Supplemental Figure 1. Structure of CYP3A7*1C variant



CYP3A7*1C variant is displayed on top; the reference sequence on bottom. The variant is a haplotype of alleles at seven SNPs (boxes), with NCBI dbSNP designations indicated. The thick line at top indicates the region to which our Taqman MBG probes hybridized. The thin line at the bottom indicates the restriction fragment length polymorphism (a SspI site) used to genotype the variant by others (Smit P, et al. J Clin Endocrinol Metab 2005;90:5313-5316). The coordinates at bottom indicate the 60-bp region of the promoter relative to the transcription start site.