

Supplementary Data

SUPPLEMENTARY TABLE S1. ALLELE FREQUENCIES

<i>SNP</i>	<i>Chromosome</i>	<i>Functional consequence</i>	<i>MAF Hapmap-CHB</i>
rs12050772	15:51252674	Intron variant	0.512
rs2289105	15:51215311	Intron variant	0.573
rs4774585	15:51324283	5'flanking region of exon2a	0.037

We selected three SNPs of the *CYP19* gene that had a minor allele frequency >0.03 , in the Chinese, using the National Center for Biotechnology Information (NCBI) SNP database (www.ncbi.nlm.nih.gov/projects/SNP) and taking advantage of prior resequencing data and functional studies.