Table of Contents for Supplemental Material

Smoking-Associated DNA Methylation Biomarkers and Their Predictive Value for All-Cause and Cardiovascular Mortality

Yan Zhang, Ben Schöttker, Ines Florath, Christian Stock, Katja Butterbach, Bernd Holleczek, Ute Mons, and Hermann Brenner

Table S1. Methylation intensity by characteristics of the study population
Table S2. Spearman correlation coefficients between methylation at 9 CpG sites
Table S3. Association between smoking behavior and methylation intensity
Table S4. Methylation intensity and smoking in relation to all-cause mortality
Table S5. Sex-specific association of the methylation score and smoking with all-cause mortality
Table S6. Evaluation of the SCORE and methylation intensity in prediction of fatal cardiovascular disease (controlling for batch effects)
Table S7. Evaluation of the SCORE and methylation intensity in prediction of fatal cardiovascular disease (excluding participants with cardiovascular disease at baseline n=216)

Figure S1. Dose-response relationships between smoking behavior and methylation intensity
Figure S2. Kaplan-Meier estimates of survival by methylation quartiles