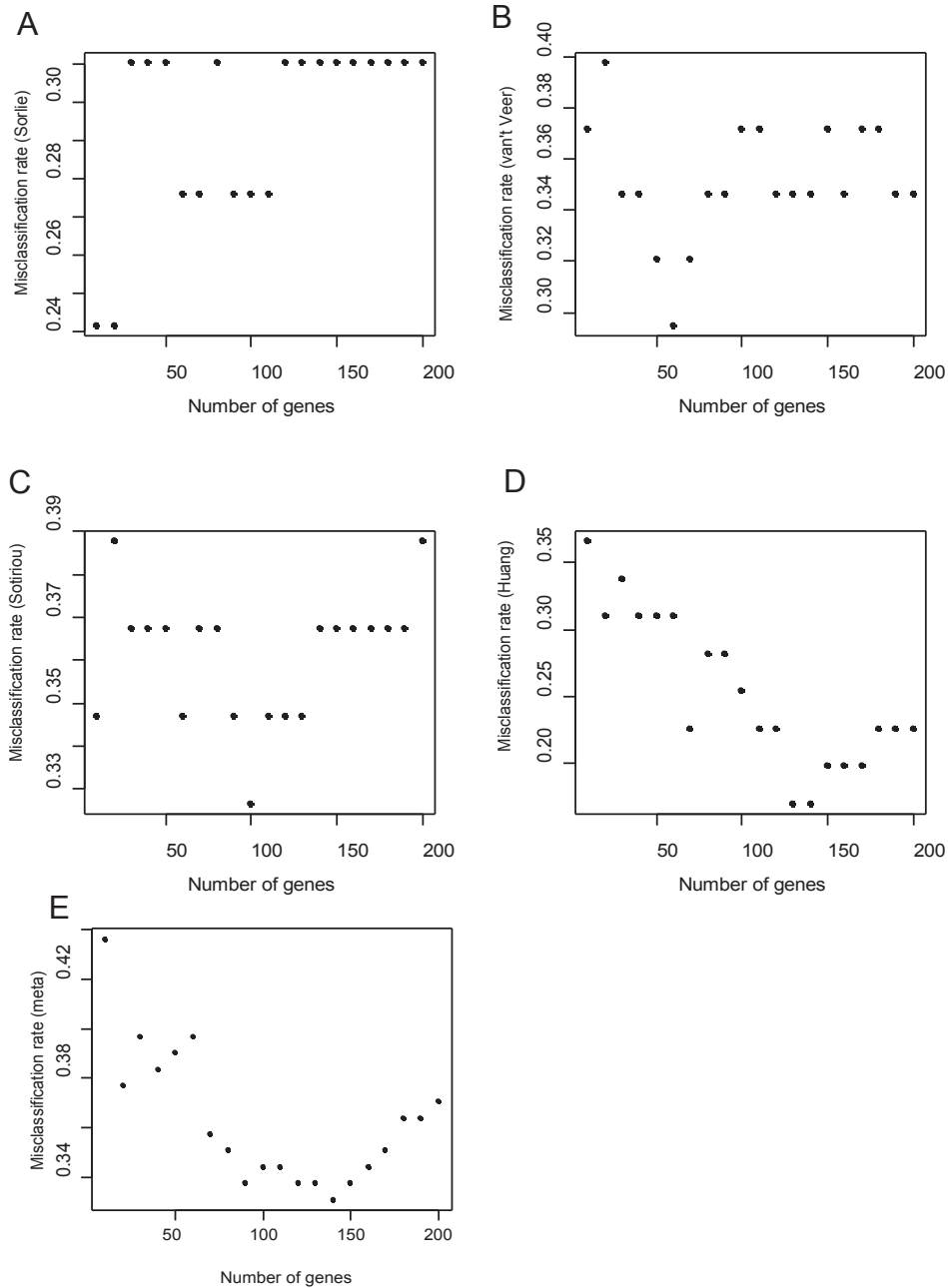




Supplementary Figure 1. Misclassification rates using study-specific gene signatures and the meta-signature of size 10 to 200. A. Sorlie et al. study signature (using 40th percentile of the risk indices as cutoff), B. van't Veer et al. study signature (using 60th percentile RI cutoff), C. Sotiriou et al. study signature (using 50th percentile RI cutoff), D. Huang et al. study signature (using 50th percentile RI cutoff), and E. the meta cohort signatures (using 50th percentile RI cutoff).