

Supplementary Figure 1

A quantile-quantile plot for association of each SNP with warfarin dose adjusted for age, sex and genotypes for *VKORC1*, *CYP2C9**2, *3 and *CYP4F2*. We omitted SNPs from the loci of *VKORC1* (chr16, 30.4–31.5Mbp), *CYP2C9* (chr10, 96.2–97.2Mbp) and *CYP4F2* (chr19, 15.3–16.3Mbp). The excess of small p -value SNPs is subtle: whereas 65.4 SNPs with $p < 0.0002$ are expected, 70 were observed (1.069 times inflated).

