**Supplemental Figure S1 A-C. The eQTL analysis of *CNOT6* mRNA expressions for rs2453176 in the 105 adjacent normal lung cancer tissue samples from the TCGA database.** A. additive model, *P* = 0.491, B. dominant model, *P* = 0.990, C. recessive model, *P* = 0.667; **D, The mRNA expression of *CNOT6* in the 107 paired lung cancer and normal adjacent tissue samples from the TCGA database** (*P* = 0.237).

**Supplementary Figure S2. The ENCODE project data of rs2453176 from UCSC browser (NCBI137/hg19).**