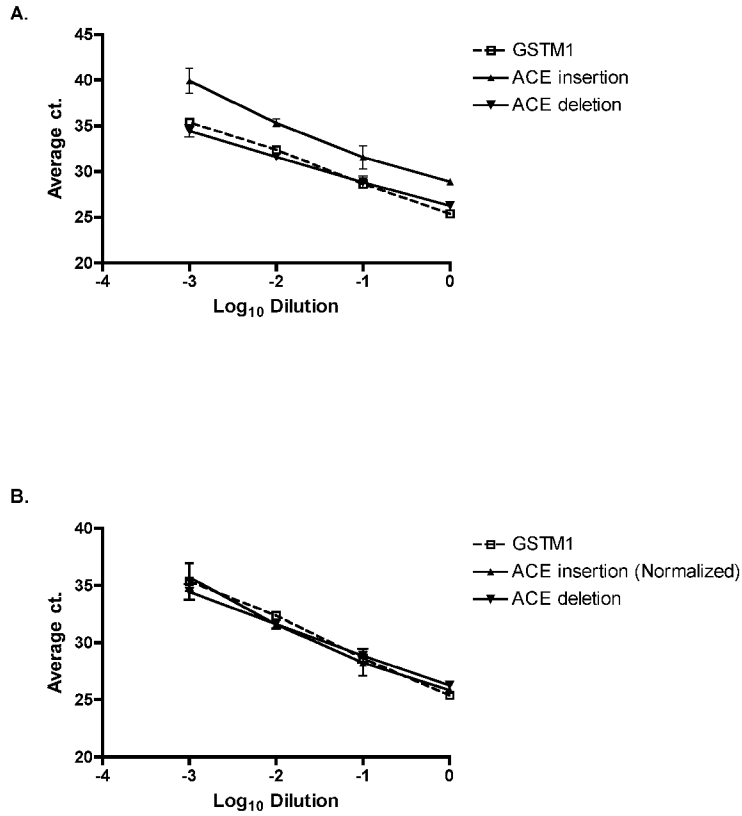


Supplemental Figure

(A) qPCR dilution profiles for ACE insertion, ACE deletion, and GSTM1 reactions using genomic DNA from a participant containing all three alleles. (B) Normalized qPCR dilution profiles. Values represent the average of duplicate measurements and error bars represent the standard error of the mean.



Supplemental Table: Primer and Probe Sequences, from Lo et al. [4]

Primer	Sequence
GST-194F	5' -GGA GAA GAT TCG TGT GGA CA-3'
GST-273R2	5' -CTG GAT TGT AGC AGA TCA TAC-3'
ACE-1721F	5' -GCT GGG ATT ACA GGC GTG ATA-3'
ACE-1428F	5' -CCA TTT CTC TAG ACC TGC TGC C-3'
ACE-1826R	5' -GCC CTT AGC TCA CCT CTG CTT-3'
Probe	Sequence
GST-215T	5' FAM-TTT GGA GAA CCA GAC CAT GGA CAA C-TAMRA3'
ACE-1745T	5' FAM-TCA CTT TTA TGT GGT TTC GCC AAT TTT ATT CCA-TAMARA3'