**Additional file 2** Figure S1. Funnel plot to diagnosis of the publication bias. Figure S2. Combined estimates for the association between APC methylation and NSCLC after trim-fill treatment. Figure S3. Sensitivity analyses of the overall effect by omitting a single study.

Figure S1. Funnel plot to diagnosis of the publication biasF:\Project\methylation\APCmeta\manuscript\clinical_epigenetics\S Figure S1. Funnel plot to diagnosis of the publication bias.tif

Figure S2. Combined estimates for the association between APC methylation and NSCLC after trim-fill treatment.

F:\Project\methylation\APCmeta\manuscript\clinical_epigenetics\S Figure S2. Combined estimates for OR in Random and Fixed Effect model after trimfill treatment.tiff

Figure S3. Sensitivity analyses of the overall effect by omitting a single study.

F:\Project\methylation\APCmeta\manuscript\clinical_epigenetics\S Figure S3. Sensitivity analyses of the overall effect by omitting a single study.tiff