; (C) Begg's test for PD-L1 rs2890658 polymorphism; (D) Egger's test for PD-L1 rs2890658 polymorphism; (E) Begg's test for PD-L1 rs10815225 polymorphism; (F) Egger's test for PD-L1 rs10815225 polymorphism; (G) Begg's test for PD-L1 rs17718883 polymorphism; (H) Egger's test for PD-L1 rs17718883 polymorphism; (I) Begg's test for PD-L1 rs2297136 polymorphism; (J) Egger's test for PD-L1 rs2297136 polymorphism.