Additional File 7.

Figure S1. Linkage disequilibrium (LD) plots of genotyped SNPs in CEBPG and PEPD for cases (A) and controls (B) separately. Case (A) and control (B) genotype frequency data for each SNP genotyped in the region of CEBPG and PEPD were used to generate the plots in Haploview software. LD plots were constructed using r^2 as the LD statistic, with black indicating complete LD and white indicating no LD. SNPs marked with an asterisk (*) were nominally associated with AAA. For details on SNPs, see Additional file 1, Table S1.